CODE BOOK

Книга кодов

for / для MIKROLATEST®

STAPHYtest 16

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Introduction

The codebook for STAPHYtest 16 is designed to identify bacteria using the MIKRO-LA-TEST STAPHYtest 16 test kit. The identification, ie interpretation of the result, can be done in three ways:

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

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

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

The codebook contains the identification results using the STAPHYtest 16 set ordered according to the value of the so-called profile code. Sorting results by the value of the profile code allows for easy and fast search. The identification quality evaluation for each profile code is based on the calculation of the probability of the result from the identification matrix data; the evaluation is detailed below.

Creating a profile code

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

VPT	URE	ARG	ORN	BGA	GLR	ESL	NIT	PHS	GAL	SUC	TRE	MAN	XYL	MLT	MNS	LAC	
1			2			3			4			5			6		

The test groups are created in accordance with the sequence of tests in the strip. Three tests are included in all groups.

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

VPT	URE	ARG	ORN	BGA	GLR	ESL	NIT	PHS	GAL	SUC	TRE	MAN	XYL	MLT	MNS	LAC	
1	2	4	1	2	4	1	2	4	1	2	4	1	2	4	1	2	

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

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

VPT +	URE -	ARG -	ORN +	BGA +	GLR +	ESL -	NIT -	PHS +	GAL +	SUC +	TRE -	MAN -	XYL +	MLT +	MNS +	LAC -	
1	0	0	1	2	4	0	0	4	1	2	0	0	2	4	1	0	
1			**7**			**4**			**3**			**6**			**1**		

In this example, a profile **174361** was created.

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

Using the codebook

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

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

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

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

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

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

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

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

List of taxa used to create a codebook

Aerococcus viridans
Dermacoccus nishinomiyaensis
Kocuria kristinae
Kocuria rosea
Kocuria varians
Macrococcus bovicus
Macrococcus carouselicus
Macrococcus caseolyticus
Macrococcus equipersicus
Micrococcus sp.
Rothia mucilaginosa
Staphylococcus arlettae
Staphylococcus aureus subsp. anaerobius
Staphylococcus aureus subsp. aureus
Staphylococcus auricularis
Staphylococcus capitis subsp. capitis
Staphylococcus capitis subsp. ureolyticus
Staphylococcus caprae
Staphylococcus carnosus subsp. carnosus
Staphylococcus carnosus subsp. utilis
Staphylococcus chromogenes
Staphylococcus cohnii subsp. cohnii
Staphylococcus cohnii subsp. urealyticum
Staphylococcus condimenti
Staphylococcus delphini
Staphylococcus epidermidis
Staphylococcus equorum
Staphylococcus felis
Staphylococcus gallinarum
Staphylococcus haemolyticus
Staphylococcus hominis subsp. hominis
Staphylococcus hominis subsp. novobiosepticus
Staphylococcus hyicus
Staphylococcus intermedius
Staphylococcus kloosii
Staphylococcus lentus
Staphylococcus lugdunensis
Staphylococcus lutrae
Staphylococcus muscae
Staphylococcus pasteuri
Staphylococcus petrasii subsp. croceilyticus
Staphylococcus petrasii subsp. jettensis
Staphylococcus petrasii subsp. petrasii
Staphylococcus petrasii subsp. pragensis
Staphylococcus piscifermentans
Staphylococcus saprophyticus subsp. bovis
Staphylococcus saprophyticus subsp. saprophyticus
Staphylococcus schleiferi subsp. coagulans
Staphylococcus schleiferi subsp. schleiferi
Staphylococcus sciuri
Staphylococcus simulans
Staphylococcus vitulinus

Staphylococcus warneri
Staphylococcus xylosus

List of tests used to create a codebook

Basic set of tests

VPT acetoin production
URE urea hydrolysis
ARG arginindihydrolase
ORN ornithindecarboxylase
BGA ß-galactosidase
GLR ß-glucuronidase
ESL aesculine hydrolysis
NIT nitrate reduction
PHS phosphatase production
GAL acid from galactose
SUC acid from saccharose
TRE acid from trehalose
MAN acid from mannitol
XYL acid from D-xylose
MLT acid from maltose
MNS acid from mannose
LAC acid from lactose

Additional (distinguishing) tests

OXI oxidase (Kovács)
PYR pyrrolidonyl arylamidase
CAT catalase production
COA coagulase production
HEM ß-haemolysis (sheep blood)
NOV novobiocin resistence
YEP yellow pigment production

Введение

Кодовая книга для СТАФИтест 16 предназначена для идентификации бактерий с использованием тестового набора MIKRO-LA-TEST СТАФИтест 16. Идентификация, т.е. интерпретация результата, может быть выполнена тремя способами:

- 1. вручную - путем сравнения результата с данными в таблице дифференцирования
- 2. с использованием программы идентификации (например, ErbaExpert)
- 3. используя эту кодовую книгу

Все упомянутые методы идентификации основаны на данных матрицы идентификации (частотной матрицы), то есть таблицы, содержащей таксоны, тесты и проценты положительных реакций конкретных тестов для каждого таксона. Таксоном может быть:

- род - *Morganella sp*.
- вид - *Morganella morganii*
- подвид - *Morganella morganii ssp. Morganii*

Кодовая книга содержит результаты идентификации с использованием набора СТАФИтест 16, упорядоченного в соответствии со значением так называемых кодов профиля. Сортировка результатов по значению кода профиля позволяет легко и быстро искать. Оценка качества идентификации для каждого кода профиля основана на вычислении вероятности результата из данных матрицы идентификации; оценка приведена ниже.

Создание кода профиля:

Для более точной идентификации целесообразно преобразовать полученные результаты каждого теста (+/-) в числовой профиль (восьмеричный код). Тесты сначала делятся на группы:

VPT	URE	ARG	ORN	BGA	GLR	ESL	NIT	PHS	GAL	SUC	TRE	MAN	XYL	MLT	MNS	LAC	
1			2			3			4			5			6		

Теперь получаем 5 групп по три и в одной группе два теста. В каждой группе только каждому положительному тесту даются числовые значения 1, 2 и 4:

первый тест тройки - 1
второй тест тройки - 2
третий тест тройки - 4
всем отрицательным результатам дается значение 0

VPT	URE	ARG	ORN	BGA	GLR	ESL	NIT	PHS	GAL	SUC	TRE	MAN	XYL	MLT	MNS	LAC	
1	2	4	1	2	4	1	2	4	1	2	4	1	2	4	1	2	

При создании кода профиля для конкретного результата необходимо выполнить следующие действия:

• положительным испытаниям присваиваются 1, 2 или 4 значения
• отрицательным испытаниям всегда присваивается нулевое значение
• полученные значения суммируются в каждой тестовой группе
• итоговая группа чисел представляет собой **код профиля**

VPT +	URE -	ARG -	ORN +	BGA +	GLR +	ESL -	NIT -	PHS +	GAL +	SUC +	TRE -	MAN -	XYL +	MLT +	MNS +	LAC -	
1	0	0	1	2	4	0	0	4	1	2	0	0	2	4	1	0	
1			**7**			**4**			**3**			**6**			**1**		

В этом примере был создан профиль **174361**.
Результаты располагаются в кодовой книге в соответствии со значением кода профиля в порядке возрастания.

ИСПОЛЬЗОВАНИЕ ИНДЕКСА ЧИСЛОВЫХ ПРОФИЛЕЙ

Найдите числовой профиль в индексе; профили систематизированы в регистре в порядке возрастания величин чисел.
Профиль сопровождается следующей информацией:

- значения 2-х индексов для каждого таксона:
  а) процент идентификации (% id) – показатель, оценивающий насколько близко полученный профиль соответствует идентифицированному таксону по сравнению с другими таксонами, включенными в банк данных;
  таксоны систематизированы в соответствии с их величинами % id.
  б) Т индекс (Tin) – показывает насколько близко профиль соответствует большинству типичных реакций данного таксона. Его величина варьирует от 0 до 1 и обратно пропорциональна количеству атипичных тестов.
- перечень атипичных тестов (T. against) для первого из перечисленных таксонов с указанием процента положительных реакций, заложенных в базу данных;
- перечень дополнительных тестов – если таксоны не достаточно хорошо идентифицируются с помощью СТАФИтест 16; дополнительные тесты приводятся с процентом положительных реакций, заложенных в базу данных;
- пояснения в случае, если качество идентификации выходит за рамки вышеперечисленных показателей;
- оценка качества идентификации, исходя из значений %id и T-index:

Процент идентификации
%id >= 99 – таксон отлично отличается от других таксонов
%id >= 95 - таксон очень хорошо отличается от других таксонов
%id >= 90 - таксон удовлетворительно отличается от других таксонов
%id < 90 - таксон неудовлетворительно отличается от других таксонов

Т-индекс
Т-индекс >= 0.75- типичный штамм
Т-индекс >= 0.50- менее типичный штамм
Т-индекс >= 0.25- атипичный штамм
Т-индекс < 0.25- полностью атипичный штамм

Т-индекс	**%id**	**качество идентификации**
>= 0.75	>= 0.99	ОТЛИЧНАЯ ИДЕНТИФИКАЦИЯ
>= 0.50	>= 0.95	ОЧЕНЬ ХОРОШАЯ ИДЕНТИФИКАЦИЯ
>= 0.25	>= 0.90	ХОРОШАЯ ИДЕНТИФИКАЦИЯ для первого таксона
>= 0	>= 0.90	ПРИЕМЛИМАЯ ИДЕНТИФИКАЦИЯ для первого таксона
	< 0.90	НИЗКОЕ РАЗЛИЧИЕ - выбранные таксоны относятся к разным родам ИДЕНТИФИКАЦИЯ ДО РОДА - выбраны более одного таксона того же рода ИДЕНТИФИКАЦИЯ ДО ВИДА - выбраны более чем один вид и подвид в одном образце

В случае, если профиль не найден в индексе профилей, то возможно:

- профиль является совсем атипичным и частота его встречаемости очень низкая
- профиль относится к таксону, который не включен в базу данных.

Список таксонов, используемых для создания кодовой книги

Aerococcus viridans
Dermacoccus nishinomiyaensis
Kocuria kristinae
Kocuria rosea
Kocuria varians
Macrococcus bovicus
Macrococcus carouselicus
Macrococcus caseolyticus
Macrococcus equipersicus
Micrococcus sp.
Rothia mucilaginosa
Staphylococcus arlettae
Staphylococcus aureus subsp. anaerobius
Staphylococcus aureus subsp. aureus
Staphylococcus auricularis
Staphylococcus capitis subsp. capitis
Staphylococcus capitis subsp. ureolyticus
Staphylococcus caprae
Staphylococcus carnosus subsp. carnosus
Staphylococcus carnosus subsp. utilis
Staphylococcus chromogenes
Staphylococcus cohnii subsp. cohnii
Staphylococcus cohnii subsp. urealyticum
Staphylococcus condimenti
Staphylococcus delphini
Staphylococcus epidermidis
Staphylococcus equorum
Staphylococcus felis
Staphylococcus gallinarum
Staphylococcus haemolyticus
Staphylococcus hominis subsp. hominis
Staphylococcus hominis subsp. novobiosepticus
Staphylococcus hyicus
Staphylococcus intermedius
Staphylococcus kloosii
Staphylococcus lentus
Staphylococcus lugdunensis
Staphylococcus lutrae
Staphylococcus muscae
Staphylococcus pasteuri
Staphylococcus petrasii subsp. croceilyticus
Staphylococcus petrasii subsp. jettensis
Staphylococcus petrasii subsp. petrasii
Staphylococcus petrasii subsp. pragensis
Staphylococcus piscifermentans
Staphylococcus saprophyticus subsp. bovis
Staphylococcus saprophyticus subsp. saprophyticus
Staphylococcus schleiferi subsp. coagulans
Staphylococcus schleiferi subsp. schleiferi
Staphylococcus sciuri
Staphylococcus simulans
Staphylococcus vitulinus
Staphylococcus warneri
Staphylococcus xylosus

Список тестов, используемых для создания кода профиля

Базовый набор тестов

VPT	Ацетоин
URE	Уреаза
ARG	Аргинин
ORN	Орнитин
BGA	β-Галактозидаза
GLR	β-Глюкуронидаза
ESL	Эскулин
NIT	Нитраты
PHS	Фосфатаза
GAL	Галактоза
SUC	Сахароза
TRE	Трегалоза
MAN	Маннитол
XYL	Ксилоза
MLT	Мальтоза
MNS	Манноза
LAC	Лактоза

Дополнительные (отличительные) тесты

OXI	Цитохромоксидаза
PYR	Пирролидониларилами даза
CAT	производство каталазы
COA	производство коагулазы
HEM	ß- гемолиз (овечья кровь)
NOV	резистентность к нообибиоцину
YEP	производство желтого пигмента

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
000000	**Low discrimination**							**000000**
Micrococcus sp.		71.00	1.00					
Dermacoccus nishinomiyaensis		25.22	1.00					
Kocuria rosea		3.23	0.80					
000001	**Low discrimination**							**000001**
Micrococcus sp.		73.91	0.65	MNS 2				
Dermacoccus nishinomiyaensis		12.99	0.59					
Kocuria rosea		8.66	0.54					
000002	**Low discrimination**							**000002**
					OXI			
Micrococcus sp.		64.48	0.59	LAC 1	89			
Dermacoccus nishinomiyaensis		22.90	0.59		93			
Kocuria varians		9.22	0.59		13			
000010	**Low discrimination**							**000010**
					NOV			
Macrococcus bovicus		72.88	0.89		99			
Micrococcus sp.		10.75	0.59		1			
Macrococcus equipersicus		6.13	0.68		99			
000011	**Low discrimination**							**000011**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		77.42	0.64	NIT 89	1	1	1	
Macrococcus bovicus		9.50	0.48		99	99	99	
Staphylococcus cohnii subsp. cohnii		5.12	0.49		1	99	12	
000012	**Low discrimination**							**000012**
					NOV			
Macrococcus bovicus		70.49	0.48	LAC 1	99			
Micrococcus sp.		10.39	0.18		1			
Macrococcus equipersicus		5.93	0.27		99			
000013	**Low discrimination**							**000013**
					OXI	PYR	NOV	
Staphylococcus capitis subsp. capitis		66.25	0.29	NIT 89	1	1	1	
Aerococcus viridans		26.02	0.42	LAC 2	1	95	1	
Macrococcus bovicus		4.02	0.07		99	1	99	
000020	**Low discrimination**							**000020**
Kocuria rosea		55.65	0.73	NIT 90				
Micrococcus sp.		28.87	0.59					
Dermacoccus nishinomiyaensis		10.26	0.59					
000021	**Low discrimination**							**000021**
					OXI			
Kocuria rosea		71.62	0.47	NIT 90	35			
Micrococcus sp.		14.41	0.24	MNS 5	89			
Kocuria varians		10.93	0.39		13			
000022	**Low discrimination**							**000022**
					OXI			
Kocuria varians		59.78	0.49		13			
Kocuria rosea		23.17	0.32		35			
Micrococcus sp.		12.02	0.18		89			
000023	**Low discrimination**							**000023**
					PYR			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		63.44	0.27	MNS 8	10		
Aerococcus viridans		21.68	0.32		95		
Kocuria rosea		14.88	0.05		1		
000030	**Low discrimination**						**000030**
					NOV	YEP	
Kocuria rosea		66.49	0.53	NIT 90	1	1	
Macrococcus bovicus		21.27	0.48	MAN 10	99	99	
Staphylococcus vitulinus		5.38	0.29		99	75	
000031	**Low discrimination**						**000031**
					NOV	YEP	
Kocuria rosea		61.00	0.27	NIT 90	1	1	
Staphylococcus capitis subsp. capitis		30.54	0.23	MAN 10	1	1	
Macrococcus bovicus		3.75	0.07	MNS 5	99	99	
000033	**Good identification**						**000033**
Aerococcus viridans		100.0	0.42	ESL 90			
000040	**Low discrimination**						**000040**
					NOV	YEP	
Micrococcus sp.		88.85	0.77	MLT 7	1	41	
Dermacoccus nishinomiyaensis		4.24	0.59		80	1	
Macrococcus bovicus		1.91	0.55		99	99	
000041	**Low discrimination**						**000041**
					NOV	OXI	
Micrococcus sp.		55.27	0.42	MLT 7	1	89	
Staphylococcus cohnii subsp. cohnii		33.42	0.57	MNS 2	99	1	
Staphylococcus auricularis		3.57	0.33		1	1	
000042	**Low discrimination**						**000042**
					OXI		
Micrococcus sp.		59.56	0.36	MLT 7	89		
Kocuria varians		24.83	0.45	LAC 1	13		
Staphylococcus auricularis		7.78	0.33		1		
000043	**Low discrimination**						**000043**
					OXI	PYR	
Aerococcus viridans		46.70	0.44	ESL 90	1	95	
Kocuria varians		21.35	0.23	TRE 95	13	10	
Micrococcus sp.		12.02	0.01		89	63	
000050	**Good identification**						**000050**
					OXI	YEP	
Macrococcus bovicus		90.11	0.96		99	99	
Macrococcus equipersicus		8.78	0.77		99	99	
Staphylococcus cohnii subsp. cohnii		0.62	0.58		1	12	
000051	**Low discrimination**						**000051**
					OXI	YEP	
Macrococcus bovicus		52.96	0.55	MNS 1	99	99	
Staphylococcus cohnii subsp. cohnii		38.76	0.59		1	12	
Macrococcus equipersicus		5.16	0.36		99	99	
000052	**Low discrimination**						**000052**
					OXI	NOV	
Macrococcus bovicus		86.28	0.55	LAC 1	99	99	
Macrococcus equipersicus		8.40	0.36		99	99	
Aerococcus viridans		4.14	0.47		1	1	
000053	**Low discrimination**						**000053**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI	NOV	PYR	YEP	
Aerococcus viridans		86.49	0.54	ESL 90	1	1	95	1	
Macrococcus bovicus		7.80	0.14	TRE 95	99	99	1	99	
Staphylococcus cohnii subsp. cohnii		5.71	0.18		1	99	1	12	
000060	**Low discrimination**								**000060**
					OXI				
Micrococcus sp.		53.05	0.36	XYL 1	89				
Kocuria varians		27.55	0.47	MLT 7	13				
Kocuria rosea		13.72	0.32		35				
000062	**Low discrimination**								**000062**
					OXI	PYR			
Kocuria varians		76.87	0.36		13	10			
Aerococcus viridans		17.84	0.37		1	95			
Micrococcus sp.		5.30	-.05		89	63			
000063	**Low discrimination**								**000063**
					PYR				
Aerococcus viridans		86.16	0.44	ESL 90	95				
Kocuria varians		13.84	0.14	TRE 95	10				
000070	**Low discrimination**								**000070**
					NOV	YEP			
Macrococcus bovicus		88.47	0.55	XYL 1	99	99			
Macrococcus equipersicus		8.62	0.36		99	99			
Kocuria rosea		1.20	0.12		1	1			
000071	**Low discrimination**								**000071**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		41.57	0.34	ESL 90	1	1	95	1	
Macrococcus bovicus		33.74	0.14	TRE 95	99	99	1	99	
Staphylococcus cohnii subsp. cohnii		24.69	0.18	LAC 90	1	99	1	12	
000072	**Low discrimination**								**000072**
					OXI	NOV			
Aerococcus viridans		82.62	0.47	ESL 90	1	1			
Macrococcus bovicus		17.38	0.14	TRE 95	99	99			
000073	**Very good identification**								**000073**
Aerococcus viridans		100.0	0.54	ESL 90					
000100	**Low discrimination**								**000100**
					OXI				
Dermacoccus nishinomiyaensis		78.02	0.85		93				
Micrococcus sp.		11.65	0.59		89				
Kocuria varians		8.86	0.73		13				
000101	**Low discrimination**								**000101**
					OXI				
Dermacoccus nishinomiyaensis		41.94	0.44	MNS 1	93				
Kocuria varians		40.99	0.52		13				
Micrococcus sp.		12.65	0.24		89				
000102	**Low discrimination**								**000102**
					OXI				
Kocuria varians		72.85	0.62		13				
Dermacoccus nishinomiyaensis		22.98	0.44		93				
Micrococcus sp.		3.43	0.18		89				
000103	**Low discrimination**								**000103**
					OXI	PYR			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		89.55	0.40	MNS 8	13	10		
Aerococcus viridans		7.17	0.32		1	95		
Dermacoccus nishinomiyaensis		3.28	0.03		93	45		
000110	**Low discrimination**							**000110**
					YEP			
Macrococcus bovicus		89.02	0.81		99			
Macrococcus equipersicus		7.48	0.61		99			
Dermacoccus nishinomiyaensis		2.07	0.44		1			
000111	**Low discrimination**							**000111**
					OXI	NOV	YEP	
Macrococcus bovicus		67.20	0.40	MNS 1	99	99	99	
Staphylococcus capitis subsp. capitis		12.91	0.23		1	1	1	
Staphylococcus cohnii subsp. cohnii		9.40	0.29		1	99	12	
000112	**Low discrimination**							**000112**
					NOV	OXI		
Macrococcus bovicus		80.23	0.40	LAC 1	99	99		
Macrococcus equipersicus		6.74	0.20		99	99		
Kocuria varians		5.92	0.21		1	13		
000113	**Acceptable identification**							**000113**
					OXI	NOV		
Aerococcus viridans		93.79	0.42	ESL 90	1	1		
Macrococcus bovicus		6.21	-.01	TRE 95	99	99		
000120	**Low discrimination**							**000120**
					OXI			
Kocuria varians		73.19	0.64		13			
Dermacoccus nishinomiyaensis		18.54	0.44		93			
Kocuria rosea		5.33	0.32		35			
000121	**Acceptable identification**							**000121**
					OXI			
Kocuria varians		93.15	0.42	MNS 8	13			
Kocuria rosea		4.11	0.05		35			
Dermacoccus nishinomiyaensis		2.74	0.03		93			
000122	**Good identification**							**000122**
					OXI	PYR		
Kocuria varians		98.27	0.53		13	10		
Dermacoccus nishinomiyaensis		0.89	0.03		93	45		
Aerococcus viridans		0.83	0.24		1	95		
000123	**Low discrimination**							**000123**
					PYR			
Kocuria varians		81.45	0.31	MNS 8	10			
Aerococcus viridans		18.55	0.32		95			
000130	**Low discrimination**							**000130**
					NOV	OXI		
Macrococcus bovicus		78.56	0.40	XYL 1	99	99		
Kocuria varians		7.22	0.23		1	13		
Macrococcus equipersicus		6.60	0.20		99	99		
000132	**Low discrimination**							**000132**
					NOV	PYR	OXI	
Aerococcus viridans		64.47	0.34	ESL 90	1	95	1	
Kocuria varians		25.57	0.12	TRE 95	1	10	13	
Macrococcus bovicus		9.97	-.01		99	1	99	
000133	**Good identification**							**000133**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.42	ESL 90			
000140	**Low discrimination**						**000140**
					NOV	OXI	
Kocuria varians		38.51	0.60		1	13	
Micrococcus sp.		17.37	0.36		1	89	
Macrococcus bovicus		15.81	0.48		99	99	
000141	**Low discrimination**						**000141**
					NOV	OXI	
Kocuria varians		47.66	0.38	MNS 8	1	13	
Staphylococcus cohnii subsp. cohnii		33.55	0.37		99	1	
Micrococcus sp.		5.04	0.01		1	89	
000142	**Low discrimination**						**000142**
					PYR		
Kocuria varians		88.47	0.49		10		
Aerococcus viridans		4.81	0.37		95		
Staphylococcus auricularis		2.05	0.14		40		
000143	**Low discrimination**						**000143**
					PYR		
Aerococcus viridans		59.32	0.44	ESL 90	95		
Kocuria varians		40.68	0.27	TRE 95	10		
000150	**Good identification**						**000150**
					OXI	YEP	
Macrococcus bovicus		90.93	0.89		99	99	
Macrococcus equipersicus		8.86	0.70		99	99	
Staphylococcus cohnii subsp. cohnii		0.16	0.39		1	12	
000151	**Low discrimination**						**000151**
					OXI	YEP	
Macrococcus bovicus		75.98	0.48	MNS 1	99	99	
Staphylococcus cohnii subsp. cohnii		14.42	0.39		1	12	
Macrococcus equipersicus		7.40	0.28		99	99	
000152	**Low discrimination**						**000152**
					OXI	NOV	
Macrococcus bovicus		81.94	0.48	LAC 1	99	99	
Aerococcus viridans		9.18	0.47		1	1	
Macrococcus equipersicus		7.98	0.28		99	99	
000153	**Very good identification**						**000153**
					OXI	NOV	
Aerococcus viridans		96.28	0.54	ESL 90	1	1	
Macrococcus bovicus		3.72	0.07	TRE 95	99	99	
000160	**Good identification**						**000160**
					NOV	OXI	
Kocuria varians		95.49	0.51		1	13	
Micrococcus sp.		1.24	-.05		1	89	
Macrococcus bovicus		1.13	0.07		99	99	
000161	**Low discrimination**						**000161**
					PYR		
Kocuria varians		88.49	0.29	MNS 8	10		
Aerococcus viridans		11.51	0.25		95		
000162	**Low discrimination**						**000162**
					PYR		
Kocuria varians		86.60	0.39		10		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		13.40	0.37		95		
000163	**Low discrimination**						**000163**
					PYR		
Aerococcus viridans		80.59	0.44	ESL 90	95		
Kocuria varians		19.42	0.17	TRE 95	10		
000170	**Low discrimination**						**000170**
					OXI	NOV	
Macrococcus bovicus		89.41	0.48	XYL 1	99	99	
Macrococcus equipersicus		8.71	0.28		99	99	
Aerococcus viridans		1.11	0.27		1	1	
000171	**Low discrimination**						**000171**
					OXI	NOV	
Aerococcus viridans		74.20	0.34	ESL 90	1	1	
Macrococcus bovicus		25.80	0.07	TRE 95	99	99	
000172	**Acceptable identification**						**000172**
					OXI	NOV	
Aerococcus viridans		91.73	0.47	ESL 90	1	1	
Macrococcus bovicus		8.27	0.07	TRE 95	99	99	
000173	**Very good identification**						**000173**
Aerococcus viridans		100.0	0.54	ESL 90			
000200	**Low discrimination**						**000200**
Micrococcus sp.		76.35	0.79	SUC 9			
Dermacoccus nishinomiyaensis		17.50	0.75				
Kocuria rosea		3.90	0.61				
000201	**Low discrimination**						**000201**
					OXI		
Micrococcus sp.		55.23	0.45	SUC 9	89		
Staphylococcus capitis subsp. capitis		21.58	0.47	MNS 2	1		
Kocuria rosea		7.27	0.34		35		
000202	**Low discrimination**						**000202**
					OXI		
Micrococcus sp.		65.00	0.38	SUC 9	89		
Dermacoccus nishinomiyaensis		14.90	0.34	LAC 1	93		
Kocuria varians		12.81	0.41		13		
000210	**Low discrimination**						**000210**
					NOV		
Macrococcus bovicus		63.48	0.76		99		
Staphylococcus vitulinus		23.76	0.62		99		
Micrococcus sp.		3.70	0.38		1		
000211	**Very good identification**						**000211**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		97.02	0.70	NIT 89	1	1	
Macrococcus bovicus		1.60	0.35		99	99	
Staphylococcus vitulinus		0.60	0.21		99	99	
000212	**Low discrimination**						**000212**
					OXI	NOV	
Macrococcus bovicus		59.99	0.35	LAC 1	99	99	
Staphylococcus vitulinus		22.45	0.21		99	99	
Aerococcus viridans		6.72	0.34		1	1	
000213	**Low discrimination**						**000213**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	PYR	
Staphylococcus capitis subsp. capitis		81.77	0.35	NIT 89	1	1	
Aerococcus viridans		17.29	0.42	LAC 2	1	95	
Kocuria kristinae		0.93	-.06		99	93	
000220	**Low discrimination**						**000220**
					OXI		
Kocuria rosea		56.47	0.53	NIT 90	35		
Micrococcus sp.		26.08	0.38	SUC 10	89		
Kocuria varians		6.41	0.43		13		
000221	**Low discrimination**						**000221**
					OXI		
Kocuria rosea		69.62	0.27	NIT 90	35		
Kocuria varians		13.04	0.21	SUC 10	13		
Micrococcus sp.		12.47	0.03	MNS 5	89		
000222	**Low discrimination**						**000222**
					OXI		
Kocuria varians		64.31	0.31	SUC 12	13		
Kocuria rosea		20.31	0.12		35		
Micrococcus sp.		9.38	-.03		89		
000223	**Low discrimination**						**000223**
					PYR		
Aerococcus viridans		71.47	0.32	ESL 90	95		
Kocuria varians		28.53	0.10	TRE 95	10		
000230	**Very good identification**						**000230**
					NOV	YEP	
Staphylococcus vitulinus		97.52	0.70	NIT 91	99	75	
Kocuria rosea		1.35	0.34		1	1	
Macrococcus bovicus		0.97	0.35		99	99	
000231	**Low discrimination**						**000231**
					NOV	OXI	
Staphylococcus vitulinus		58.63	0.29	NIT 91	99	99	
Staphylococcus capitis subsp. capitis		35.45	0.29	MNS 1	1	1	
Kocuria rosea		4.24	0.07		1	35	
000232	**Low discrimination**						**000232**
					NOV	PYR	
Staphylococcus vitulinus		88.92	0.29	NIT 91	99	1	
Aerococcus viridans		9.85	0.34	LAC 1	1	95	
Kocuria rosea		1.23	-.08		1	1	
000233	**Good identification**						**000233**
Aerococcus viridans		100.0	0.42	ESL 90			
000240	**Low discrimination**						**000240**
					COA	OXI	
Micrococcus sp.		44.80	0.56	SUC 9	1	89	
Staphylococcus aureus subsp. anaerobius		40.52	0.57	MLT 7	99	1	
Staphylococcus auricularis		4.96	0.52		1	1	
000241	**Low discrimination**						**000241**
					OXI		
Kocuria kristinae		85.20	0.54	VPT 90	99		
Micrococcus sp.		6.13	0.21	TRE 95	89		
Staphylococcus auricularis		3.70	0.33		1		
000242	**Low discrimination**						**000242**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	OXI		
Staphylococcus auricularis		27.24	0.33	NIT 90	1	1		
Micrococcus sp.		22.35	0.15	TRE 92	1	89		
Staphylococcus aureus subsp. anaerobius		20.22	0.16	LAC 10	99	1		
000243	**Low discrimination**							**000243**
					OXI			
Kocuria kristinae		87.30	0.42	VPT 90	99			
Aerococcus viridans		9.98	0.44	TRE 95	1			
Staphylococcus auricularis		1.68	0.13		1			
000250	**Very good identification**							**000250**
					NOV			
Macrococcus bovicus		95.68	0.84		99			
Macrococcus equipersicus		3.69	0.56		99			
Micrococcus sp.		0.18	0.15		1			
000251	**Low discrimination**							**000251**
					OXI	NOV	PYR	
Macrococcus bovicus		62.58	0.43	MNS 1	99	99	1	
Staphylococcus capitis subsp. capitis		16.41	0.29		1	1	1	
Kocuria kristinae		13.63	0.26		99	1	93	
000252	**Low discrimination**							**000252**
					OXI	NOV		
Macrococcus bovicus		81.26	0.43	LAC 1	99	99		
Aerococcus viridans		15.61	0.47		1	1		
Macrococcus equipersicus		3.13	0.15		99	99		
000253	**Low discrimination**							**000253**
					OXI	NOV	PYR	
Aerococcus viridans		87.41	0.54	ESL 90	1	1	95	
Kocuria kristinae		10.62	0.13	TRE 95	99	1	93	
Macrococcus bovicus		1.97	0.02		99	99	1	
000262	**Low discrimination**							**000262**
					PYR			
Aerococcus viridans		62.98	0.37	ESL 90	95			
Kocuria varians		37.02	0.18	TRE 95	10			
000263	**Acceptable identification**							**000263**
					OXI			
Aerococcus viridans		91.88	0.44	ESL 90	1			
Kocuria kristinae		8.12	0.01	TRE 95	99			
000270	**Low discrimination**							**000270**
Macrococcus bovicus		66.69	0.43	XYL 1				
Staphylococcus vitulinus		28.92	0.29					
Macrococcus equipersicus		2.57	0.15					
000271	**Low discrimination**							**000271**
					OXI	NOV		
Aerococcus viridans		83.13	0.34	ESL 90	1	1		
Macrococcus bovicus		16.87	0.02	TRE 95	99	99		
000272	**Low discrimination**							**000272**
					OXI	NOV	YEP	
Aerococcus viridans		86.41	0.47	ESL 90	1	1	1	
Staphylococcus arlettae		9.05	0.15	TRE 95	1	99	99	
Macrococcus bovicus		4.55	0.02		99	99	99	
000273	**Good identification**							**000273**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		94.39	0.54	ESL 90	1	1		
Staphylococcus arlettae		5.62	0.18	TRE 95	99	99		
000300	**Low discrimination**							**000300**
					OXI			
Dermacoccus nishinomiyaensis		60.08	0.61	SUC 6	93			
Kocuria varians		14.57	0.56		13			
Micrococcus sp.		13.90	0.38		89			
000301	**Low discrimination**							**000301**
					OXI	PYR		
Kocuria varians		35.74	0.34	SUC 12	13	10		
Kocuria kristinae		30.68	0.25	MNS 8	99	93		
Dermacoccus nishinomiyaensis		17.12	0.20		93	45		
000302	**Low discrimination**							**000302**
					OXI			
Kocuria varians		80.61	0.44	SUC 12	13			
Dermacoccus nishinomiyaensis		11.90	0.20		93			
Micrococcus sp.		2.75	-.03		89			
000310	**Good identification**							**000310**
Macrococcus bovicus		90.26	0.69					
Macrococcus carouselicus		5.08	0.39					
Macrococcus equipersicus		3.00	0.40					
000311	**Low discrimination**							**000311**
					OXI	PYR	NOV	
Staphylococcus capitis subsp. capitis		54.30	0.29	NIT 89	1	1	1	
Macrococcus bovicus		38.04	0.28	GAL 1	99	1	99	
Aerococcus viridans		2.58	0.22		1	95	1	
000312	**Low discrimination**							**000312**
					OXI	NOV		
Macrococcus bovicus		71.88	0.28	LAC 1	99	99		
Aerococcus viridans		18.79	0.34		1	1		
Macrococcus carouselicus		4.05	-.02		99	99		
000313	**Good identification**							**000313**
Aerococcus viridans		100.0	0.42	ESL 90				
000320	**Low discrimination**							**000320**
					OXI			
Kocuria varians		81.76	0.46	SUC 12	13			
Dermacoccus nishinomiyaensis		9.69	0.20		93			
Kocuria rosea		4.86	0.12		35			
000322	**Acceptable identification**							**000322**
					PYR			
Kocuria varians		94.14	0.35	SUC 12	10			
Aerococcus viridans		5.86	0.24		95			
000323	**Low discrimination**							**000323**
					PYR			
Aerococcus viridans		62.55	0.32	ESL 90	95			
Kocuria varians		37.45	0.13	TRE 95	10			
000330	**Low discrimination**							**000330**
Staphylococcus vitulinus		66.82	0.29	NIT 91				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		28.30	0.28	GAL 1				
Macrococcus carouselicus		1.59	-.02					
000332	**Good identification**							**000332**
Aerococcus viridans		100.0	0.34	ESL 90				
000333	**Good identification**							**000333**
Aerococcus viridans		100.0	0.42	ESL 90				
000340	**Low discrimination**							**000340**
					NOV	OXI		
Kocuria varians		28.68	0.42	SUC 12	1	13		
Macrococcus bovicus		21.58	0.35		99	99		
Staphylococcus auricularis		11.43	0.33		1	1		
000341	**Acceptable identification**							**000341**
					NOV	OXI	PYR	
Kocuria kristinae		93.24	0.44	VPT 90	1	99	93	
Kocuria varians		2.65	0.20	TRE 95	1	13	10	
Staphylococcus hominis subsp. novobiosepticus		1.87	0.07		99	1	1	
000342	**Low discrimination**							**000342**
					NOV	PYR		
Kocuria varians		48.00	0.31	SUC 12	1	10		
Staphylococcus hominis subsp. novobiosepticus		22.50	0.14		99	1		
Aerococcus viridans		19.13	0.37		1	95		
000343	**Low discrimination**							**000343**
					OXI	NOV	PYR	
Kocuria kristinae		69.63	0.32	VPT 90	99	1	93	
Aerococcus viridans		23.87	0.44	TRE 95	1	1	95	
Staphylococcus hominis subsp. novobiosepticus		3.01	0.02		1	99	1	
000350	**Very good identification**							**000350**
					OXI	NOV		
Macrococcus bovicus		96.22	0.76		99	99		
Macrococcus equipersicus		3.71	0.49		99	99		
Aerococcus viridans		0.05	0.27		1	1		
000351	**Low discrimination**							**000351**
					OXI	NOV	PYR	
Macrococcus bovicus		75.52	0.35	MNS 1	99	99	1	
Kocuria kristinae		12.80	0.16		99	1	93	
Aerococcus viridans		8.77	0.34		1	1	95	
000352	**Low discrimination**							**000352**
					OXI	NOV		
Macrococcus bovicus		67.27	0.35	LAC 1	99	99		
Aerococcus viridans		30.14	0.47		1	1		
Macrococcus equipersicus		2.59	0.08		99	99		
000353	**Very good identification**							**000353**
					OXI			
Aerococcus viridans		96.11	0.54	ESL 90	1			
Kocuria kristinae		3.89	0.03	TRE 95	99			
000360	**Good identification**							**000360**
					PYR			
Kocuria varians		96.57	0.33	SUC 12	10			
Aerococcus viridans		3.43	0.17		95			
000362	**Low discrimination**							**000362**

Code Taxon	ID score	T index	tests against	Differentiation tests			Code
				PYR			
Aerococcus viridans	53.15	0.37	ESL 90	95			
Kocuria varians	46.85	0.21	TRE 95	10			
000363 **Good identification**							**000363**
Aerococcus viridans	100.0	0.44	ESL 90				
000370 **Acceptable identification**							**000370**
				OXI	NOV		
Macrococcus bovicus	91.89	0.35	XYL 1	99	99		
Aerococcus viridans	4.58	0.27		1	1		
Macrococcus equipersicus	3.54	0.08		99	99		
000371 **Good identification**							**000371**
Aerococcus viridans	100.0	0.34	ESL 90				
000372 **Low discrimination**							**000372**
				NOV	YEP		
Aerococcus viridans	62.76	0.47	ESL 90	1	1		
Staphylococcus arlettae	37.24	0.31	TRE 95	99	99		
000373 **Low discrimination**							**000373**
				NOV	YEP		
Aerococcus viridans	74.79	0.54	ESL 90	1	1		
Staphylococcus arlettae	25.21	0.34	TRE 95	99	99		
000400 **Low discrimination**							**000400**
				NOV	OXI		
Staphylococcus cohnii subsp. cohnii	44.11	0.87		99	1		
Macrococcus carouselicus	19.15	0.69		99	99		
Micrococcus sp.	16.62	0.59		1	89		
000401 **Very good identification**							**000401**
				NOV	OXI		
Staphylococcus cohnii subsp. cohnii	96.59	0.88		99	1		
Staphylococcus auricularis	1.63	0.48		1	1		
Micrococcus sp.	0.69	0.24		1	89		
000402 **Low discrimination**							**000402**
				NOV	PYR		
Staphylococcus auricularis	38.05	0.48	NIT 90	1	40		
Staphylococcus cohnii subsp. cohnii	21.05	0.46	LAC 10	99	1		
Aerococcus viridans	9.55	0.51		1	95		
000403 **Low discrimination**							**000403**
				NOV	PYR		
Staphylococcus cohnii subsp. cohnii	44.79	0.47	LAC 1	99	1		
Aerococcus viridans	43.75	0.58		1	95		
Staphylococcus auricularis	8.30	0.28		1	40		
000410 **Low discrimination**							**000410**
				OXI	YEP		
Macrococcus bovicus	68.32	0.92		99	99		
Staphylococcus cohnii subsp. cohnii	22.68	0.89		1	12		
Macrococcus carouselicus	4.95	0.65		99	67		
000411 **Very good identification**							**000411**
				OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii	96.38	0.90		1	99	12	
Macrococcus bovicus	2.71	0.51		99	99	99	
Staphylococcus capitis subsp. capitis	0.30	0.29		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
000412	**Low discrimination**								**000412**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		54.73	0.51	LAC 1	99	99	1	99	
Aerococcus viridans		19.42	0.60		1	1	95	1	
Staphylococcus cohnii subsp. cohnii		18.17	0.48		1	99	1	12	
000413	**Low discrimination**								**000413**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		69.13	0.68	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		30.03	0.49		1	99	1	12	
Macrococcus bovicus		0.84	0.10		99	99	1	99	
000420	**Low discrimination**								**000420**
					NOV				
Staphylococcus cohnii subsp. cohnii		28.47	0.46	XYL 1	99				
Kocuria rosea		20.68	0.32		1				
Kocuria varians		14.24	0.37		1				
000421	**Low discrimination**								**000421**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		83.28	0.47	XYL 1	99	1			
Aerococcus viridans		9.04	0.39		1	95			
Kocuria varians		3.34	0.16		1	10			
000422	**Low discrimination**								**000422**
					PYR				
Aerococcus viridans		72.80	0.51	ESL 90	95				
Kocuria varians		22.65	0.26		10				
Staphylococcus auricularis		2.93	0.07		40				
000423	**Very good identification**								**000423**
					NOV	PYR			
Aerococcus viridans		97.85	0.58	ESL 90	1	95			
Kocuria varians		1.14	0.04		1	10			
Staphylococcus cohnii subsp. cohnii		1.01	0.05		99	1			
000430	**Low discrimination**								**000430**
					OXI	YEP			
Macrococcus bovicus		61.89	0.51	XYL 1	99	99			
Staphylococcus cohnii subsp. cohnii		20.54	0.48		1	12			
Staphylococcus vitulinus		5.87	0.24		99	75			
000431	**Low discrimination**								**000431**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		77.89	0.49	XYL 1	1	99	1	12	
Aerococcus viridans		19.92	0.48		1	1	95	1	
Macrococcus bovicus		2.19	0.10		99	99	1	99	
000432	**Very good identification**								**000432**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		96.35	0.60	ESL 90	1	1	95	1	
Macrococcus bovicus		2.74	0.10		99	99	1	99	
Staphylococcus cohnii subsp. cohnii		0.91	0.07		1	99	1	12	
000433	**Very good identification**								**000433**
					NOV	PYR			
Aerococcus viridans		99.56	0.68	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		0.44	0.08		99	1			
000440	**Low discrimination**								**000440**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		84.79	0.97		1	99	12		

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus auricularis		10.26	0.75		1	1	1		
Macrococcus bovicus		2.42	0.59		99	99	99		
000441	**Very good identification**								**000441**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		98.15	0.98		1	99			
Staphylococcus auricularis		1.22	0.55		1	1			
Kocuria kristinae		0.43	0.39		99	1			
000442	**Low discrimination**								**000442**
					NOV	PYR			
Staphylococcus auricularis		44.53	0.55	NIT 90	1	40			
Staphylococcus cohnii subsp. cohnii		33.44	0.56	LAC 10	99	1			
Aerococcus viridans		19.15	0.63		1	95			
000443	**Low discrimination**								**000443**
					NOV	PYR			
Aerococcus viridans		49.76	0.71	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		40.33	0.57		99	1			
Staphylococcus auricularis		5.51	0.36		1	40			
000450	**Low discrimination**								**000450**
					OXI	YEP			
Macrococcus bovicus		65.90	1.00		99	99			
Staphylococcus cohnii subsp. cohnii		29.68	0.99		1	12			
Macrococcus equipersicus		4.28	0.77		99	99			
000451	**Very good identification**								**000451**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		97.51	1.00		1	99	1	12	
Macrococcus bovicus		2.02	0.59		99	99	1	99	
Aerococcus viridans		0.32	0.61		1	1	95	1	
000452	**Low discrimination**								**000452**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		45.51	0.59	LAC 1	99	99	1	99	
Aerococcus viridans		27.68	0.73		1	1	95	1	
Staphylococcus cohnii subsp. cohnii		20.50	0.58		1	99	1	12	
000453	**Low discrimination**								**000453**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		73.87	0.80	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		25.40	0.59		1	99	1	12	
Macrococcus bovicus		0.53	0.18		99	99	1	99	
000460	**Low discrimination**								**000460**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		78.51	0.56	XYL 1	99	1			
Staphylococcus auricularis		9.50	0.34		1	40			
Aerococcus viridans		5.00	0.43		1	95			
000461	**Low discrimination**								**000461**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		86.45	0.57	XYL 1	99	1			
Aerococcus viridans		11.85	0.51		1	95			
Staphylococcus auricularis		1.07	0.14		1	40			
000462	**Good identification**								**000462**
					NOV	PYR			
Aerococcus viridans		94.50	0.63	ESL 90	1	95			
Staphylococcus auricularis		2.22	0.14		1	40			
Staphylococcus cohnii subsp. cohnii		1.67	0.15		99	1			

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
000463	**Very good identification**								**000463**
					NOV	PYR			
Aerococcus viridans		99.19	0.71	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		0.81	0.16		99	1			
000470	**Low discrimination**								**000470**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		62.63	0.59	XYL 1	99	99	1	99	
Staphylococcus cohnii subsp. cohnii		28.21	0.58		1	99	1	12	
Aerococcus viridans		4.23	0.53		1	1	95	1	
000471	**Low discrimination**								**000471**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		74.42	0.59	XYL 1	1	99	1	12	
Aerococcus viridans		24.04	0.61		1	1	95	1	
Macrococcus bovicus		1.54	0.18		99	99	1	99	
000472	**Good identification**								**000472**
					OXI	NOV	YEP		
Aerococcus viridans		94.88	0.73	ESL 90	1	1	1		
Staphylococcus kloosii		2.31	0.40		1	99	99		
Macrococcus bovicus		1.58	0.18		99	99	99		
000473	**Very good identification**								**000473**
					NOV	YEP	PYR		
Aerococcus viridans		99.34	0.80	ESL 90	1	1	95		
Staphylococcus cohnii subsp. cohnii		0.35	0.18		99	12	1		
Staphylococcus arlettae		0.31	0.18		99	99	1		
000500	**Low discrimination**								**000500**
					OXI				
Macrococcus carouselicus		47.81	0.62	ESL 94	99				
Staphylococcus cohnii subsp. cohnii		28.55	0.67		1				
Dermacoccus nishinomiyaensis		6.55	0.44		93				
000501	**Good identification**								**000501**
					NOV				
Staphylococcus cohnii subsp. cohnii		93.80	0.68	GAL 10	99				
Staphylococcus auricularis		1.58	0.28		1				
Kocuria varians		1.46	0.29		1				
000502	**Low discrimination**								**000502**
					PYR				
Kocuria varians		37.74	0.39	TRE 7	10				
Aerococcus viridans		28.42	0.51		95				
Staphylococcus auricularis		12.58	0.28		40				
000503	**Low discrimination**								**000503**
					NOV	PYR			
Aerococcus viridans		83.51	0.58	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		9.50	0.27		99	1			
Kocuria varians		4.13	0.17		1	10			
000510	**Low discrimination**								**000510**
					OXI	YEP			
Macrococcus bovicus		82.23	0.85		99	99			
Staphylococcus cohnii subsp. cohnii		7.08	0.69		1	12			
Macrococcus carouselicus		5.95	0.58		99	67			
000511	**Low discrimination**								**000511**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		87.30	0.70	GAL 10	1	99	1	12	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Macrococcus bovicus		9.46	0.44		99	99	1	99	
Aerococcus viridans		2.03	0.48		1	1	95	1	
000512	**Low discrimination**								**000512**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		48.19	0.44	LAC 1	99	99	1	99	
Aerococcus viridans		39.89	0.60		1	1	95	1	
Staphylococcus cohnii subsp. cohnii		4.15	0.28		1	99	1	12	
000513	**Good identification**								**000513**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		94.92	0.68	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		4.58	0.29		1	99	1	12	
Macrococcus bovicus		0.50	0.03		99	99	1	99	
000520	**Low discrimination**								**000520**
					NOV	OXI			
Kocuria varians		64.78	0.41	TRE 7	1	13			
Macrococcus carouselicus		16.07	0.20		99	99			
Staphylococcus cohnii subsp. cohnii		9.59	0.26		99	1			
000521	**Low discrimination**								**000521**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		39.70	0.27	GAL 10	99	1			
Aerococcus viridans		38.78	0.39	XYL 1	1	95			
Kocuria varians		21.52	0.19		1	10			
000522	**Low discrimination**								**000522**
					PYR				
Aerococcus viridans		68.18	0.51	ESL 90	95				
Kocuria varians		31.82	0.30		10				
000523	**Very good identification**								**000523**
					PYR				
Aerococcus viridans		98.29	0.58	ESL 90	95				
Kocuria varians		1.71	0.08		10				
000530	**Low discrimination**								**000530**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		76.10	0.44	XYL 1	99	99	1	99	
Aerococcus viridans		7.00	0.41		1	1	95	1	
Staphylococcus cohnii subsp. cohnii		6.55	0.28		1	99	1	12	
000531	**Low discrimination**								**000531**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		67.50	0.48	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		29.32	0.29	LAC 90	1	99	1	12	
Macrococcus bovicus		3.18	0.03		99	99	1	99	
000532	**Very good identification**								**000532**
					OXI	NOV	YEP		
Aerococcus viridans		97.83	0.60	ESL 90	1	1	1		
Macrococcus bovicus		1.19	0.03		99	99	99		
Staphylococcus kloosii		0.97	0.20		1	99	99		
000533	**Very good identification**								**000533**
Aerococcus viridans		100.0	0.68	ESL 90					
000540	**Low discrimination**								**000540**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		76.94	0.78	GAL 10	1	99	12		
Staphylococcus auricularis		9.31	0.55		1	1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Macrococcus bovicus		8.47	0.51		99	99	99		
000541	**Very good identification**								**000541**
					OXI	NOV	PYR		
Staphylococcus cohnii subsp. cohnii		96.10	0.78	GAL 10	1	99	1		
Kocuria kristinae		1.27	0.30		99	1	93		
Aerococcus viridans		1.20	0.51		1	1	95		
000542	**Low discrimination**								**000542**
					NOV	PYR			
Aerococcus viridans		59.83	0.63	ESL 90	1	95			
Staphylococcus auricularis		15.46	0.36		1	40			
Staphylococcus cohnii subsp. cohnii		11.61	0.36		99	1			
000543	**Low discrimination**								**000543**
					OXI	NOV	PYR		
Aerococcus viridans		88.14	0.71	ESL 90	1	1	95		
Staphylococcus cohnii subsp. cohnii		7.94	0.37		1	99	1		
Kocuria kristinae		2.60	0.17		99	1	93		
000550	**Low discrimination**								**000550**
					OXI	YEP			
Macrococcus bovicus		84.30	0.92		99	99			
Staphylococcus cohnii subsp. cohnii		9.85	0.80		1	12			
Macrococcus equipersicus		5.47	0.70		99	99			
000551	**Low discrimination**								**000551**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		89.68	0.80	GAL 10	1	99	1	12	
Macrococcus bovicus		7.16	0.51		99	99	1	99	
Aerococcus viridans		2.63	0.61		1	1	95	1	
000552	**Low discrimination**								**000552**
					OXI	NOV	YEP		
Aerococcus viridans		49.46	0.73	ESL 90	1	1	1		
Macrococcus bovicus		34.85	0.51		99	99	99		
Staphylococcus kloosii		9.35	0.59		1	99	99		
000553	**Very good identification**								**000553**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		95.96	0.80	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		3.67	0.39		1	99	1	12	
Macrococcus bovicus		0.29	0.10		99	99	1	99	
000560	**Low discrimination**								**000560**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		43.31	0.36	GAL 10	99	1			
Aerococcus viridans		24.80	0.43	XYL 1	1	95			
Kocuria varians		20.27	0.27		1	10			
000561	**Low discrimination**								**000561**
					NOV	PYR			
Aerococcus viridans		54.32	0.51	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		44.03	0.37	LAC 90	99	1			
Kocuria varians		1.65	0.06		1	10			
000562	**Very good identification**								**000562**
					PYR				
Aerococcus viridans		97.50	0.63	ESL 90	95				
Kocuria varians		2.50	0.16		10				
000563	**Very good identification**								**000563**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Aerococcus viridans		100.0	0.71	ESL 90					
000570	**Low discrimination**								**000570**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		71.36	0.51	XYL 1	99	99	1	99	
Aerococcus viridans		11.25	0.53		1	1	95	1	
Staphylococcus cohnii subsp. cohnii		8.33	0.39		1	99	1	12	
000571	**Low discrimination**								**000571**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		72.92	0.61	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		25.08	0.39	LAC 90	1	99	1	12	
Macrococcus bovicus		2.00	0.10		99	99	1	99	
000572	**Good identification**								**000572**
					NOV	YEP			
Aerococcus viridans		92.11	0.73	ESL 90	1	1			
Staphylococcus kloosii		4.36	0.46		99	99			
Staphylococcus arlettae		2.88	0.31		99	99			
000573	**Very good identification**								**000573**
					NOV	YEP			
Aerococcus viridans		98.26	0.80	ESL 90	1	1			
Staphylococcus arlettae		1.74	0.34		99	99			
000600	**Low discrimination**								**000600**
					NOV	OXI			
Macrococcus carouselicus		77.91	0.75	ESL 94	99	99			
Staphylococcus auricularis		13.40	0.67		1	1			
Micrococcus sp.		3.29	0.38		1	89			
000601	**Low discrimination**								**000601**
					OXI	NOV			
Kocuria kristinae		80.67	0.61	VPT 90	99	1			
Staphylococcus auricularis		8.18	0.47	MLT 90	1	1			
Staphylococcus cohnii subsp. cohnii		5.31	0.47		1	99			
000602	**Low discrimination**								**000602**
					OXI	NOV			
Staphylococcus auricularis		52.27	0.47	NIT 90	1	1			
Macrococcus carouselicus		27.62	0.34	LAC 10	99	99			
Aerococcus viridans		14.21	0.51		1	1			
000603	**Low discrimination**								**000603**
					OXI				
Kocuria kristinae		76.64	0.48	VPT 90	99				
Aerococcus viridans		19.71	0.58	MLT 90	1				
Staphylococcus auricularis		3.45	0.27		1				
000610	**Low discrimination**								**000610**
Macrococcus bovicus		56.56	0.80						
Macrococcus carouselicus		33.25	0.71						
Staphylococcus vitulinus		7.94	0.56						
000611	**Low discrimination**								**000611**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		28.06	0.49	SUC 1	1	99	12		
Macrococcus bovicus		19.51	0.39		99	99	99		
Staphylococcus capitis subsp. capitis		16.08	0.35		1	1	1		
000612	**Low discrimination**								**000612**
					OXI	NOV			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		44.34	0.60	ESL 90	1	1		
Macrococcus bovicus		31.25	0.39		99	99		
Macrococcus carouselicus		18.37	0.30		99	99		
000613	**Good identification**							**000613**
					OXI	PYR		
Aerococcus viridans		93.76	0.68	ESL 90	1	95		
Kocuria kristinae		5.07	0.20		99	93		
Staphylococcus capitis subsp. capitis		0.48	0.00		1	1		
000620	**Low discrimination**							**000620**
					OXI	NOV		
Macrococcus carouselicus		40.46	0.34	ESL 94	99	99		
Staphylococcus muscae		23.06	0.18	XYL 1	1	1		
Staphylococcus vitulinus		16.70	0.24		99	99		
000622	**Acceptable identification**							**000622**
					PYR			
Aerococcus viridans		92.63	0.51	ESL 90	95			
Kocuria varians		3.93	0.08		10			
Staphylococcus auricularis		3.44	0.06		40			
000623	**Very good identification**							**000623**
					OXI			
Aerococcus viridans		96.22	0.58	ESL 90	1			
Kocuria kristinae		3.78	0.07		99			
000630	**Very good identification**							**000630**
Staphylococcus vitulinus		95.45	0.65	NIT 91				
Macrococcus bovicus		2.54	0.39					
Macrococcus carouselicus		1.49	0.30					
000632	**Low discrimination**							**000632**
					OXI	NOV	YEP	
Aerococcus viridans		77.13	0.60	ESL 90	1	1	1	
Staphylococcus vitulinus		20.62	0.24		99	99	75	
Staphylococcus arlettae		1.70	0.15		1	99	99	
000633	**Very good identification**							**000633**
					NOV	YEP		
Aerococcus viridans		98.76	0.68	ESL 90	1	1		
Staphylococcus arlettae		1.24	0.18		99	99		
000640	**Low discrimination**							**000640**
					NOV	PYR		
Staphylococcus petrasii subsp. pragensis		83.46	0.74	VPT 95	1	99		
Staphylococcus auricularis		12.93	0.74		1	40		
Staphylococcus cohnii subsp. cohnii		1.17	0.56		99	1		
000641	**Good identification**							**000641**
					OXI	NOV		
Kocuria kristinae		93.61	0.80	VPT 90	99	1		
Staphylococcus auricularis		2.46	0.55		1	1		
Staphylococcus cohnii subsp. cohnii		2.17	0.57		1	99		
000642	**Low discrimination**							**000642**
Staphylococcus auricularis		44.24	0.55	NIT 90				
Staphylococcus petrasii subsp. pragensis		25.96	0.33	LAC 10				
Aerococcus viridans		20.62	0.63					
000643	**Low discrimination**							**000643**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI				
Kocuria kristinae		88.69	0.68	VPT 90	99				
Aerococcus viridans		10.14	0.71		1				
Staphylococcus auricularis		1.04	0.35		1				
000650	**Good identification**								**000650**
					OXI	YEP			
Macrococcus bovicus		93.87	0.88		99	99			
Macrococcus equipersicus		2.41	0.56		99	99			
Staphylococcus cohnii subsp. cohnii		1.71	0.58		1	12			
000651	**Low discrimination**								**000651**
					OXI	NOV	PYR	YEP	
Kocuria kristinae		42.99	0.52	VPT 90	99	1	93	15	
Staphylococcus cohnii subsp. cohnii		30.41	0.59	MAN 4	1	99	1	12	
Macrococcus bovicus		15.58	0.47		99	99	1	99	
000652	**Low discrimination**								**000652**
					OXI	NOV	PYR		
Aerococcus viridans		60.47	0.73	ESL 90	1	1	95		
Macrococcus bovicus		24.86	0.47		99	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		9.74	0.29		1	99	7		
000653	**Low discrimination**								**000653**
					OXI	NOV	PYR		
Aerococcus viridans		88.50	0.80	ESL 90	1	1	95		
Kocuria kristinae		10.76	0.40		99	1	93		
Staphylococcus cohnii subsp. cohnii		0.31	0.18		1	99	1		
000660	**Low discrimination**								**000660**
Staphylococcus petrasii subsp. pragensis		78.11	0.33	VPT 95					
Staphylococcus auricularis		12.10	0.33	XYL 1					
Aerococcus viridans		6.89	0.43						
000661	**Low discrimination**								**000661**
					OXI				
Kocuria kristinae		72.51	0.39	VPT 90	99				
Aerococcus viridans		22.79	0.51	XYL 1	1				
Staphylococcus auricularis		1.91	0.13		1				
000662	**Very good identification**								**000662**
					NOV	YEP			
Aerococcus viridans		95.15	0.63	ESL 90	1	1			
Staphylococcus auricularis		2.06	0.13		1	1			
Staphylococcus arlettae		1.57	0.15		99	99			
000663	**Good identification**								**000663**
					OXI	NOV	YEP		
Aerococcus viridans		91.09	0.71	ESL 90	1	1	1		
Kocuria kristinae		8.05	0.27		99	1	15		
Staphylococcus arlettae		0.86	0.18		1	99	99		
000670	**Low discrimination**								**000670**
					OXI	NOV			
Macrococcus bovicus		66.40	0.47	XYL 1	99	99			
Aerococcus viridans		17.95	0.53		1	1			
Staphylococcus vitulinus		10.80	0.24		99	99			
000671	**Low discrimination**								**000671**
					OXI	NOV	PYR		
Aerococcus viridans		87.54	0.61	ESL 90	1	1	95		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		3.87	0.11	LAC 90	99	1	93	
Staphylococcus cohnii subsp. cohnii		2.74	0.18		1	99	1	
000672	**Low discrimination**							**000672**
					OXI	NOV	YEP	
Aerococcus viridans		64.43	0.73	ESL 90	1	1	1	
Staphylococcus arlettae		35.15	0.57		1	99	99	
Macrococcus bovicus		0.27	0.05		99	99	99	
000673	**Low discrimination**							**000673**
					OXI	NOV	YEP	
Aerococcus viridans		76.27	0.80	ESL 90	1	1	1	
Staphylococcus arlettae		23.64	0.59		1	99	99	
Kocuria kristinae		0.09	-.02		99	1	15	
000700	**Good identification**							**000700**
					OXI	NOV		
Macrococcus carouselicus		92.84	0.68	ESL 94	99	99		
Staphylococcus auricularis		4.14	0.47		1	1		
Macrococcus bovicus		1.02	0.31		99	99		
000701	**Low discrimination**							**000701**
					OXI	NOV	PYR	
Kocuria kristinae		86.20	0.51	VPT 90	99	1	93	
Macrococcus carouselicus		5.94	0.27	MLT 90	99	99	1	
Staphylococcus auricularis		2.91	0.27		1	1	40	
000702	**Low discrimination**							**000702**
					OXI	NOV		
Aerococcus viridans		40.76	0.51	ESL 90	1	1		
Macrococcus carouselicus		33.96	0.27		99	99		
Staphylococcus auricularis		16.66	0.27		1	1		
000703	**Low discrimination**							**000703**
					OXI			
Kocuria kristinae		55.74	0.39	VPT 90	99			
Aerococcus viridans		42.99	0.58	MLT 90	1			
Staphylococcus auricularis		0.84	0.08		1			
000710	**Low discrimination**							**000710**
Macrococcus bovicus		61.69	0.73					
Macrococcus carouselicus		36.27	0.64					
Macrococcus equipersicus		1.37	0.40					
000711	**Low discrimination**							**000711**
					OXI	NOV		
Macrococcus bovicus		28.98	0.31	MNS 1	99	99		
Aerococcus viridans		24.87	0.48		1	1		
Macrococcus carouselicus		17.04	0.23		99	99		
000712	**Low discrimination**							**000712**
					OXI	NOV		
Aerococcus viridans		67.28	0.60	ESL 90	1	1		
Macrococcus bovicus		20.32	0.31		99	99		
Macrococcus carouselicus		11.95	0.23		99	99		
000713	**Very good identification**							**000713**
					OXI	NOV	PYR	
Aerococcus viridans		97.38	0.68	ESL 90	1	1	95	
Kocuria kristinae		1.75	0.10		99	1	93	
Staphylococcus lentus		0.86	0.12		99	99	6	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
000720	**Low discrimination**							**000720**
					NOV	PYR	OXI	
Macrococcus carouselicus		69.02	0.27	ESL 94	99	1	99	
Kocuria varians		18.69	0.23	XYL 1	1	10	13	
Aerococcus viridans		9.21	0.31		1	95	1	
000721	**Low discrimination**							**000721**
					OXI			
Aerococcus viridans		67.96	0.39	ESL 90	1			
Kocuria kristinae		32.04	0.10	LAC 90	99			
000722	**Acceptable identification**							**000722**
					PYR			
Aerococcus viridans		94.02	0.51	ESL 90	95			
Kocuria varians		5.98	0.12		10			
000723	**Very good identification**							**000723**
					OXI			
Aerococcus viridans		98.71	0.58	ESL 90	1			
Kocuria kristinae		1.29	-.03		99			
000730	**Low discrimination**							**000730**
Macrococcus bovicus		34.92	0.31	XYL 1				
Staphylococcus vitulinus		30.92	0.24					
Macrococcus carouselicus		20.53	0.23					
000731	**Good identification**							**000731**
					OXI	NOV		
Aerococcus viridans		95.52	0.48	ESL 90	1	1		
Staphylococcus sciuri		4.48	0.05	LAC 90	99	99		
000732	**Low discrimination**							**000732**
					NOV	YEP		
Aerococcus viridans		88.90	0.60	ESL 90	1	1		
Staphylococcus arlettae		11.11	0.31		99	99		
000733	**Good identification**							**000733**
					OXI	NOV	YEP	
Aerococcus viridans		92.61	0.68	ESL 90	1	1	1	
Staphylococcus arlettae		6.57	0.34		1	99	99	
Staphylococcus lentus		0.82	0.12		99	99	1	
000740	**Low discrimination**							**000740**
					OXI	NOV		
Staphylococcus auricularis		42.22	0.55	NIT 90	1	1		
Staphylococcus petrasii subsp. pragensis		24.78	0.33	GAL 10	1	1		
Macrococcus bovicus		10.40	0.39		99	99		
000741	**Very good identification**							**000741**
					OXI			
Kocuria kristinae		97.17	0.71	VPT 90	99			
Aerococcus viridans		0.93	0.51		1			
Staphylococcus auricularis		0.85	0.35		1			
000742	**Low discrimination**							**000742**
					OXI	NOV		
Aerococcus viridans		66.74	0.63	ESL 90	1	1		
Staphylococcus auricularis		15.91	0.35		1	1		
Macrococcus caseolyticus		4.98	0.14		99	99		
000743	**Low discrimination**							**000743**
					OXI			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		74.26	0.58	VPT 90	99			
Aerococcus viridans		25.46	0.71		1			
Staphylococcus auricularis		0.29	0.15		1			
000750	**Very good identification**							**000750**
					OXI	NOV		
Macrococcus bovicus		95.78	0.80		99	99		
Macrococcus equipersicus		2.46	0.49		99	99		
Aerococcus viridans		0.61	0.53		1	1		
000751	**Low discrimination**							**000751**
					OXI	NOV	PYR	
Kocuria kristinae		40.76	0.42	VPT 90	99	1	93	
Aerococcus viridans		27.95	0.61	MAN 4	1	1	95	
Macrococcus bovicus		18.99	0.39		99	99	1	
000752	**Low discrimination**							**000752**
					OXI	NOV	YEP	
Aerococcus viridans		79.29	0.73	ESL 90	1	1	1	
Macrococcus bovicus		13.97	0.39		99	99	99	
Staphylococcus arlettae		2.48	0.31		1	99	99	
000753	**Good identification**							**000753**
					OXI	NOV	YEP	
Aerococcus viridans		93.94	0.80	ESL 90	1	1	1	
Kocuria kristinae		3.81	0.30		99	1	15	
Staphylococcus arlettae		1.67	0.34		1	99	99	
000760	**Low discrimination**							**000760**
Aerococcus viridans		68.07	0.43	ESL 90				
Staphylococcus auricularis		13.28	0.13	LAC 90				
Staphylococcus petrasii subsp. pragensis		7.79	-.09					
000761	**Low discrimination**							**000761**
					OXI			
Kocuria kristinae		51.47	0.30	VPT 90	99			
Aerococcus viridans		48.53	0.51	XYL 1	1			
000762	**Good identification**							**000762**
					NOV	YEP		
Aerococcus viridans		91.43	0.63	ESL 90	1	1		
Staphylococcus arlettae		8.57	0.31		99	99		
000763	**Good identification**							**000763**
					OXI	NOV	YEP	
Aerococcus viridans		92.36	0.71	ESL 90	1	1	1	
Staphylococcus arlettae		4.92	0.34		1	99	99	
Kocuria kristinae		2.72	0.17		99	1	15	
000770	**Low discrimination**							**000770**
					OXI	NOV	YEP	
Macrococcus bovicus		53.82	0.39	XYL 1	99	99	99	
Aerococcus viridans		33.94	0.53		1	1	1	
Staphylococcus arlettae		9.54	0.31		1	99	99	
000771	**Low discrimination**							**000771**
					OXI	NOV	YEP	
Aerococcus viridans		80.86	0.61	ESL 90	1	1	1	
Staphylococcus arlettae		12.91	0.34	LAC 90	1	99	99	
Staphylococcus sciuri		3.56	0.17		99	99	57	
000772	**Low discrimination**							**000772**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus arlettae		75.44	0.72	GLR 92	99	99		
Aerococcus viridans		24.40	0.73		1	1		
Staphylococcus kloosii		0.11	0.26		99	99		
000773	**Low discrimination**							**000773**
					OXI	NOV	YEP	
Staphylococcus arlettae		63.57	0.75	GLR 92	1	99	99	
Aerococcus viridans		36.19	0.80		1	1	1	
Staphylococcus lentus		0.16	0.18		99	99	1	
001000	**Low discrimination**							**001000**
					NOV			
Macrococcus carouselicus		43.79	0.69	TRE 94	99			
Micrococcus sp.		38.01	0.59		1			
Dermacoccus nishinomiyaensis		13.50	0.59		80			
001002	**Low discrimination**							**001002**
					NOV	OXI		
Macrococcus carouselicus		34.60	0.28	TRE 94	99	99		
Micrococcus sp.		30.04	0.18	LAC 1	1	89		
Aerococcus viridans		17.12	0.44		1	1		
001003	**Good identification**							**001003**
Aerococcus viridans		100.0	0.52	TRE 95				
001010	**Low discrimination**							**001010**
Macrococcus equipersicus		44.21	0.75	URE 86				
Macrococcus bovicus		27.51	0.69					
Macrococcus carouselicus		26.88	0.65					
001011	**Low discrimination**							**001011**
Macrococcus equipersicus		32.16	0.34	URE 86				
Macrococcus bovicus		20.01	0.28	MNS 1				
Macrococcus carouselicus		19.55	0.24					
001012	**Low discrimination**							**001012**
					OXI	NOV		
Aerococcus viridans		38.76	0.54	TRE 95	1	1		
Macrococcus equipersicus		27.46	0.34		99	99		
Macrococcus bovicus		17.08	0.28		99	99		
001013	**Very good identification**							**001013**
Aerococcus viridans		100.0	0.61	TRE 95				
001020	**Low discrimination**							**001020**
					NOV			
Kocuria rosea		40.27	0.32	ESL 1	1			
Macrococcus carouselicus		24.07	0.28	NIT 90	99			
Micrococcus sp.		20.89	0.18		1			
001021	**Low discrimination**							**001021**
					PYR			
Aerococcus viridans		59.29	0.32	TRE 95	95			
Kocuria rosea		40.71	0.05	LAC 90	1			
001022	**Acceptable identification**							**001022**
					PYR			
Aerococcus viridans		91.90	0.44	TRE 95	95			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		8.10	0.08		10		
001023	**Good identification**						**001023**
Aerococcus viridans		100.0	0.52	TRE 95			
001030	**Low discrimination**						**001030**
Staphylococcus vitulinus		59.90	0.38	NIT 91			
Macrococcus equipersicus		15.64	0.34	SUC 99			
Macrococcus bovicus		9.73	0.28				
001031	**Acceptable identification**						**001031**
					OXI	NOV	
Aerococcus viridans		90.44	0.42	TRE 95	1	1	
Staphylococcus vitulinus		9.56	-.03	LAC 90	99	99	
001032	**Very good identification**						**001032**
					OXI	NOV	
Aerococcus viridans		97.33	0.54	TRE 95	1	1	
Staphylococcus vitulinus		2.67	-.03		99	99	
001033	**Very good identification**						**001033**
Aerococcus viridans		100.0	0.61	TRE 95			
001040	**Low discrimination**						**001040**
					NOV		
Macrococcus equipersicus		60.16	0.57	URE 86	99		
Micrococcus sp.		26.27	0.36	MAN 95	1		
Macrococcus bovicus		6.20	0.36		99		
001041	**Low discrimination**						**001041**
					NOV	OXI	
Aerococcus viridans		55.77	0.44	TRE 95	1	1	
Macrococcus equipersicus		16.27	0.16	LAC 90	99	99	
Micrococcus sp.		14.35	0.01		1	89	
001042	**Low discrimination**						**001042**
					NOV	OXI	
Aerococcus viridans		89.09	0.56	TRE 95	1	1	
Macrococcus equipersicus		6.74	0.16		99	99	
Micrococcus sp.		2.94	-.05		1	89	
001043	**Very good identification**						**001043**
Aerococcus viridans		100.0	0.64	TRE 95			
001050	**Low discrimination**						**001050**
					OXI	NOV	
Macrococcus equipersicus		64.85	0.84	URE 86	99	99	
Macrococcus bovicus		34.82	0.77		99	99	
Aerococcus viridans		0.15	0.47		1	1	
001051	**Low discrimination**						**001051**
					OXI	NOV	
Macrococcus equipersicus		47.31	0.43	URE 86	99	99	
Aerococcus viridans		25.61	0.54	MNS 1	1	1	
Macrococcus bovicus		25.40	0.36		99	99	
001052	**Low discrimination**						**001052**
					OXI	NOV	
Aerococcus viridans		57.60	0.66	TRE 95	1	1	
Macrococcus equipersicus		27.59	0.43		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Micrococcus bovicus		14.81	0.36		99	99		
001053	**Very good identification**							**001053**
					OXI	NOV		
Aerococcus viridans		99.79	0.74	TRE 95	1	1		
Micrococcus equipersicus		0.21	0.02		99	99		
001060	**Low discrimination**							**001060**
					NOV	OXI		
Aerococcus viridans		46.90	0.37	TRE 95	1	1		
Micrococcus equipersicus		31.92	0.16	LAC 90	99	99		
Micrococcus sp.		13.94	-.05		1	89		
001061	**Good identification**							**001061**
Aerococcus viridans		100.0	0.44	TRE 95				
001062	**Very good identification**							**001062**
Aerococcus viridans		100.0	0.56	TRE 95				
001063	**Very good identification**							**001063**
Aerococcus viridans		100.0	0.64	TRE 95				
001070	**Low discrimination**							**001070**
					OXI	NOV		
Micrococcus equipersicus		56.08	0.43	URE 86	99	99		
Micrococcus bovicus		30.11	0.36	XYL 1	99	99		
Aerococcus viridans		13.01	0.47		1	1		
001071	**Very good identification**							**001071**
					OXI	NOV		
Aerococcus viridans		98.17	0.54	TRE 95	1	1		
Micrococcus equipersicus		1.83	0.02	LAC 90	99	99		
001072	**Very good identification**							**001072**
					OXI	NOV		
Aerococcus viridans		99.52	0.66	TRE 95	1	1		
Micrococcus equipersicus		0.48	0.02		99	99		
001073	**Very good identification**							**001073**
Aerococcus viridans		100.0	0.74	TRE 95				
001100	**Low discrimination**							**001100**
					YEP			
Micrococcus carouselicus		80.83	0.62	TRE 94	67			
Dermacoccus nishinomiyaensis		11.08	0.44		1			
Micrococcus equipersicus		4.47	0.41		99			
001102	**Low discrimination**							**001102**
					NOV	PYR	OXI	
Aerococcus viridans		41.51	0.44	TRE 95	1	95	1	
Micrococcus carouselicus		35.96	0.20		99	1	99	
Kocuria varians		15.62	0.21		1	10	13	
001103	**Good identification**							**001103**
Aerococcus viridans		100.0	0.52	TRE 95				
001110	**Low discrimination**							**001110**
Micrococcus equipersicus		44.74	0.67	URE 86				
Micrococcus bovicus		27.83	0.62					

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Micrococcus carouselicus		27.20	0.58					
001112	**Low discrimination**							**001112**
					OXI	NOV		
Aerococcus viridans		59.62	0.54	TRE 95	1	1		
Micrococcus equipersicus		18.11	0.26		99	99		
Micrococcus bovicus		11.26	0.21		99	99		
001113	**Very good identification**							**001113**
Aerococcus viridans		100.0	0.61	TRE 95				
001121	**Good identification**							**001121**
Aerococcus viridans		100.0	0.32	TRE 95				
001122	**Low discrimination**							**001122**
					PYR			
Aerococcus viridans		88.32	0.44	TRE 95	95			
Kocuria varians		11.68	0.12		10			
001123	**Good identification**							**001123**
Aerococcus viridans		100.0	0.52	TRE 95				
001131	**Good identification**							**001131**
Aerococcus viridans		100.0	0.42	TRE 95				
001132	**Very good identification**							**001132**
Aerococcus viridans		100.0	0.54	TRE 95				
001133	**Very good identification**							**001133**
Aerococcus viridans		100.0	0.61	TRE 95				
001140	**Low discrimination**							**001140**
Micrococcus equipersicus		80.16	0.50	URE 86				
Micrococcus bovicus		8.26	0.28	MAN 95				
Micrococcus carouselicus		5.41	0.20					
001141	**Low discrimination**							**001141**
					OXI	NOV		
Aerococcus viridans		88.89	0.44	TRE 95	1	1		
Micrococcus equipersicus		11.11	0.09	LAC 90	99	99		
001142	**Good identification**							**001142**
					NOV	PYR	OXI	
Aerococcus viridans		94.96	0.56	TRE 95	1	95	1	
Micrococcus equipersicus		3.08	0.09		99	1	99	
Kocuria varians		1.96	0.08		1	10	13	
001143	**Very good identification**							**001143**
Aerococcus viridans		100.0	0.64	TRE 95				
001150	**Low discrimination**							**001150**
					OXI	NOV		
Micrococcus equipersicus		64.73	0.76	URE 86	99	99		
Micrococcus bovicus		34.75	0.69		99	99		
Aerococcus viridans		0.35	0.47		1	1		
001151	**Low discrimination**							**001151**
					OXI	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		45.11	0.54	TRE 95	1	1		
Macrococcus equipersicus		35.72	0.35	LAC 90	99	99		
Macrococcus bovicus		19.18	0.28		99	99		
001152	**Low discrimination**							**001152**
					OXI	NOV		
Aerococcus viridans		75.79	0.66	TRE 95	1	1		
Macrococcus equipersicus		15.56	0.35		99	99		
Macrococcus bovicus		8.35	0.28		99	99		
001153	**Very good identification**							**001153**
Aerococcus viridans		100.0	0.74	TRE 95				
001160	**Low discrimination**							**001160**
					NOV	PYR	OXI	
Aerococcus viridans		65.65	0.37	TRE 95	1	95	1	
Macrococcus equipersicus		19.15	0.09	LAC 90	99	1	99	
Kocuria varians		15.20	0.09		1	10	13	
001161	**Good identification**							**001161**
Aerococcus viridans		100.0	0.44	TRE 95				
001162	**Very good identification**							**001162**
Aerococcus viridans		100.0	0.56	TRE 95				
001163	**Very good identification**							**001163**
Aerococcus viridans		100.0	0.64	TRE 95				
001170	**Low discrimination**							**001170**
					OXI	NOV		
Macrococcus equipersicus		48.12	0.35	URE 86	99	99		
Aerococcus viridans		26.05	0.47	XYL 1	1	1		
Macrococcus bovicus		25.83	0.28		99	99		
001171	**Very good identification**							**001171**
Aerococcus viridans		100.0	0.54	TRE 95				
001172	**Very good identification**							**001172**
Aerococcus viridans		100.0	0.66	TRE 95				
001173	**Very good identification**							**001173**
Aerococcus viridans		100.0	0.74	TRE 95				
001200	**Good identification**							**001200**
					NOV			
Macrococcus carouselicus		93.83	0.75	TRE 94	99			
Micrococcus sp.		3.97	0.38		1			
Dermacoccus nishinomiyaensis		0.91	0.34		80			
001201	**Low discrimination**							**001201**
					NOV	PYR		
Macrococcus carouselicus		69.50	0.34	TRE 94	99	1		
Kocuria kristinae		17.08	0.22	MNS 1	1	93		
Micrococcus sp.		5.94	0.03		1	63		
001202	**Low discrimination**							**001202**
					NOV	OXI		
Macrococcus carouselicus		77.76	0.34	TRE 94	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		18.95	0.44	LAC 1	1	1	
Micrococcus sp.		3.29	-.03		1	89	
001203	**Acceptable identification**						**001203**
					OXI		
Aerococcus viridans		90.25	0.52	TRE 95	1		
Kocuria kristinae		9.75	0.10		99		
001210	**Low discrimination**						**001210**
Staphylococcus vitulinus		48.71	0.70	NIT 91			
Macrococcus carouselicus		42.45	0.71				
Macrococcus bovicus		5.35	0.57				
001211	**Low discrimination**						**001211**
					OXI	NOV	
Staphylococcus vitulinus		33.54	0.29	NIT 91	99	99	
Macrococcus carouselicus		29.23	0.30	MNS 1	99	99	
Staphylococcus capitis subsp. capitis		20.28	0.29		1	1	
001212	**Low discrimination**						**001212**
					OXI	NOV	
Staphylococcus vitulinus		32.81	0.29	NIT 91	99	99	
Aerococcus viridans		32.70	0.54	LAC 1	1	1	
Macrococcus carouselicus		28.60	0.30		99	99	
001213	**Very good identification**						**001213**
Aerococcus viridans		100.0	0.61	TRE 95			
001220	**Low discrimination**						**001220**
					NOV		
Staphylococcus vitulinus		62.87	0.38	NIT 91	99		
Macrococcus carouselicus		31.70	0.34	MAN 99	99		
Kocuria rosea		2.90	0.12		1		
001221	**Low discrimination**						**001221**
					OXI	NOV	
Aerococcus viridans		75.92	0.32	TRE 95	1	1	
Staphylococcus vitulinus		24.08	-.03	LAC 90	99	99	
001222	**Acceptable identification**						**001222**
					OXI	NOV	
Aerococcus viridans		92.40	0.44	TRE 95	1	1	
Staphylococcus vitulinus		7.60	-.03		99	99	
001223	**Good identification**						**001223**
Aerococcus viridans		100.0	0.52	TRE 95			
001230	**Very good identification**						**001230**
					OXI	NOV	
Staphylococcus vitulinus		99.56	0.79	NIT 91	99	99	
Macrococcus carouselicus		0.32	0.30		99	99	
Aerococcus viridans		0.04	0.34		1	1	
001231	**Acceptable identification**						**001231**
					OXI	NOV	
Staphylococcus vitulinus		90.05	0.38	NIT 91	99	99	
Aerococcus viridans		8.60	0.42	MNS 1	1	1	
Staphylococcus sciuri		1.34	0.09		99	99	
001232	**Low discrimination**						**001232**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus vitulinus		73.07	0.38	NIT 91	99	99		
Aerococcus viridans		26.93	0.54	LAC 1	1	1		
001233	**Very good identification**							**001233**
					OXI	NOV		
Aerococcus viridans		98.84	0.61	TRE 95	1	1		
Staphylococcus vitulinus		1.16	-.03		99	99		
001240	**Low discrimination**							**001240**
					NOV			
Macrococcus carouselicus		37.02	0.34	TRE 94	99			
Macrococcus equipersicus		26.71	0.37	MLT 1	99			
Micrococcus sp.		11.66	0.15		1			
001241	**Low discrimination**							**001241**
					OXI			
Kocuria kristinae		89.75	0.42	VPT 90	99			
Aerococcus viridans		10.26	0.44	TRE 95	1			
001242	**Very good identification**							**001242**
Aerococcus viridans		100.0	0.56	TRE 95				
001243	**Low discrimination**							**001243**
					OXI			
Aerococcus viridans		80.45	0.64	TRE 95	1			
Kocuria kristinae		19.56	0.29		99			
001250	**Low discrimination**							**001250**
Macrococcus bovicus		54.67	0.64	ESL 10				
Macrococcus equipersicus		40.29	0.63					
Staphylococcus vitulinus		2.16	0.29					
001251	**Low discrimination**							**001251**
					OXI	NOV		
Aerococcus viridans		57.25	0.54	TRE 95	1	1		
Macrococcus bovicus		14.20	0.23	LAC 90	99	99		
Staphylococcus sciuri		11.14	0.23		99	99		
001252	**Low discrimination**							**001252**
					OXI	NOV		
Aerococcus viridans		89.96	0.66	TRE 95	1	1		
Macrococcus bovicus		5.78	0.23		99	99		
Macrococcus equipersicus		4.26	0.22		99	99		
001253	**Very good identification**							**001253**
					OXI	NOV	PYR	
Aerococcus viridans		99.32	0.74	TRE 95	1	1	95	
Staphylococcus sciuri		0.35	0.07		99	99	1	
Kocuria kristinae		0.34	0.01		99	1	93	
001260	**Low discrimination**							**001260**
					OXI	NOV		
Aerococcus viridans		84.39	0.37	TRE 95	1	1		
Staphylococcus vitulinus		15.61	-.03	LAC 90	99	99		
001261	**Acceptable identification**							**001261**
					OXI			
Aerococcus viridans		91.88	0.44	TRE 95	1			
Kocuria kristinae		8.12	0.01	LAC 90	99			
001262	**Very good identification**							**001262**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.56	TRE 95				
001263	**Very good identification**							**001263**
Aerococcus viridans		100.0	0.64	TRE 95				
001270	**Low discrimination**							**001270**
					OXI	NOV		
Staphylococcus vitulinus		74.89	0.38	NIT 91	99	99		
Aerococcus viridans		12.27	0.47	MLT 1	1	1		
Macrococcus bovicus		7.10	0.23		99	99		
001271	**Low discrimination**							**001271**
					OXI	NOV		
Aerococcus viridans		85.24	0.54	TRE 95	1	1		
Staphylococcus sciuri		12.51	0.21	LAC 90	99	99		
Staphylococcus vitulinus		2.25	-.03		99	99		
001272	**Very good identification**							**001272**
					OXI	NOV	YEP	
Aerococcus viridans		98.19	0.66	TRE 95	1	1	1	
Staphylococcus arlettae		1.14	0.15		1	99	99	
Staphylococcus vitulinus		0.67	-.03		99	99	75	
001273	**Very good identification**							**001273**
					OXI	NOV	YEP	
Aerococcus viridans		99.08	0.74	TRE 95	1	1	1	
Staphylococcus arlettae		0.66	0.18		1	99	99	
Staphylococcus sciuri		0.26	0.05		99	99	57	
001300	**Very good identification**							**001300**
					YEP			
Macrococcus carouselicus		98.96	0.68	TRE 94	67			
Dermacoccus nishinomiyaensis		0.43	0.20		1			
Macrococcus equipersicus		0.27	0.20		99			
001301	**Low discrimination**							**001301**
					OXI	NOV	PYR	
Macrococcus carouselicus		74.71	0.27	TRE 94	99	99	1	
Kocuria kristinae		14.28	0.12	MNS 1	99	1	93	
Aerococcus viridans		11.01	0.32		1	1	95	
001302	**Low discrimination**							**001302**
					NOV	PYR	OXI	
Macrococcus carouselicus		62.59	0.27	TRE 94	99	1	99	
Aerococcus viridans		35.59	0.44	LAC 1	1	95	1	
Kocuria varians		1.83	0.03		1	10	13	
001303	**Good identification**							**001303**
					OXI			
Aerococcus viridans		96.52	0.52	TRE 95	1			
Kocuria kristinae		3.48	0.00		99			
001310	**Low discrimination**							**001310**
Macrococcus carouselicus		80.90	0.64	TRE 94				
Macrococcus bovicus		10.19	0.49					
Macrococcus equipersicus		6.48	0.47					
001312	**Low discrimination**							**001312**
					OXI	NOV		
Aerococcus viridans		68.87	0.54	TRE 95	1	1		
Macrococcus carouselicus		25.81	0.23		99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus bovicus		3.25	0.08		99	99	
001313	**Very good identification**						**001313**
					OXI	NOV	
Aerococcus viridans		97.47	0.61	TRE 95	1	1	
Staphylococcus lentus		1.82	0.12		99	99	
Staphylococcus sciuri		0.71	0.01		99	99	
001320	**Low discrimination**						**001320**
					OXI	NOV	
Macrococcus carouselicus		87.24	0.27	TRE 94	99	99	
Aerococcus viridans		5.51	0.24	XYL 1	1	1	
Staphylococcus vitulinus		4.08	-.03		99	99	
001321	**Good identification**						**001321**
Aerococcus viridans		100.0	0.32	TRE 95			
001322	**Good identification**						**001322**
Aerococcus viridans		100.0	0.44	TRE 95			
001323	**Good identification**						**001323**
Aerococcus viridans		100.0	0.52	TRE 95			
001330	**Low discrimination**						**001330**
					OXI	NOV	
Staphylococcus vitulinus		82.81	0.38	NIT 91	99	99	
Macrococcus carouselicus		11.44	0.23	GAL 1	99	99	
Aerococcus viridans		3.39	0.34		1	1	
001331	**Low discrimination**						**001331**
					OXI	NOV	
Aerococcus viridans		70.99	0.42	TRE 95	1	1	
Staphylococcus sciuri		21.50	0.15	LAC 90	99	99	
Staphylococcus vitulinus		7.51	-.03		99	99	
001332	**Very good identification**						**001332**
					OXI	NOV	
Aerococcus viridans		97.33	0.54	TRE 95	1	1	
Staphylococcus vitulinus		2.67	-.03		99	99	
001333	**Very good identification**						**001333**
					OXI	NOV	
Aerococcus viridans		98.17	0.61	TRE 95	1	1	
Staphylococcus lentus		1.84	0.12		99	99	
001340	**Low discrimination**						**001340**
					OXI	NOV	
Macrococcus carouselicus		45.24	0.27	TRE 94	99	99	
Macrococcus equipersicus		32.65	0.29	MLT 1	99	99	
Aerococcus viridans		11.43	0.37		1	1	
001341	**Low discrimination**						**001341**
					OXI		
Kocuria kristinae		73.54	0.32	VPT 90	99		
Aerococcus viridans		25.21	0.44	TRE 95	1		
Rothia mucilaginosa		1.24	-.09		1		
001342	**Very good identification**						**001342**
					NOV	PYR	
Aerococcus viridans		98.88	0.56	TRE 95	1	95	
Staphylococcus hominis subsp. novobiosepticus		1.12	-.08		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
001343	**Good identification**							**001343**
					OXI			
Aerococcus viridans		92.51	0.64	TRE 95	1			
Kocuria kristinae		7.50	0.19		99			
001350	**Low discrimination**							**001350**
					OXI	NOV		
Macrococcus bovicus		55.06	0.57	ESL 10	99	99		
Macrococcus equipersicus		40.57	0.56		99	99		
Aerococcus viridans		2.24	0.47		1	1		
001351	**Low discrimination**							**001351**
					OXI	NOV		
Aerococcus viridans		61.47	0.54	TRE 95	1	1		
Staphylococcus sciuri		23.21	0.29	LAC 90	99	99		
Macrococcus bovicus		6.53	0.16		99	99		
001352	**Very good identification**							**001352**
					OXI	NOV		
Aerococcus viridans		95.43	0.66	TRE 95	1	1		
Macrococcus bovicus		2.63	0.16		99	99		
Macrococcus equipersicus		1.94	0.14		99	99		
001353	**Very good identification**							**001353**
					OXI	NOV		
Aerococcus viridans		98.39	0.74	TRE 95	1	1		
Staphylococcus lentus		0.93	0.18		99	99		
Staphylococcus sciuri		0.67	0.13		99	99		
001360	**Good identification**							**001360**
Aerococcus viridans		100.0	0.37	TRE 95				
001361	**Good identification**							**001361**
Aerococcus viridans		100.0	0.44	TRE 95				
001362	**Very good identification**							**001362**
Aerococcus viridans		100.0	0.56	TRE 95				
001363	**Very good identification**							**001363**
Aerococcus viridans		100.0	0.64	TRE 95				
001370	**Low discrimination**							**001370**
					OXI	NOV		
Aerococcus viridans		63.51	0.47	TRE 95	1	1		
Macrococcus bovicus		15.75	0.16	LAC 90	99	99		
Macrococcus equipersicus		11.60	0.14		99	99		
001371	**Low discrimination**							**001371**
					OXI	NOV		
Aerococcus viridans		76.40	0.54	TRE 95	1	1		
Staphylococcus sciuri		21.76	0.27	LAC 90	99	99		
Staphylococcus lentus		1.84	0.07		99	99		
001372	**Good identification**							**001372**
					NOV	YEP		
Aerococcus viridans		93.82	0.66	TRE 95	1	1		
Staphylococcus arlettae		6.19	0.31		99	99		
001373	**Very good identification**							**001373**
					OXI	NOV	YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		95.05	0.74	TRE 95	1	1	1	
Staphylococcus arlettae		3.56	0.34		1	99	99	
Staphylococcus lentus		0.90	0.18		99	99	1	
001400	**Excellent identification**							**001400**
					OXI	YEP		
Macrococcus carouselicus		99.26	0.94		99	67		
Macrococcus equipersicus		0.35	0.49		99	99		
Staphylococcus cohnii subsp. cohnii		0.15	0.46		1	12		
001401	**Low discrimination**							**001401**
					OXI	NOV	PYR	
Macrococcus carouselicus		75.35	0.53	MNS 1	99	99	1	
Staphylococcus cohnii subsp. cohnii		12.00	0.47		1	99	1	
Aerococcus viridans		11.72	0.58		1	1	95	
001402	**Low discrimination**							**001402**
					OXI	NOV		
Macrococcus carouselicus		62.20	0.53	LAC 1	99	99		
Aerococcus viridans		37.32	0.70		1	1		
Macrococcus equipersicus		0.22	0.08		99	99		
001403	**Very good identification**							**001403**
					OXI	NOV	PYR	
Aerococcus viridans		99.06	0.78		1	1	95	
Macrococcus carouselicus		0.72	0.12		99	99	1	
Staphylococcus cohnii subsp. cohnii		0.11	0.05		1	99	1	
001410	**Low discrimination**							**001410**
Macrococcus carouselicus		82.65	0.90					
Macrococcus equipersicus		8.68	0.75					
Macrococcus bovicus		8.10	0.73					
001411	**Low discrimination**							**001411**
					OXI	NOV	PYR	
Macrococcus carouselicus		44.40	0.49	MNS 1	99	99	1	
Aerococcus viridans		32.40	0.68		1	1	95	
Staphylococcus cohnii subsp. cohnii		14.08	0.49		1	99	1	
001412	**Low discrimination**							**001412**
					OXI	NOV		
Aerococcus viridans		69.90	0.80		1	1		
Macrococcus carouselicus		24.83	0.49		99	99		
Macrococcus equipersicus		2.61	0.34		99	99		
001413	**Very good identification**							**001413**
					OXI	NOV	PYR	
Aerococcus viridans		99.80	0.88		1	1	95	
Macrococcus carouselicus		0.15	0.08		99	99	1	
Staphylococcus cohnii subsp. cohnii		0.05	0.08		1	99	1	
001420	**Acceptable identification**							**001420**
					OXI	NOV		
Macrococcus carouselicus		93.18	0.53	XYL 1	99	99		
Aerococcus viridans		6.21	0.51		1	1		
Macrococcus equipersicus		0.33	0.08		99	99		
001421	**Good identification**							**001421**
					OXI	NOV	PYR	
Aerococcus viridans		93.00	0.58	LAC 90	1	1	95	
Macrococcus carouselicus		6.04	0.12		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		0.96	0.05		1	99	1	
001422	**Very good identification**							**001422**
					OXI	NOV		
Aerococcus viridans		98.34	0.70		1	1		
Macrococcus carouselicus		1.66	0.12		99	99		
001423	**Very good identification**							**001423**
Aerococcus viridans		100.0	0.78					
001430	**Low discrimination**							**001430**
					OXI	NOV		
Macrococcus carouselicus		57.31	0.49	XYL 1	99	99		
Aerococcus viridans		17.93	0.60		1	1		
Staphylococcus vitulinus		12.96	0.33		99	99		
001431	**Very good identification**							**001431**
					OXI	NOV	PYR	
Aerococcus viridans		97.91	0.68	LAC 90	1	1	95	
Macrococcus carouselicus		1.36	0.08		99	99	1	
Staphylococcus cohnii subsp. cohnii		0.43	0.08		1	99	1	
001432	**Very good identification**							**001432**
					OXI	NOV		
Aerococcus viridans		99.56	0.80		1	1		
Macrococcus carouselicus		0.36	0.08		99	99		
Staphylococcus vitulinus		0.08	-.08		99	99		
001433	**Very good identification**							**001433**
Aerococcus viridans		100.0	0.88					
001440	**Low discrimination**							**001440**
					OXI	YEP		
Macrococcus carouselicus		34.63	0.53	MLT 1	99	67		
Macrococcus equipersicus		32.74	0.57		99	99		
Staphylococcus cohnii subsp. cohnii		16.12	0.56		1	12		
001441	**Low discrimination**							**001441**
					OXI	NOV	PYR	
Aerococcus viridans		51.47	0.71	LAC 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		41.72	0.57		1	99	1	
Kocuria kristinae		4.55	0.27		99	1	93	
001442	**Very good identification**							**001442**
					OXI	NOV		
Aerococcus viridans		98.68	0.83		1	1		
Macrococcus carouselicus		0.42	0.12		99	99		
Macrococcus equipersicus		0.39	0.16		99	99		
001443	**Very good identification**							**001443**
					OXI	NOV	PYR	
Aerococcus viridans		99.67	0.90		1	1	95	
Kocuria kristinae		0.25	0.15		99	1	93	
Staphylococcus cohnii subsp. cohnii		0.09	0.16		1	99	1	
001450	**Low discrimination**							**001450**
					OXI	NOV		
Macrococcus equipersicus		52.02	0.84	URE 86	99	99		
Macrococcus bovicus		41.89	0.80		99	99		
Aerococcus viridans		2.32	0.73		1	1		
001451	**Low discrimination**							**001451**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI	NOV	PYR	YEP	
Aerococcus viridans		65.47	0.80	LAC 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		22.51	0.59		1	99	1	12	
Macrococcus equipersicus		6.37	0.43		99	99	1	99	
001452	**Good identification**								**001452**
					OXI	NOV			
Aerococcus viridans		94.51	0.92		1	1			
Macrococcus equipersicus		2.38	0.43		99	99			
Macrococcus bovicus		1.92	0.39		99	99			
001453	**Excellent identification**								**001453**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		99.94	1.00		1	1	95	1	
Staphylococcus cohnii subsp. cohnii		0.04	0.18		1	99	1	12	
Macrococcus equipersicus		0.01	0.02		99	99	1	99	
001460	**Good identification**								**001460**
					OXI	NOV			
Aerococcus viridans		91.17	0.63	LAC 90	1	1			
Macrococcus carouselicus		3.45	0.12		99	99			
Macrococcus equipersicus		3.27	0.16		99	99			
001461	**Very good identification**								**001461**
					NOV	PYR			
Aerococcus viridans		99.19	0.71	LAC 90	1	95			
Staphylococcus cohnii subsp. cohnii		0.81	0.16		99	1			
001462	**Very good identification**								**001462**
Aerococcus viridans		100.0	0.83						
001463	**Very good identification**								**001463**
Aerococcus viridans		100.0	0.90						
001470	**Low discrimination**								**001470**
					OXI	NOV			
Aerococcus viridans		68.99	0.73	LAC 90	1	1			
Macrococcus equipersicus		15.65	0.43		99	99			
Macrococcus bovicus		12.61	0.39		99	99			
001471	**Very good identification**								**001471**
					OXI	NOV	PYR		
Aerococcus viridans		99.23	0.80	LAC 90	1	1	95		
Staphylococcus cohnii subsp. cohnii		0.35	0.18		1	99	1		
Staphylococcus sciuri		0.25	0.11		99	99	1		
001472	**Very good identification**								**001472**
					NOV	YEP			
Aerococcus viridans		99.62	0.92		1	1			
Staphylococcus kloosii		0.27	0.40		99	99			
Staphylococcus arlettae		0.06	0.15		99	99			
001473	**Excellent identification**								**001473**
					NOV	YEP			
Aerococcus viridans		99.97	1.00		1	1			
Staphylococcus arlettae		0.04	0.18		99	99			
001500	**Very good identification**								**001500**
					OXI	NOV			
Macrococcus carouselicus		99.37	0.86		99	99			
Macrococcus equipersicus		0.35	0.41		99	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		0.16	0.51		1	1		
001501	**Low discrimination**							**001501**
					OXI	NOV	PYR	
Macrococcus carouselicus		70.80	0.45	MNS 1	99	99	1	
Aerococcus viridans		25.70	0.58		1	1	95	
Staphylococcus cohnii subsp. cohnii		2.92	0.27		1	99	1	
001502	**Low discrimination**							**001502**
					OXI	NOV		
Aerococcus viridans		58.25	0.70		1	1		
Macrococcus carouselicus		41.61	0.45		99	99		
Macrococcus equipersicus		0.15	0.00		99	99		
001503	**Very good identification**							**001503**
					OXI	NOV		
Aerococcus viridans		99.69	0.78		1	1		
Macrococcus carouselicus		0.31	0.04		99	99		
001510	**Low discrimination**							**001510**
Macrococcus carouselicus		82.55	0.82					
Macrococcus equipersicus		8.67	0.67					
Macrococcus bovicus		8.09	0.65					
001511	**Low discrimination**							**001511**
					OXI	NOV		
Aerococcus viridans		56.77	0.68	LAC 90	1	1		
Macrococcus carouselicus		33.33	0.41		99	99		
Macrococcus equipersicus		3.50	0.26		99	99		
001512	**Low discrimination**							**001512**
					OXI	NOV		
Aerococcus viridans		84.21	0.80		1	1		
Macrococcus carouselicus		12.82	0.41		99	99		
Macrococcus equipersicus		1.35	0.26		99	99		
001513	**Very good identification**							**001513**
					OXI	NOV		
Aerococcus viridans		99.84	0.88		1	1		
Staphylococcus lentus		0.10	0.12		99	99		
Macrococcus carouselicus		0.07	0.00		99	99		
001520	**Low discrimination**							**001520**
					OXI	NOV		
Macrococcus carouselicus		86.28	0.45	XYL 1	99	99		
Aerococcus viridans		13.42	0.51		1	1		
Macrococcus equipersicus		0.31	0.00		99	99		
001521	**Very good identification**							**001521**
					OXI	NOV		
Aerococcus viridans		97.29	0.58	LAC 90	1	1		
Macrococcus carouselicus		2.71	0.04		99	99		
001522	**Very good identification**							**001522**
					OXI	NOV		
Aerococcus viridans		99.28	0.70		1	1		
Macrococcus carouselicus		0.72	0.04		99	99		
001523	**Very good identification**							**001523**
Aerococcus viridans		100.0	0.78					
001530	**Low discrimination**							**001530**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI	NOV			
Macrococcus carouselicus		51.42	0.41	XYL 1	99	99			
Aerococcus viridans		37.53	0.60		1	1			
Macrococcus equipersicus		5.40	0.26		99	99			
001531	**Very good identification**								**001531**
					OXI	NOV			
Aerococcus viridans		98.90	0.68	LAC 90	1	1			
Macrococcus carouselicus		0.59	0.00		99	99			
Staphylococcus sciuri		0.52	0.05		99	99			
001532	**Very good identification**								**001532**
					OXI	NOV	YEP		
Aerococcus viridans		99.74	0.80		1	1	1		
Macrococcus carouselicus		0.15	0.00		99	99	67		
Staphylococcus kloosii		0.11	0.20		1	99	99		
001533	**Very good identification**								**001533**
					OXI	NOV			
Aerococcus viridans		99.90	0.88		1	1			
Staphylococcus lentus		0.10	0.12		99	99			
001540	**Low discrimination**								**001540**
					OXI	NOV			
Macrococcus carouselicus		35.06	0.45	MLT 1	99	99			
Macrococcus equipersicus		33.15	0.50		99	99			
Aerococcus viridans		21.81	0.63		1	1			
001541	**Low discrimination**								**001541**
					OXI	NOV	PYR		
Aerococcus viridans		88.05	0.71	LAC 90	1	1	95		
Staphylococcus cohnii subsp. cohnii		7.93	0.37		1	99	1		
Kocuria kristinae		2.59	0.17		99	1	93		
001542	**Very good identification**								**001542**
					OXI	NOV			
Aerococcus viridans		99.59	0.83		1	1			
Macrococcus carouselicus		0.18	0.04		99	99			
Macrococcus equipersicus		0.17	0.09		99	99			
001543	**Very good identification**								**001543**
					OXI				
Aerococcus viridans		99.92	0.90		1				
Kocuria kristinae		0.08	0.05		99				
001550	**Low discrimination**								**001550**
					OXI	NOV			
Macrococcus equipersicus		50.72	0.76	URE 86	99	99			
Macrococcus bovicus		40.85	0.73		99	99			
Aerococcus viridans		5.27	0.73		1	1			
001551	**Low discrimination**								**001551**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		89.01	0.80	LAC 90	1	1	95	1	
Macrococcus equipersicus		3.71	0.35		99	99	1	99	
Staphylococcus cohnii subsp. cohnii		3.40	0.39		1	99	1	12	
001552	**Very good identification**								**001552**
					OXI	NOV	YEP		
Aerococcus viridans		96.07	0.92		1	1	1		
Staphylococcus kloosii		2.02	0.59		1	99	99		
Macrococcus equipersicus		1.04	0.35		99	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
001553	**Excellent identification**							**001553**
					OXI	NOV	YEP	
Aerococcus viridans		99.93	1.00		1	1	1	
Staphylococcus lentus		0.05	0.18		99	99	1	
Staphylococcus kloosii		0.01	0.18		1	99	99	
001560	**Very good identification**							**001560**
					OXI	NOV		
Aerococcus viridans		96.94	0.63	LAC 90	1	1		
Macrococcus carouselicus		1.57	0.04		99	99		
Macrococcus equipersicus		1.49	0.09		99	99		
001561	**Very good identification**							**001561**
Aerococcus viridans		100.0	0.71	LAC 90				
001562	**Very good identification**							**001562**
Aerococcus viridans		100.0	0.83					
001563	**Very good identification**							**001563**
Aerococcus viridans		100.0	0.90					
001570	**Low discrimination**							**001570**
					OXI	NOV		
Aerococcus viridans		81.79	0.73	LAC 90	1	1		
Macrococcus equipersicus		7.95	0.35		99	99		
Macrococcus bovicus		6.40	0.32		99	99		
001571	**Very good identification**							**001571**
					OXI	NOV		
Aerococcus viridans		99.39	0.80	LAC 90	1	1		
Staphylococcus sciuri		0.49	0.17		99	99		
Staphylococcus lentus		0.13	0.07		99	99		
001572	**Very good identification**							**001572**
					NOV	YEP		
Aerococcus viridans		99.14	0.92		1	1		
Staphylococcus kloosii		0.52	0.46		99	99		
Staphylococcus arlettae		0.34	0.31		99	99		
001573	**Excellent identification**							**001573**
					OXI	NOV	YEP	
Aerococcus viridans		99.75	1.00		1	1	1	
Staphylococcus arlettae		0.20	0.34		1	99	99	
Staphylococcus lentus		0.05	0.18		99	99	1	
001600	**Excellent identification**							**001600**
					OXI	NOV		
Macrococcus carouselicus		99.90	1.00		99	99		
Aerococcus viridans		0.03	0.51		1	1		
Staphylococcus vitulinus		0.03	0.24		99	99		
001601	**Low discrimination**							**001601**
					OXI	NOV	PYR	
Macrococcus carouselicus		72.51	0.59	MNS 1	99	99	1	
Kocuria kristinae		21.61	0.48		99	1	93	
Aerococcus viridans		5.56	0.58		1	1	95	
001602	**Low discrimination**							**001602**
					OXI	NOV		
Macrococcus carouselicus		77.07	0.59	LAC 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		22.78	0.70		1	1		
Staphylococcus auricularis		0.09	0.06		1	1		
001603	**Low discrimination**							**001603**
					OXI	NOV	PYR	
Aerococcus viridans		89.07	0.78		1	1	95	
Kocuria kristinae		9.62	0.36		99	1	93	
Macrococcus carouselicus		1.31	0.18		99	99	1	
001610	**Good identification**							**001610**
Macrococcus carouselicus		94.33	0.96					
Staphylococcus vitulinus		3.89	0.65					
Macrococcus bovicus		1.14	0.60					
001611	**Low discrimination**							**001611**
					OXI	NOV		
Macrococcus carouselicus		63.88	0.55	MNS 1	99	99		
Aerococcus viridans		22.97	0.68		1	1		
Staphylococcus sciuri		8.08	0.43		99	99		
001612	**Low discrimination**							**001612**
					OXI	NOV		
Aerococcus viridans		56.65	0.80		1	1		
Macrococcus carouselicus		40.86	0.55		99	99		
Staphylococcus vitulinus		1.68	0.24		99	99		
001613	**Very good identification**							**001613**
					OXI	NOV		
Aerococcus viridans		98.82	0.88		1	1		
Staphylococcus sciuri		0.63	0.26		99	99		
Macrococcus carouselicus		0.31	0.14		99	99		
001620	**Good identification**							**001620**
					OXI	NOV		
Macrococcus carouselicus		90.58	0.59	XYL 1	99	99		
Staphylococcus vitulinus		6.45	0.33		99	99		
Aerococcus viridans		2.97	0.51		1	1		
001621	**Low discrimination**							**001621**
					OXI	NOV	PYR	
Aerococcus viridans		82.98	0.58	LAC 90	1	1	95	
Macrococcus carouselicus		10.94	0.18		99	99	1	
Kocuria kristinae		3.26	0.07		99	1	93	
001622	**Very good identification**							**001622**
					OXI	NOV		
Aerococcus viridans		96.47	0.70		1	1		
Macrococcus carouselicus		3.30	0.18		99	99		
Staphylococcus vitulinus		0.24	-.08		99	99		
001623	**Very good identification**							**001623**
					OXI			
Aerococcus viridans		99.89	0.78		1			
Kocuria kristinae		0.11	-.05		99			
001630	**Good identification**							**001630**
					OXI	NOV		
Staphylococcus vitulinus		90.31	0.74	NIT 91	99	99		
Macrococcus carouselicus		8.19	0.55		99	99		
Aerococcus viridans		1.26	0.60		1	1		
001631	**Low discrimination**							**001631**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Aerococcus viridans		62.36	0.68	LAC 90	1	1		
Staphylococcus vitulinus		19.32	0.33		99	99		
Staphylococcus sciuri		16.56	0.40		99	99		
001632	**Good identification**							**001632**
					OXI	NOV		
Aerococcus viridans		91.62	0.80		1	1		
Staphylococcus vitulinus		7.36	0.33		99	99		
Macrococcus carouselicus		0.67	0.14		99	99		
001633	**Very good identification**							**001633**
					OXI	NOV	YEP	
Aerococcus viridans		99.25	0.88		1	1	1	
Staphylococcus sciuri		0.48	0.24		99	99	57	
Staphylococcus arlettae		0.14	0.18		1	99	99	
001640	**Low discrimination**							**001640**
					OXI	NOV		
Macrococcus carouselicus		70.57	0.59	MLT 1	99	99		
Staphylococcus petrasii subsp. pragensis		11.67	0.33		1	1		
Aerococcus viridans		9.27	0.63		1	1		
001641	**Low discrimination**							**001641**
					OXI	NOV	PYR	
Kocuria kristinae		88.93	0.68	VPT 90	99	1	93	
Aerococcus viridans		10.16	0.71		1	1	95	
Macrococcus carouselicus		0.34	0.18		99	99	1	
001642	**Very good identification**							**001642**
					OXI	NOV	PYR	
Aerococcus viridans		98.22	0.83		1	1	95	
Macrococcus carouselicus		0.84	0.18		99	99	1	
Kocuria kristinae		0.56	0.15		99	1	93	
001643	**Low discrimination**							**001643**
					OXI	NOV	PYR	
Aerococcus viridans		80.41	0.90		1	1	95	
Kocuria kristinae		19.55	0.56		99	1	93	
Staphylococcus sciuri		0.05	0.07		99	99	1	
001650	**Low discrimination**							**001650**
Macrococcus bovicus		47.34	0.68	ESL 10				
Macrococcus equipersicus		23.26	0.63					
Macrococcus carouselicus		16.99	0.55					
001651	**Low discrimination**							**001651**
					OXI	NOV	PYR	
Aerococcus viridans		66.96	0.80	LAC 90	1	1	95	
Staphylococcus sciuri		22.16	0.55		99	99	1	
Kocuria kristinae		8.14	0.40		99	1	93	
001652	**Very good identification**							**001652**
					OXI	NOV		
Aerococcus viridans		98.74	0.92		1	1		
Macrococcus bovicus		0.50	0.27		99	99		
Macrococcus equipersicus		0.25	0.22		99	99		
001653	**Very good identification**							**001653**
					OXI	NOV	PYR	
Aerococcus viridans		98.99	1.00		1	1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		0.59	0.38		99	99	1	
Kocuria kristinae		0.33	0.27		99	1	93	
001660	**Good identification**							**001660**
					OXI	NOV		
Aerococcus viridans		90.76	0.63	LAC 90	1	1		
Macrococcus carouselicus		6.98	0.18		99	99		
Staphylococcus petrasii subsp. pragensis		1.15	-.09		1	1		
001661	**Low discrimination**							**001661**
					OXI	NOV	PYR	
Aerococcus viridans		89.96	0.71	LAC 90	1	1	95	
Kocuria kristinae		7.95	0.27		99	1	93	
Staphylococcus sciuri		2.08	0.21		99	99	1	
001662	**Very good identification**							**001662**
					NOV	YEP		
Aerococcus viridans		99.82	0.83		1	1		
Staphylococcus arlettae		0.18	0.15		99	99		
001663	**Very good identification**							**001663**
					OXI	NOV	YEP	
Aerococcus viridans		99.61	0.90		1	1	1	
Kocuria kristinae		0.25	0.15		99	1	15	
Staphylococcus arlettae		0.10	0.18		1	99	99	
001670	**Low discrimination**							**001670**
					OXI	NOV		
Aerococcus viridans		74.35	0.73	LAC 90	1	1		
Staphylococcus vitulinus		13.44	0.33		99	99		
Staphylococcus sciuri		5.36	0.33		99	99		
001671	**Low discrimination**							**001671**
					OXI	NOV	YEP	
Aerococcus viridans		79.61	0.80	LAC 90	1	1	1	
Staphylococcus sciuri		19.88	0.52		99	99	57	
Staphylococcus arlettae		0.25	0.18		1	99	99	
001672	**Good identification**							**001672**
					OXI	NOV	YEP	
Aerococcus viridans		94.13	0.92		1	1	1	
Staphylococcus arlettae		5.71	0.57		1	99	99	
Staphylococcus sciuri		0.12	0.17		99	99	57	
001673	**Very good identification**							**001673**
					OXI	NOV	YEP	
Aerococcus viridans		96.20	1.00		1	1	1	
Staphylococcus arlettae		3.31	0.59		1	99	99	
Staphylococcus sciuri		0.44	0.36		99	99	57	
001700	**Very good identification**							**001700**
					OXI	NOV		
Macrococcus carouselicus		99.87	0.92		99	99		
Aerococcus viridans		0.08	0.51		1	1		
Rothia mucilaginosa		0.03	0.15		1	1		
001701	**Low discrimination**							**001701**
					OXI	NOV	PYR	
Macrococcus carouselicus		66.64	0.51	MNS 1	99	99	1	
Kocuria kristinae		15.44	0.39		99	1	93	
Aerococcus viridans		11.91	0.58		1	1	95	
001702	**Low discrimination**							**001702**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Macrococcus carouselicus		59.19	0.51	LAC 1	99	99		
Aerococcus viridans		40.81	0.70		1	1		
001703	**Very good identification**							**001703**
					OXI	NOV	PYR	
Aerococcus viridans		95.57	0.78		1	1	95	
Kocuria kristinae		3.44	0.26		99	1	93	
Macrococcus carouselicus		0.60	0.10		99	99	1	
001710	**Very good identification**							**001710**
Macrococcus carouselicus		97.80	0.88					
Macrococcus bovicus		1.18	0.53					
Macrococcus equipersicus		0.50	0.47					
001711	**Low discrimination**							**001711**
					OXI	NOV		
Macrococcus carouselicus		37.71	0.47	MNS 1	99	99		
Aerococcus viridans		31.63	0.68		1	1		
Staphylococcus sciuri		21.61	0.49		99	99		
001712	**Low discrimination**							**001712**
					OXI	NOV		
Aerococcus viridans		75.74	0.80		1	1		
Macrococcus carouselicus		23.41	0.47		99	99		
Macrococcus bovicus		0.28	0.12		99	99		
001713	**Low discrimination**							**001713**
					OXI	NOV		
Aerococcus viridans		89.96	0.88		1	1		
Staphylococcus lentus		8.76	0.53		99	99		
Staphylococcus sciuri		1.11	0.32		99	99		
001720	**Acceptable identification**							**001720**
					OXI	NOV		
Macrococcus carouselicus		92.74	0.51	XYL 1	99	99		
Aerococcus viridans		7.11	0.51		1	1		
Staphylococcus vitulinus		0.16	-.08		99	99		
001721	**Low discrimination**							**001721**
					OXI	NOV		
Aerococcus viridans		89.49	0.58	LAC 90	1	1		
Macrococcus carouselicus		5.06	0.10		99	99		
Staphylococcus sciuri		4.28	0.15		99	99		
001722	**Very good identification**							**001722**
					OXI	NOV		
Aerococcus viridans		98.56	0.70		1	1		
Macrococcus carouselicus		1.44	0.10		99	99		
001723	**Very good identification**							**001723**
					OXI	NOV		
Aerococcus viridans		99.71	0.78		1	1		
Staphylococcus lentus		0.29	0.12		99	99		
001730	**Low discrimination**							**001730**
					OXI	NOV		
Macrococcus carouselicus		59.17	0.47	XYL 1	99	99		
Aerococcus viridans		21.27	0.60		1	1		
Staphylococcus vitulinus		15.38	0.33		99	99		
001731	**Low discrimination**							**001731**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Aerococcus viridans		56.25	0.68	LAC 90	1	1		
Staphylococcus sciuri		28.99	0.46		99	99		
Staphylococcus lentus		13.91	0.41		99	99		
001732	**Very good identification**							**001732**
					OXI	NOV	YEP	
Aerococcus viridans		97.77	0.80		1	1	1	
Staphylococcus arlettae		1.36	0.31		1	99	99	
Macrococcus carouselicus		0.31	0.06		99	99	67	
001733	**Low discrimination**							**001733**
					OXI	NOV		
Aerococcus viridans		89.72	0.88		1	1		
Staphylococcus lentus		8.74	0.53		99	99		
Staphylococcus sciuri		0.84	0.30		99	99		
001740	**Low discrimination**							**001740**
					OXI	NOV		
Macrococcus carouselicus		68.09	0.51	MLT 1	99	99		
Aerococcus viridans		20.87	0.63		1	1		
Rothia mucilaginosa		4.17	0.23		1	1		
001741	**Low discrimination**							**001741**
					OXI			
Kocuria kristinae		68.61	0.58	VPT 90	99			
Aerococcus viridans		23.52	0.71		1			
Rothia mucilaginosa		6.04	0.33		1			
001742	**Very good identification**							**001742**
					OXI	NOV	PYR	
Aerococcus viridans		99.45	0.83		1	1	95	
Macrococcus carouselicus		0.36	0.10		99	99	1	
Kocuria kristinae		0.19	0.05		99	1	93	
001743	**Good identification**							**001743**
					OXI	NOV	PYR	
Aerococcus viridans		92.26	0.90		1	1	95	
Kocuria kristinae		7.48	0.46		99	1	93	
Staphylococcus lentus		0.14	0.18		99	99	6	
001750	**Low discrimination**							**001750**
					OXI	NOV		
Macrococcus bovicus		40.45	0.60	ESL 10	99	99		
Aerococcus viridans		20.87	0.73		1	1		
Macrococcus equipersicus		19.87	0.56		99	99		
001751	**Low discrimination**							**001751**
					OXI	NOV		
Aerococcus viridans		54.80	0.80	LAC 90	1	1		
Staphylococcus sciuri		35.21	0.61		99	99		
Staphylococcus lentus		6.88	0.48		99	99		
001752	**Very good identification**							**001752**
					OXI	NOV	YEP	
Aerococcus viridans		98.61	0.92		1	1	1	
Staphylococcus arlettae		0.34	0.31		1	99	99	
Staphylococcus sciuri		0.33	0.26		99	99	57	
001753	**Good identification**							**001753**
					OXI	NOV		
Aerococcus viridans		93.97	1.00		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		4.65	0.59		99	99		
Staphylococcus sciuri		1.09	0.44		99	99		
001760	**Very good identification**							**001760**
					OXI	NOV		
Aerococcus viridans		95.61	0.63	LAC 90	1	1		
Macrococcus carouselicus		3.15	0.10		99	99		
Staphylococcus sciuri		1.24	0.08		99	99		
001761	**Good identification**							**001761**
					OXI	NOV	PYR	
Aerococcus viridans		92.52	0.71	LAC 90	1	1	95	
Staphylococcus sciuri		4.16	0.27		99	99	1	
Kocuria kristinae		2.73	0.17		99	1	93	
001762	**Very good identification**							**001762**
					NOV	YEP		
Aerococcus viridans		98.97	0.83		1	1		
Staphylococcus arlettae		1.03	0.31		99	99		
001763	**Very good identification**							**001763**
					OXI	NOV	YEP	
Aerococcus viridans		99.10	0.90		1	1	1	
Staphylococcus arlettae		0.59	0.34		1	99	99	
Staphylococcus lentus		0.15	0.18		99	99	1	
001770	**Low discrimination**							**001770**
					OXI	NOV	YEP	
Aerococcus viridans		82.21	0.73	LAC 90	1	1	1	
Staphylococcus sciuri		11.49	0.39		99	99	57	
Staphylococcus arlettae		2.57	0.31		1	99	99	
001771	**Low discrimination**							**001771**
					OXI	NOV		
Aerococcus viridans		61.35	0.80	LAC 90	1	1		
Staphylococcus sciuri		29.74	0.58		99	99		
Staphylococcus lentus		7.70	0.48		99	99		
001772	**Low discrimination**							**001772**
					OXI	NOV	YEP	
Aerococcus viridans		74.20	0.92		1	1	1	
Staphylococcus arlettae		25.49	0.72		1	99	99	
Staphylococcus sciuri		0.19	0.23		99	99	57	
001773	**Low discrimination**							**001773**
					OXI	NOV	YEP	
Aerococcus viridans		79.75	1.00		1	1	1	
Staphylococcus arlettae		15.57	0.75		1	99	99	
Staphylococcus lentus		3.94	0.59		99	99	1	
002000	**Low discrimination**							**002000**
Kocuria rosea		59.09	1.00					
Dermacoccus nishinomiyaensis		28.88	0.95					
Micrococcus sp.		7.61	0.74					
002001	**Low discrimination**							**002001**
Kocuria rosea		70.22	0.74	MNS 5				
Staphylococcus capitis subsp. capitis		11.21	0.60					
Kocuria varians		7.09	0.62					
002002	**Low discrimination**							**002002**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Kocuria varians		49.89	0.73		13			
Kocuria rosea		29.22	0.59		35			
Dermacoccus nishinomiyaensis		14.28	0.54		93			
002003	**Low discrimination**							**002003**
Kocuria varians		62.93	0.51	MNS 8				
Kocuria rosea		22.31	0.33					
Staphylococcus capitis subsp. capitis		7.20	0.25					
002010	**Low discrimination**							**002010**
					YEP	NOV		
Kocuria rosea		79.52	0.80	MAN 10	1	1		
Macrococcus bovicus		13.33	0.69		99	99		
Dermacoccus nishinomiyaensis		3.53	0.54		1	80		
002011	**Good identification**							**002011**
					NOV	YEP		
Staphylococcus capitis subsp. capitis		94.69	0.83		1	1		
Kocuria rosea		4.96	0.54		1	1		
Macrococcus bovicus		0.16	0.28		99	99		
002012	**Low discrimination**							**002012**
					NOV	YEP	OXI	
Kocuria rosea		70.31	0.39	MAN 10	1	1	35	
Macrococcus bovicus		11.78	0.28	LAC 1	99	99	99	
Kocuria varians		10.91	0.32		1	80	13	
002013	**Good identification**							**002013**
Staphylococcus capitis subsp. capitis		96.85	0.48	LAC 2				
Kocuria rosea		2.51	0.13					
Kocuria varians		0.64	0.10					
002020	**Good identification**							**002020**
					OXI			
Kocuria rosea		93.89	0.92		35			
Kocuria varians		4.71	0.75		13			
Dermacoccus nishinomiyaensis		1.08	0.54		93			
002021	**Good identification**							**002021**
Kocuria rosea		91.71	0.66	MNS 5				
Kocuria varians		7.60	0.53					
Staphylococcus capitis subsp. capitis		0.35	0.19					
002022	**Low discrimination**							**002022**
					OXI			
Kocuria varians		57.93	0.63		13			
Kocuria rosea		41.38	0.51		35			
Dermacoccus nishinomiyaensis		0.48	0.13		93			
002023	**Low discrimination**							**002023**
Kocuria varians		69.81	0.42	MNS 8				
Kocuria rosea		30.19	0.25					
002030	**Good identification**							**002030**
					NOV	OXI		
Kocuria rosea		90.83	0.73	MAN 10	1	35		
Staphylococcus vitulinus		8.26	0.50		99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		0.41	0.34		1	13		
002031	**Low discrimination**							**002031**
					NOV	OXI		
Kocuria rosea		67.80	0.47	MAN 10	1	35		
Staphylococcus capitis subsp. capitis		30.51	0.42	MNS 5	1	1		
Staphylococcus vitulinus		1.18	0.09		99	99		
002032	**Low discrimination**							**002032**
					NOV	OXI		
Kocuria rosea		82.09	0.32	MAN 10	1	35		
Kocuria varians		10.45	0.22	LAC 1	1	13		
Staphylococcus vitulinus		7.46	0.09		99	99		
002040	**Low discrimination**							**002040**
Staphylococcus auricularis		33.62	0.73	TRE 92				
Kocuria varians		22.95	0.70					
Kocuria rosea		17.27	0.59					
002041	**Low discrimination**							**002041**
Staphylococcus auricularis		49.81	0.53	TRE 92				
Kocuria varians		26.60	0.49	MNS 10				
Kocuria rosea		12.12	0.33					
002042	**Low discrimination**							**002042**
Kocuria varians		60.77	0.59					
Staphylococcus auricularis		35.08	0.53					
Kocuria rosea		1.64	0.18					
002043	**Low discrimination**							**002043**
Kocuria varians		57.55	0.37	MNS 8				
Staphylococcus auricularis		42.45	0.33					
002050	**Very good identification**							**002050**
					NOV	YEP		
Macrococcus bovicus		95.14	0.77	NIT 10	99	99		
Kocuria rosea		2.46	0.39		1	1		
Macrococcus equipersicus		0.84	0.36		99	99		
002051	**Low discrimination**							**002051**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		60.08	0.42	MLT 1	1	1	1	
Macrococcus bovicus		23.38	0.36		99	99	99	
Staphylococcus cohnii subsp. cohnii		11.59	0.36		1	99	12	
002052	**Low discrimination**							**002052**
					NOV	OXI		
Macrococcus bovicus		85.97	0.36	NIT 10	99	99		
Kocuria varians		7.49	0.18	LAC 1	1	13		
Staphylococcus auricularis		4.32	0.12		1	1		
002060	**Low discrimination**							**002060**
Kocuria varians		50.22	0.61					
Kocuria rosea		46.10	0.51					
Staphylococcus auricularis		2.12	0.32					
002061	**Low discrimination**							**002061**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		62.13	0.39	MNS 8				
Kocuria rosea		34.52	0.25					
Staphylococcus auricularis		3.34	0.12					
002062	**Good identification**							**002062**
Kocuria varians		95.29	0.50					
Kocuria rosea		3.13	0.10					
Staphylococcus auricularis		1.58	0.12					
002063	**Good identification**							**002063**
Kocuria varians		100.0	0.28	MNS 8				
002070	**Low discrimination**							**002070**
					NOV	YEP	OXI	
Kocuria rosea		47.57	0.32	MAN 10	1	1	35	
Micrococcus bovicus		43.40	0.36	MLT 1	99	99	99	
Kocuria varians		4.71	0.20		1	80	13	
002100	**Low discrimination**							**002100**
					OXI			
Dermacoccus nishinomiyaensis		47.18	0.80		93			
Kocuria varians		46.48	0.88		13			
Kocuria rosea		5.12	0.59		35			
002101	**Low discrimination**							**002101**
					OXI			
Kocuria varians		82.49	0.66	MNS 8	13			
Dermacoccus nishinomiyaensis		9.73	0.39		93			
Kocuria rosea		5.50	0.33		35			
002102	**Very good identification**							**002102**
					OXI			
Kocuria varians		95.71	0.76		13			
Dermacoccus nishinomiyaensis		3.48	0.39		93			
Staphylococcus auricularis		0.39	0.26		1			
002103	**Very good identification**							**002103**
					OXI			
Kocuria varians		99.07	0.55	MNS 8	13			
Staphylococcus auricularis		0.51	0.06		1			
Dermacoccus nishinomiyaensis		0.42	-.02		93			
002110	**Low discrimination**							**002110**
					YEP	NOV		
Micrococcus bovicus		71.77	0.62	NIT 10	99	99		
Kocuria rosea		10.09	0.39		1	1		
Dermacoccus nishinomiyaensis		8.46	0.39		1	80		
002111	**Low discrimination**							**002111**
					NOV	OXI		
Staphylococcus capitis subsp. capitis		82.61	0.42	GAL 1	1	1		
Micrococcus bovicus		5.90	0.21		99	99		
Kocuria varians		5.90	0.25		1	13		
002112	**Low discrimination**							**002112**
					NOV	YEP	OXI	
Kocuria varians		72.04	0.35	MAN 1	1	80	13	
Micrococcus bovicus		22.22	0.21		99	99	99	
Kocuria rosea		3.13	-.02		1	1	35	
002120	**Low discrimination**							**002120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Kocuria varians		85.90	0.78		13			
Kocuria rosea		11.54	0.51		35			
Dermacoccus nishinomiyaensis		2.51	0.39		93			
002121	**Good identification**							**002121**
					OXI			
Kocuria varians		92.19	0.56	MNS 8	13			
Kocuria rosea		7.50	0.25		35			
Dermacoccus nishinomiyaensis		0.31	-.02		93			
002122	**Very good identification**							**002122**
					OXI			
Kocuria varians		99.42	0.67		13			
Kocuria rosea		0.48	0.10		35			
Dermacoccus nishinomiyaensis		0.10	-.02		93			
002123	**Good identification**							**002123**
Kocuria varians		100.0	0.45	MNS 8				
002130	**Low discrimination**							**002130**
					NOV	YEP	OXI	
Kocuria rosea		51.15	0.32	GAL 1	1	1	35	
Kocuria varians		34.62	0.37	MAN 10	1	80	13	
Macrococcus bovicus		8.57	0.21		99	99	99	
002140	**Low discrimination**							**002140**
					OXI			
Kocuria varians		84.65	0.74		13			
Staphylococcus auricularis		9.19	0.53		1			
Dermacoccus nishinomiyaensis		3.95	0.39		93			
002141	**Low discrimination**							**002141**
					NOV			
Kocuria varians		86.58	0.52	MNS 8	1			
Staphylococcus auricularis		12.01	0.33		1			
Staphylococcus cohnii subsp. cohnii		0.94	0.14		99			
002142	**Good identification**							**002142**
					NOV	OXI		
Kocuria varians		93.38	0.63		1	13		
Staphylococcus auricularis		3.99	0.33		1	1		
Macrococcus caseolyticus		2.21	0.18		99	99		
002143	**Acceptable identification**							**002143**
Kocuria varians		94.82	0.41	MNS 8				
Staphylococcus auricularis		5.18	0.14					
002150	**Very good identification**							**002150**
					NOV	OXI		
Macrococcus bovicus		97.68	0.69	NIT 10	99	99		
Kocuria varians		1.07	0.33		1	13		
Macrococcus equipersicus		0.87	0.28		99	99		
002151	**Low discrimination**							**002151**
					NOV	OXI	YEP	
Macrococcus bovicus		77.93	0.28	NIT 10	99	99	99	
Staphylococcus cohnii subsp. cohnii		10.02	0.16	MNS 1	99	1	12	
Kocuria varians		7.33	0.11		1	13	80	
002152	**Low discrimination**							**002152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	OXI	
Macrococcus bovicus		76.63	0.28	NIT 10	99	99	
Kocuria varians		23.37	0.22	LAC 1	1	13	
002160	**Very good identification**						**002160**
Kocuria varians		98.95	0.65				
Kocuria rosea		0.61	0.10				
Staphylococcus auricularis		0.31	0.12				
002161	**Good identification**						**002161**
Kocuria varians		100.0	0.43	MNS 8			
002162	**Very good identification**						**002162**
Kocuria varians		100.0	0.53				
002163	**Good identification**						**002163**
Kocuria varians		100.0	0.32	MNS 8			
002170	**Low discrimination**						**002170**
					NOV	OXI	
Macrococcus bovicus		72.47	0.28	NIT 10	99	99	
Kocuria varians		27.53	0.24	XYL 1	1	13	
002200	**Low discrimination**						**002200**
Kocuria rosea		64.13	0.80	SUC 10			
Dermacoccus nishinomiyaensis		18.01	0.70				
Micrococcus sp.		7.35	0.53				
002201	**Low discrimination**						**002201**
Staphylococcus capitis subsp. capitis		65.91	0.65	MAN 93			
Kocuria rosea		24.70	0.54				
Staphylococcus auricularis		3.65	0.45				
002202	**Low discrimination**						**002202**
Kocuria varians		48.92	0.55	SUC 12			
Kocuria rosea		23.35	0.39				
Staphylococcus auricularis		17.98	0.45				
002203	**Low discrimination**						**002203**
Staphylococcus capitis subsp. capitis		46.24	0.31	MAN 93			
Kocuria varians		29.68	0.33	LAC 2			
Staphylococcus auricularis		13.94	0.25				
002210	**Low discrimination**						**002210**
					NOV	YEP	
Staphylococcus vitulinus		88.87	0.82		99	75	
Kocuria rosea		6.92	0.61		1	1	
Macrococcus bovicus		2.61	0.57		99	99	
002211	**Very good identification**						**002211**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		98.89	0.88		1	1	
Staphylococcus vitulinus		0.76	0.41		99	99	
Kocuria rosea		0.31	0.34		1	35	
002212	**Low discrimination**						**002212**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus vitulinus		86.64	0.41	LAC 1	99	75	
Kocuria rosea		6.75	0.20		1	1	
Macrococcus bovicus		2.54	0.16		99	99	
002213	**Very good identification**						**002213**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		99.47	0.54	LAC 2	1	1	
Staphylococcus vitulinus		0.38	0.00		99	99	
Kocuria rosea		0.15	-.07		1	35	
002220	**Low discrimination**						**002220**
					NOV	OXI	
Kocuria rosea		85.44	0.73	SUC 10	1	35	
Staphylococcus vitulinus		7.77	0.50		99	99	
Kocuria varians		5.26	0.57		1	13	
002221	**Low discrimination**						**002221**
Kocuria rosea		84.35	0.47	SUC 10			
Kocuria varians		8.57	0.35	MNS 5			
Staphylococcus capitis subsp. capitis		5.31	0.24				
002222	**Low discrimination**						**002222**
					NOV	OXI	
Kocuria varians		60.77	0.46	SUC 12	1	13	
Kocuria rosea		35.37	0.32		1	35	
Staphylococcus vitulinus		3.22	0.09		99	99	
002230	**Very good identification**						**002230**
					NOV	OXI	
Staphylococcus vitulinus		98.76	0.91		99	99	
Kocuria rosea		1.22	0.53		1	35	
Kocuria varians		0.01	0.16		1	13	
002231	**Low discrimination**						**002231**
					NOV	OXI	
Staphylococcus vitulinus		64.51	0.50	MNS 1	99	99	
Staphylococcus capitis subsp. capitis		31.21	0.47		1	1	
Kocuria rosea		4.15	0.27		1	35	
002232	**Good identification**						**002232**
					NOV	OXI	
Staphylococcus vitulinus		98.59	0.50	LAC 1	99	99	
Kocuria rosea		1.22	0.12		1	35	
Kocuria varians		0.19	0.04		1	13	
002240	**Low discrimination**						**002240**
Staphylococcus auricularis		74.55	0.72	TRE 92			
Kocuria varians		7.52	0.53				
Kocuria rosea		4.61	0.39				
002241	**Low discrimination**						**002241**
					OXI	PYR	
Staphylococcus auricularis		68.51	0.52	TRE 92	1	40	
Kocuria kristinae		13.20	0.31	MNS 10	99	93	
Kocuria varians		5.41	0.31		13	10	
002242	**Low discrimination**						**002242**
Staphylococcus auricularis		52.35	0.52	TRE 92			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		22.15	0.47	LAC 10				
Kocuria varians		13.40	0.41					
002243	**Low discrimination**							**002243**
					OXI	NOV		
Staphylococcus auricularis		45.76	0.33	TRE 92	1	1		
Kocuria kristinae		19.84	0.19	MNS 10	99	1		
Staphylococcus hominis subsp. novobiosepticus		11.39	0.12	LAC 10	1	99		
002250	**Low discrimination**							**002250**
					OXI	PYR		
Macrococcus bovicus		82.35	0.64	NIT 10	99	1		
Staphylococcus vitulinus		12.14	0.41		99	1		
Staphylococcus haemolyticus		1.57	0.33		1	98		
002251	**Acceptable identification**							**002251**
					OXI	NOV		
Staphylococcus capitis subsp. capitis		90.57	0.47	MLT 1	1	1		
Macrococcus bovicus		4.74	0.23		99	99		
Staphylococcus sciuri		1.84	0.17		99	99		
002252	**Low discrimination**							**002252**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		32.66	0.30	URE 99	1	1	1	
Staphylococcus haemolyticus		29.88	0.28	MAN 13	1	98	2	
Macrococcus bovicus		28.10	0.23		99	1	99	
002260	**Low discrimination**							**002260**
Kocuria varians		47.00	0.43	SUC 12				
Kocuria rosea		35.16	0.32					
Staphylococcus auricularis		13.40	0.31					
002262	**Low discrimination**							**002262**
Kocuria varians		86.22	0.32	SUC 12				
Staphylococcus auricularis		9.69	0.11					
Staphylococcus hominis subsp. hominis		4.10	0.06					
002270	**Good identification**							**002270**
					NOV	YEP		
Staphylococcus vitulinus		96.23	0.50	MLT 1	99	75		
Macrococcus bovicus		2.44	0.23		99	99		
Kocuria rosea		1.19	0.12		1	1		
002300	**Low discrimination**							**002300**
					OXI			
Kocuria varians		60.32	0.70	SUC 12	13			
Dermacoccus nishinomiyaensis		28.66	0.55		93			
Kocuria rosea		5.41	0.39		35			
002301	**Low discrimination**							**002301**
Kocuria varians		72.07	0.48	SUC 12				
Staphylococcus capitis subsp. capitis		10.45	0.24	MNS 8				
Staphylococcus auricularis		6.37	0.25					
002302	**Very good identification**							**002302**
					OXI			
Kocuria varians		95.06	0.59	SUC 12	13			
Staphylococcus auricularis		2.59	0.25		1			
Dermacoccus nishinomiyaensis		1.62	0.14		93			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
002303	**Low discrimination**						**002303**
Kocuria varians		80.97	0.37	SUC 12			
Staphylococcus hyicus		16.21	0.13	MNS 8			
Staphylococcus auricularis		2.82	0.05				
002310	**Low discrimination**						**002310**
					NOV	OXI	
Macrococcus bovicus		49.78	0.49	NIT 10	99	99	
Staphylococcus vitulinus		39.96	0.41		99	99	
Kocuria varians		3.15	0.29		1	13	
002311	**Good identification**						**002311**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		96.51	0.47	GAL 1	1	1	
Staphylococcus sciuri		1.01	0.11		99	99	
Macrococcus bovicus		0.93	0.08		99	99	
002320	**Low discrimination**						**002320**
					OXI		
Kocuria varians		88.10	0.60	SUC 12	13		
Kocuria rosea		9.64	0.32		35		
Dermacoccus nishinomiyaensis		1.20	0.14		93		
002321	**Acceptable identification**						**002321**
Kocuria varians		93.79	0.39	SUC 12			
Kocuria rosea		6.21	0.05	MNS 8			
002322	**Good identification**						**002322**
Kocuria varians		100.0	0.49	SUC 12			
002323	**Good identification**						**002323**
Kocuria varians		100.0	0.27	SUC 12			
002330	**Good identification**						**002330**
					NOV	OXI	
Staphylococcus vitulinus		97.35	0.50	GAL 1	99	99	
Kocuria rosea		1.20	0.12		1	35	
Kocuria varians		1.00	0.19		1	13	
002340	**Low discrimination**						**002340**
					NOV		
Kocuria varians		36.86	0.56	SUC 12	1		
Staphylococcus auricularis		27.08	0.52		1		
Staphylococcus hominis subsp. novobiosepticus		16.69	0.39		99		
002341	**Low discrimination**						**002341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		28.48	0.27	VPT 90	99		
Kocuria varians		21.89	0.34	URE 99	1		
Staphylococcus auricularis		20.55	0.33		1		
002342	**Low discrimination**						**002342**
					NOV		
Staphylococcus hominis subsp. hominis		30.40	0.48	URE 99	1		
Kocuria varians		25.47	0.45		1		
Staphylococcus hominis subsp. novobiosepticus		22.01	0.34		99		
002343	**Low discrimination**						**002343**
					NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		44.15	0.33	VPT 96	1			
Staphylococcus hominis subsp. novobiosepticus		29.41	0.21	URE 92	99			
Kocuria varians		11.84	0.23		1			
002350	**Good identification**							**002350**
					OXI	PYR	NOV	
Micrococcus bovicus		90.51	0.57	NIT 10	99	1	99	
Staphylococcus haemolyticus		4.55	0.34		1	98	2	
Staphylococcus hominis subsp. hominis		2.29	0.30		1	1	1	
002352	**Low discrimination**							**002352**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		40.68	0.31	URE 99	1	1	1	
Staphylococcus haemolyticus		38.74	0.29	MAN 13	1	98	2	
Micrococcus bovicus		13.85	0.16		99	1	99	
002360	**Good identification**							**002360**
					NOV			
Kocuria varians		95.93	0.47	SUC 12	1			
Staphylococcus auricularis		2.03	0.11		1			
Staphylococcus hominis subsp. novobiosepticus		1.25	-.02		99			
002362	**Acceptable identification**							**002362**
					NOV			
Kocuria varians		94.41	0.36	SUC 12	1			
Staphylococcus hominis subsp. hominis		3.24	0.07		1			
Staphylococcus hominis subsp. novobiosepticus		2.35	-.07		99			
002400	**Low discrimination**							**002400**
Staphylococcus auricularis		64.19	0.87					
Staphylococcus carnosus subsp. utilis		17.27	0.59					
Kocuria rosea		6.69	0.59					
002401	**Low discrimination**							**002401**
					NOV			
Staphylococcus auricularis		61.22	0.67	MNS 10	1			
Staphylococcus cohnii subsp. cohnii		30.39	0.64		99			
Kocuria rosea		3.02	0.33		1			
002402	**Low discrimination**							**002402**
Staphylococcus auricularis		85.66	0.67	LAC 10				
Kocuria varians		10.33	0.50					
Staphylococcus carnosus subsp. utilis		2.10	0.18					
002403	**Low discrimination**							**002403**
					NOV			
Staphylococcus auricularis		87.31	0.48	MNS 10	1			
Kocuria varians		8.24	0.28	LAC 10	1			
Staphylococcus cohnii subsp. cohnii		3.94	0.23		99			
002410	**Low discrimination**							**002410**
					OXI	YEP		
Micrococcus bovicus		71.78	0.73	NIT 10	99	99		
Staphylococcus cohnii subsp. cohnii		16.14	0.66		1	12		
Micrococcus carouselicus		2.99	0.41		99	67		
002411	**Low discrimination**							**002411**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		71.22	0.67	NIT 7	1	99	12	
Staphylococcus capitis subsp. capitis		23.85	0.48		1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		2.95	0.32		99	99	99	
002412	**Low discrimination**							**002412**
					OXI	NOV	YEP	
Macrococcus bovicus		55.46	0.32	NIT 10	99	99	99	
Staphylococcus auricularis		21.38	0.26	LAC 1	1	1	1	
Staphylococcus cohnii subsp. cohnii		12.47	0.25		1	99	12	
002420	**Low discrimination**							**002420**
Kocuria rosea		56.39	0.51	TRE 1				
Kocuria varians		21.06	0.52					
Staphylococcus auricularis		12.75	0.46					
002422	**Low discrimination**							**002422**
Kocuria varians		74.94	0.40	TRE 7				
Staphylococcus auricularis		17.88	0.26					
Kocuria rosea		7.19	0.10					
002430	**Low discrimination**							**002430**
					NOV	YEP		
Staphylococcus vitulinus		73.80	0.45	SUC 99	99	75		
Kocuria rosea		14.58	0.32		1	1		
Macrococcus bovicus		8.55	0.32		99	99		
002433	**Good identification**							**002433**
Aerococcus viridans		100.0	0.27	ESL 90				
002440	**Good identification**							**002440**
					NOV			
Staphylococcus auricularis		92.56	0.95		1			
Staphylococcus cohnii subsp. cohnii		6.40	0.74		99			
Kocuria varians		0.41	0.47		1			
002441	**Low discrimination**							**002441**
					NOV			
Staphylococcus auricularis		59.51	0.75	MNS 10	1			
Staphylococcus cohnii subsp. cohnii		40.09	0.75		99			
Kocuria varians		0.21	0.26		1			
002442	**Very good identification**							**002442**
					NOV			
Staphylococcus auricularis		97.45	0.75	LAC 10	1			
Kocuria varians		1.11	0.36		1			
Staphylococcus cohnii subsp. cohnii		0.61	0.33		99			
002443	**Good identification**							**002443**
					NOV	PYR		
Staphylococcus auricularis		92.00	0.55	MNS 10	1	40		
Staphylococcus cohnii subsp. cohnii		5.63	0.34	LAC 10	99	1		
Aerococcus viridans		0.93	0.30		1	95		
002450	**Low discrimination**							**002450**
					OXI	NOV	YEP	
Macrococcus bovicus		74.05	0.80	NIT 10	99	99	99	
Staphylococcus cohnii subsp. cohnii		22.60	0.76		1	99	12	
Staphylococcus auricularis		2.60	0.53		1	1	1	
002451	**Very good identification**							**002451**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		95.52	0.77	NIT 7	1	99	12	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		2.92	0.39		99	99	99	
Staphylococcus auricularis		1.13	0.34		1	1	1	
002452	**Low discrimination**							**002452**
					OXI	NOV	YEP	
Macrococcus bovicus		54.11	0.39	NIT 10	99	99	99	
Staphylococcus auricularis		20.87	0.34	LAC 1	1	1	1	
Staphylococcus cohnii subsp. cohnii		16.51	0.35		1	99	12	
002453	**Low discrimination**							**002453**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		62.13	0.36	NIT 7	99	1		
Aerococcus viridans		24.25	0.39	LAC 1	1	95		
Staphylococcus auricularis		8.05	0.14		1	40		
002460	**Low discrimination**							**002460**
					NOV			
Staphylococcus auricularis		80.22	0.53	XYL 1	1			
Kocuria varians		12.47	0.38		1			
Staphylococcus cohnii subsp. cohnii		5.54	0.33		99			
002461	**Low discrimination**							**002461**
					NOV			
Staphylococcus auricularis		55.70	0.34	XYL 1	1			
Staphylococcus cohnii subsp. cohnii		37.52	0.34	MNS 10	99			
Kocuria varians		6.77	0.16		1			
002462	**Low discrimination**							**002462**
					PYR			
Staphylococcus auricularis		69.34	0.34	XYL 1	40			
Kocuria varians		27.35	0.27	LAC 10	10			
Aerococcus viridans		3.31	0.22		95			
002463	**Low discrimination**							**002463**
					PYR			
Aerococcus viridans		43.41	0.30	ESL 90	95			
Staphylococcus auricularis		43.25	0.14	NIT 1	40			
Kocuria varians		13.35	0.05		10			
002470	**Low discrimination**							**002470**
					OXI	YEP		
Macrococcus bovicus		72.59	0.39	NIT 10	99	99		
Staphylococcus cohnii subsp. cohnii		22.15	0.35	XYL 1	1	12		
Staphylococcus vitulinus		2.71	0.04		99	75		
002471	**Low discrimination**							**002471**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		89.12	0.36	NIT 7	1	99	1	
Staphylococcus sciuri		4.30	0.05	XYL 1	99	99	1	
Aerococcus viridans		3.86	0.20		1	1	95	
002472	**Low discrimination**							**002472**
					OXI	NOV		
Aerococcus viridans		84.55	0.32	ESL 90	1	1		
Macrococcus bovicus		15.45	-.02	NIT 1	99	99		
002473	**Good identification**							**002473**
Aerococcus viridans		100.0	0.39	ESL 90				
002500	**Low discrimination**							**002500**
					OXI			
Staphylococcus auricularis		52.41	0.67	GAL 10	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		33.60	0.65		13			
Dermacoccus nishinomiyaensis		4.58	0.39		93			
002501	**Low discrimination**							**002501**
					NOV			
Staphylococcus auricularis		48.85	0.48	GAL 10	1			
Kocuria varians		24.50	0.43	MNS 10	1			
Staphylococcus cohnii subsp. cohnii		24.24	0.45		99			
002502	**Low discrimination**							**002502**
					NOV	OXI		
Kocuria varians		58.87	0.53	TRE 7	1	13		
Staphylococcus auricularis		36.18	0.48		1	1		
Macrococcus caseolyticus		4.06	0.18		99	99		
002503	**Low discrimination**							**002503**
Kocuria varians		52.50	0.31	TRE 7				
Staphylococcus auricularis		41.23	0.28	MNS 8				
Staphylococcus hyicus		2.21	-.07					
002510	**Low discrimination**							**002510**
					OXI	YEP		
Macrococcus bovicus		89.04	0.65	NIT 10	99	99		
Staphylococcus cohnii subsp. cohnii		5.19	0.46		1	12		
Macrococcus carouselicus		3.70	0.33		99	67		
002511	**Low discrimination**							**002511**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		81.71	0.47	NIT 7	1	99	12	
Macrococcus bovicus		13.07	0.24	GAL 10	99	99	99	
Staphylococcus capitis subsp. capitis		2.49	0.07		1	1	1	
002513	**Low discrimination**							**002513**
					OXI	NOV	PYR	
Aerococcus viridans		44.72	0.27	ESL 90	1	1	95	
Staphylococcus lentus		39.20	0.12	NIT 1	99	99	6	
Staphylococcus cohnii subsp. cohnii		16.08	0.06		1	99	1	
002520	**Good identification**							**002520**
Kocuria varians		93.46	0.55	TRE 7				
Staphylococcus auricularis		4.19	0.26					
Kocuria rosea		1.69	0.10					
002521	**Acceptable identification**							**002521**
					NOV			
Kocuria varians		92.10	0.33	TRE 7	1			
Staphylococcus auricularis		5.28	0.07	MNS 8	1			
Staphylococcus cohnii subsp. cohnii		2.62	0.04		99			
002522	**Good identification**							**002522**
Kocuria varians		98.26	0.44	TRE 7				
Staphylococcus auricularis		1.74	0.07					
002533	**Low discrimination**							**002533**
					OXI	NOV		
Aerococcus viridans		53.29	0.27	ESL 90	1	1		
Staphylococcus lentus		46.71	0.12	NIT 1	99	99		
002540	**Low discrimination**							**002540**
					OXI	NOV	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		66.69	0.75	GAL 10	1	1	1	
Macrococcus caseolyticus		23.55	0.55		99	99	99	
Staphylococcus cohnii subsp. cohnii		4.61	0.54		1	99	12	
002541	**Low discrimination**							**002541**
					NOV			
Staphylococcus auricularis		55.85	0.55	GAL 10	1			
Staphylococcus cohnii subsp. cohnii		37.62	0.55	MNS 10	99			
Kocuria varians		2.64	0.29		1			
002542	**Low discrimination**							**002542**
					NOV	OXI		
Macrococcus caseolyticus		80.03	0.59	VPT 99	99	99		
Staphylococcus auricularis		16.79	0.55		1	1		
Kocuria varians		2.57	0.40		1	13		
002543	**Low discrimination**							**002543**
					NOV	OXI		
Staphylococcus auricularis		57.95	0.36	GAL 10	1	1		
Macrococcus caseolyticus		25.11	0.18	MNS 10	99	99		
Kocuria varians		6.94	0.18	LAC 10	1	13		
002550	**Low discrimination**							**002550**
					OXI	PYR	YEP	
Macrococcus bovicus		88.60	0.73	NIT 10	99	1	99	
Staphylococcus cohnii subsp. cohnii		7.01	0.57		1	1	12	
Staphylococcus saprophyticus subsp. bovis		2.61	0.43		1	99	80	
002551	**Low discrimination**							**002551**
					OXI	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		80.95	0.57	NIT 7	1	1	12	
Macrococcus bovicus		9.54	0.32	GAL 10	99	1	99	
Staphylococcus saprophyticus subsp. bovis		6.96	0.31		1	99	80	
002552	**Low discrimination**							**002552**
					OXI	PYR		
Macrococcus bovicus		38.44	0.32	NIT 10	99	1		
Staphylococcus saprophyticus subsp. bovis		28.05	0.31	LAC 1	1	99		
Macrococcus caseolyticus		18.32	0.18		99	80		
002553	**Low discrimination**							**002553**
					OXI	NOV	PYR	
Aerococcus viridans		42.81	0.39	ESL 90	1	1	95	
Staphylococcus saprophyticus subsp. bovis		25.95	0.18	NIT 1	1	99	99	
Staphylococcus lentus		19.05	0.18		99	99	6	
002560	**Low discrimination**							**002560**
					NOV	OXI		
Kocuria varians		59.60	0.42	TRE 7	1	13		
Staphylococcus auricularis		28.41	0.34		1	1		
Macrococcus caseolyticus		10.03	0.14		99	99		
002562	**Low discrimination**							**002562**
					NOV	OXI		
Kocuria varians		46.22	0.30	TRE 7	1	13		
Macrococcus caseolyticus		41.37	0.18		99	99		
Staphylococcus auricularis		8.68	0.14		1	1		
002563	**Low discrimination**							**002563**
					PYR			
Aerococcus viridans		68.43	0.30	ESL 90	95			
Kocuria varians		31.57	0.08	NIT 1	10			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
002570	**Acceptable identification**							**002570**
					OXI	PYR	YEP	
Macrococcus bovicus		90.20	0.32	NIT 10	99	1	99	
Staphylococcus cohnii subsp. cohnii		7.14	0.15	XYL 1	1	1	12	
Staphylococcus saprophyticus subsp. bovis		2.66	0.02		1	99	80	
002572	**Good identification**							**002572**
Aerococcus viridans		100.0	0.32	ESL 90				
002573	**Low discrimination**							**002573**
					OXI	NOV		
Aerococcus viridans		69.21	0.39	ESL 90	1	1		
Staphylococcus lentus		30.79	0.18	NIT 1	99	99		
002600	**Good identification**							**002600**
					NOV	OXI		
Staphylococcus auricularis		92.00	0.86		1	1		
Macrococcus carouselicus		3.79	0.51		99	99		
Kocuria rosea		1.15	0.39		1	35		
002601	**Low discrimination**							**002601**
					OXI	PYR		
Staphylococcus auricularis		87.67	0.67	MNS 10	1	40		
Kocuria kristinae		7.24	0.38		99	93		
Staphylococcus capitis subsp. capitis		2.81	0.31		1	1		
002602	**Very good identification**							**002602**
					NOV	OXI		
Staphylococcus auricularis		97.07	0.67	LAC 10	1	1		
Kocuria varians		1.73	0.32		1	13		
Macrococcus carouselicus		0.36	0.10		99	99		
002603	**Low discrimination**							**002603**
					OXI	PYR		
Staphylococcus auricularis		78.00	0.47	MNS 10	1	40		
Kocuria kristinae		14.48	0.25	LAC 10	99	93		
Staphylococcus hyicus		4.34	0.13		1	1		
002610	**Low discrimination**							**002610**
Staphylococcus vitulinus		89.39	0.77					
Macrococcus bovicus		7.00	0.60					
Macrococcus carouselicus		2.36	0.47					
002611	**Low discrimination**							**002611**
					OXI	NOV		
Staphylococcus capitis subsp. capitis		72.93	0.54	TRE 2	1	1		
Staphylococcus vitulinus		15.52	0.36		99	99		
Staphylococcus sciuri		6.31	0.36		99	99		
002612	**Low discrimination**							**002612**
					OXI	NOV		
Staphylococcus vitulinus		78.54	0.36	LAC 1	99	99		
Staphylococcus auricularis		8.76	0.26		1	1		
Macrococcus bovicus		6.15	0.19		99	99		
002620	**Low discrimination**							**002620**
					OXI	NOV		
Staphylococcus muscae		88.98	0.59	PHS 99	1	1		
Staphylococcus vitulinus		6.58	0.45		99	99		
Staphylococcus auricularis		2.44	0.45		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
002630	**Very good identification**							**002630**
					OXI	NOV		
Staphylococcus vitulinus		99.78	0.86		99	99		
Staphylococcus muscae		0.14	0.18		1	1		
Macrococcus bovicus		0.03	0.19		99	99		
002631	**Low discrimination**							**002631**
					OXI	NOV		
Staphylococcus vitulinus		88.42	0.45	MNS 1	99	99		
Staphylococcus sciuri		10.03	0.34		99	99		
Staphylococcus capitis subsp. capitis		1.55	0.13		1	1		
002632	**Good identification**							**002632**
					OXI	NOV		
Staphylococcus vitulinus		99.63	0.45	LAC 1	99	99		
Aerococcus viridans		0.37	0.19		1	1		
002640	**Low discrimination**							**002640**
Staphylococcus auricularis		89.40	0.94					
Staphylococcus petrasii subsp. pragensis		7.12	0.54					
Staphylococcus haemolyticus		2.01	0.61					
002641	**Low discrimination**							**002641**
					OXI	NOV		
Staphylococcus auricularis		74.24	0.74	MNS 10	1	1		
Kocuria kristinae		23.63	0.57		99	1		
Staphylococcus cohnii subsp. cohnii		0.55	0.34		1	99		
002642	**Low discrimination**							**002642**
					PYR			
Staphylococcus auricularis		80.05	0.74	LAC 10	40			
Staphylococcus hominis subsp. hominis		9.32	0.58		1			
Staphylococcus haemolyticus		9.11	0.56		98			
002643	**Low discrimination**							**002643**
					OXI			
Staphylococcus auricularis		55.37	0.55	MNS 10	1			
Kocuria kristinae		39.66	0.45	LAC 10	99			
Staphylococcus haemolyticus		1.75	0.25		1			
002650	**Low discrimination**							**002650**
					OXI	NOV		
Macrococcus bovicus		67.99	0.68	NIT 10	99	99		
Staphylococcus haemolyticus		11.50	0.56		1	2		
Staphylococcus auricularis		8.80	0.53		1	1		
002651	**Low discrimination**							**002651**
					OXI	NOV	PYR	
Staphylococcus sciuri		51.56	0.48	ESL 99	99	99	1	
Kocuria kristinae		11.52	0.29		99	1	93	
Staphylococcus auricularis		8.77	0.33		1	1	40	
002652	**Low discrimination**							**002652**
					PYR			
Staphylococcus haemolyticus		57.70	0.51	VPT 86	98			
Staphylococcus hominis subsp. hominis		15.03	0.41	ARG 92	1			
Staphylococcus auricularis		8.72	0.33		40			
002653	**Low discrimination**							**002653**
					OXI	NOV	PYR	
Staphylococcus sciuri		38.23	0.32	ESL 99	99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		14.48	0.39	LAC 14	1	1	95	
Kocuria kristinae		13.11	0.16		99	1	93	
002660	**Low discrimination**							**002660**
					PYR			
Staphylococcus auricularis		76.59	0.53	XYL 1	40			
Staphylococcus muscae		12.09	0.18		10			
Staphylococcus petrasii subsp. pragensis		6.10	0.13		99			
002661	**Low discrimination**							**002661**
					OXI	NOV	PYR	
Staphylococcus auricularis		68.71	0.33	XYL 1	1	1	40	
Kocuria kristinae		21.87	0.16	MNS 10	99	1	93	
Staphylococcus sciuri		9.42	0.15		99	99	1	
002662	**Low discrimination**							**002662**
					PYR			
Staphylococcus auricularis		74.61	0.33	XYL 1	40			
Staphylococcus hominis subsp. hominis		8.69	0.16	LAC 10	1			
Staphylococcus haemolyticus		8.49	0.15		98			
002663	**Low discrimination**							**002663**
					OXI			
Aerococcus viridans		38.78	0.30	ESL 90	1			
Staphylococcus auricularis		35.67	0.13	NIT 1	1			
Kocuria kristinae		25.55	0.04		99			
002670	**Low discrimination**							**002670**
Staphylococcus vitulinus		87.42	0.45	MLT 1				
Macrococcus bovicus		5.91	0.27					
Staphylococcus sciuri		4.61	0.27					
002671	**Good identification**							**002671**
Staphylococcus sciuri		95.88	0.46	ESL 99				
Staphylococcus vitulinus		2.27	0.04					
Staphylococcus lentus		0.98	0.07					
002672	**Low discrimination**							**002672**
					OXI	NOV		
Aerococcus viridans		31.00	0.32	ESL 90	1	1		
Staphylococcus vitulinus		20.95	0.04	NIT 1	99	99		
Staphylococcus sciuri		17.81	0.11		99	99		
002673	**Low discrimination**							**002673**
					OXI	NOV		
Staphylococcus sciuri		51.60	0.30	ESL 99	99	99		
Aerococcus viridans		25.90	0.39	LAC 14	1	1		
Staphylococcus lentus		11.53	0.18		99	99		
002700	**Low discrimination**							**002700**
					OXI	NOV		
Staphylococcus auricularis		71.75	0.67	GAL 10	1	1		
Macrococcus carouselicus		11.41	0.43		99	99		
Rothia mucilaginosa		8.17	0.37		1	1		
002701	**Low discrimination**							**002701**
					OXI			
Rothia mucilaginosa		68.44	0.47	VPT 95	1			
Staphylococcus auricularis		22.27	0.47	ESL 90	1			
Kocuria kristinae		5.51	0.28		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
002702	**Low discrimination**						**002702**
					NOV	OXI	
Staphylococcus auricularis		70.28	0.47	GAL 10	1	1	
Kocuria varians		16.89	0.35	LAC 10	1	13	
Macrococcus caseolyticus		5.70	0.14		99	99	
002703	**Low discrimination**						**002703**
					OXI	PYR	
Staphylococcus hyicus		71.45	0.34	ARG 96	1	1	
Staphylococcus auricularis		12.42	0.27	PHS 92	1	40	
Kocuria kristinae		6.92	0.15		99	93	
002710	**Low discrimination**						**002710**
Macrococcus bovicus		57.65	0.53	NIT 10			
Macrococcus carouselicus		19.47	0.39				
Staphylococcus vitulinus		17.35	0.36				
002711	**Low discrimination**						**002711**
					OXI	NOV	
Staphylococcus sciuri		50.93	0.42	ESL 99	99	99	
Staphylococcus lentus		37.24	0.41		99	99	
Staphylococcus capitis subsp. capitis		3.06	0.13		1	1	
002713	**Acceptable identification**						**002713**
					OXI	NOV	
Staphylococcus lentus		92.11	0.53	ESL 99	99	99	
Staphylococcus sciuri		5.78	0.26		99	99	
Aerococcus viridans		1.06	0.27		1	1	
002730	**Good identification**						**002730**
Staphylococcus vitulinus		95.80	0.45	GAL 1			
Staphylococcus sciuri		2.61	0.21				
Macrococcus bovicus		1.19	0.12				
002731	**Low discrimination**						**002731**
Staphylococcus sciuri		49.63	0.40	ESL 99			
Staphylococcus lentus		48.10	0.41				
Staphylococcus vitulinus		2.28	0.04				
002733	**Acceptable identification**						**002733**
					OXI	NOV	
Staphylococcus lentus		94.08	0.53	ESL 99	99	99	
Staphylococcus sciuri		4.46	0.24		99	99	
Aerococcus viridans		1.08	0.27		1	1	
002740	**Low discrimination**						**002740**
					OXI	NOV	
Staphylococcus auricularis		57.75	0.74	GAL 10	1	1	
Macrococcus caseolyticus		14.73	0.52		99	99	
Staphylococcus haemolyticus		13.17	0.62		1	2	
002741	**Low discrimination**						**002741**
					OXI		
Rothia mucilaginosa		59.23	0.54	VPT 95	1		
Staphylococcus auricularis		19.27	0.55	ESL 90	1		
Kocuria kristinae		18.41	0.47		99		
002742	**Low discrimination**						**002742**
					OXI	PYR	NOV

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus caseolyticus		50.67	0.55	VPT 99	99	80	99	
Staphylococcus haemolyticus		17.00	0.57		1	98	2	
Staphylococcus hominis subsp. hominis		16.72	0.58		1	1	1	
002743	**Low discrimination**							**002743**
					OXI			
Kocuria kristinae		44.67	0.35	VPT 90	99			
Staphylococcus auricularis		20.79	0.35	NIT 7	1			
Staphylococcus haemolyticus		6.68	0.26		1			
002750	**Low discrimination**							**002750**
					OXI	NOV	PYR	
Macrococcus bovicus		48.09	0.60	NIT 10	99	99	1	
Staphylococcus saprophyticus subsp. bovis		22.69	0.55		1	99	99	
Staphylococcus haemolyticus		21.40	0.57		1	2	98	
002751	**Low discrimination**							**002751**
					OXI	PYR		
Staphylococcus sciuri		53.81	0.54	ESL 99	99	1		
Staphylococcus lentus		21.23	0.48		99	6		
Staphylococcus saprophyticus subsp. bovis		16.57	0.43		1	99		
002752	**Low discrimination**							**002752**
					PYR	NOV		
Staphylococcus haemolyticus		52.25	0.52	VPT 86	98	2		
Staphylococcus saprophyticus subsp. bovis		24.62	0.43	ARG 92	99	99		
Staphylococcus hominis subsp. hominis		13.08	0.41		1	1		
002753	**Low discrimination**							**002753**
					OXI	PYR		
Staphylococcus lentus		81.59	0.59	ESL 99	99	6		
Staphylococcus sciuri		9.49	0.38		99	1		
Staphylococcus saprophyticus subsp. bovis		4.49	0.31		1	99		
002760	**Low discrimination**							**002760**
					NOV	OXI		
Staphylococcus auricularis		48.85	0.33	GAL 10	1	1		
Kocuria varians		15.14	0.24	XYL 1	1	13		
Macrococcus caseolyticus		12.46	0.11		99	99		
002770	**Low discrimination**							**002770**
					OXI	PYR		
Staphylococcus sciuri		59.40	0.33	ESL 99	99	1		
Macrococcus bovicus		16.80	0.19	MNS 89	99	1		
Staphylococcus saprophyticus subsp. bovis		7.93	0.14		1	99		
002771	**Low discrimination**							**002771**
					OXI			
Staphylococcus sciuri		64.57	0.52	ESL 99	99			
Staphylococcus lentus		33.77	0.48		99			
Staphylococcus xylosus		1.10	0.11		1			
002772	**Low discrimination**							**002772**
					OXI	YEP		
Staphylococcus arlettae		44.43	0.31	GLR 92	1	99		
Staphylococcus sciuri		16.03	0.17	NIT 1	99	57		
Staphylococcus lentus		14.92	0.18		99	1		
002773	**Low discrimination**							**002773**
					OXI	YEP		
Staphylococcus lentus		85.01	0.59	ESL 99	99	1		
Staphylococcus sciuri		7.46	0.35		99	57		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		3.39	0.34		1	99	
003000	**Low discrimination**						**003000**
					NOV		
Kocuria rosea		53.20	0.59	ESL 1	1		
Dermacoccus nishinomiyaensis		26.00	0.54		80		
Macrococcus carouselicus		9.57	0.45		99		
003001	**Low discrimination**						**003001**
Kocuria rosea		68.65	0.33	ESL 1			
Staphylococcus capitis subsp. capitis		10.96	0.19	MNS 5			
Kocuria varians		6.94	0.21				
003002	**Low discrimination**						**003002**
					OXI		
Kocuria varians		47.46	0.32	ESL 1	13		
Kocuria rosea		27.80	0.18		35		
Dermacoccus nishinomiyaensis		13.59	0.13		93		
003010	**Low discrimination**						**003010**
Staphylococcus vitulinus		47.76	0.50	SUC 99			
Macrococcus bovicus		22.82	0.50				
Macrococcus carouselicus		12.81	0.41				
003011	**Low discrimination**						**003011**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		88.49	0.42	ESL 1	1	1	
Kocuria rosea		4.63	0.13		1	35	
Staphylococcus vitulinus		3.43	0.09		99	99	
003020	**Acceptable identification**						**003020**
					NOV	OXI	
Kocuria rosea		91.22	0.51	ESL 1	1	35	
Kocuria varians		4.57	0.34		1	13	
Staphylococcus vitulinus		2.49	0.18		99	99	
003030	**Very good identification**						**003030**
					NOV	YEP	
Staphylococcus vitulinus		95.80	0.59	SUC 99	99	75	
Kocuria rosea		3.94	0.32		1	1	
Macrococcus bovicus		0.17	0.08		99	99	
003040	**Low discrimination**						**003040**
Staphylococcus auricularis		28.62	0.32	ESL 1			
Kocuria varians		19.53	0.29	TRE 92			
Kocuria rosea		14.70	0.18				
003050	**Low discrimination**						**003050**
Macrococcus bovicus		84.52	0.57	ESL 10			
Macrococcus equipersicus		14.31	0.43	NIT 10			
Staphylococcus vitulinus		0.77	0.09				
003053	**Low discrimination**						**003053**
					OXI	NOV	
Aerococcus viridans		85.29	0.33	NIT 1	1	1	
Staphylococcus sciuri		14.71	0.01	TRE 95	99	99	
003073	**Good identification**						**003073**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.33	NIT 1			
003100	**Low discrimination**						**003100**
					OXI	NOV	
Dermacoccus nishinomiyaensis		33.75	0.39	ESL 1	93	80	
Kocuria varians		33.26	0.47		13	1	
Macrococcus carouselicus		27.95	0.37		99	99	
003102	**Acceptable identification**						**003102**
					OXI	NOV	
Kocuria varians		93.77	0.35	ESL 1	13	1	
Dermacoccus nishinomiyaensis		3.41	-.02		93	80	
Macrococcus carouselicus		2.82	-.04		99	99	
003110	**Low discrimination**						**003110**
Macrococcus bovicus		55.84	0.42	ESL 10			
Macrococcus carouselicus		31.35	0.33	NIT 10			
Macrococcus equipersicus		8.16	0.26				
003120	**Low discrimination**						**003120**
					OXI		
Kocuria varians		84.20	0.37	ESL 1	13		
Kocuria rosea		11.31	0.10		35		
Dermacoccus nishinomiyaensis		2.46	-.02		93		
003140	**Low discrimination**						**003140**
					NOV	OXI	
Kocuria varians		72.08	0.33	ESL 1	1	13	
Staphylococcus auricularis		7.82	0.12		1	1	
Macrococcus bovicus		7.41	0.08		99	99	
003150	**Low discrimination**						**003150**
Macrococcus bovicus		85.16	0.50	ESL 10			
Macrococcus equipersicus		14.42	0.35	NIT 10			
Staphylococcus sciuri		0.42	0.04				
003153	**Low discrimination**						**003153**
					OXI	NOV	
Aerococcus viridans		43.98	0.33	NIT 1	1	1	
Staphylococcus lentus		41.31	0.18	TRE 95	99	99	
Staphylococcus sciuri		14.72	0.07		99	99	
003173	**Low discrimination**						**003173**
					OXI	NOV	
Aerococcus viridans		45.62	0.33	NIT 1	1	1	
Staphylococcus lentus		42.86	0.18	TRE 95	99	99	
Staphylococcus sciuri		11.52	0.04		99	99	
003200	**Low discrimination**						**003200**
					NOV		
Staphylococcus vitulinus		44.42	0.50	MAN 99	99		
Macrococcus carouselicus		37.84	0.51		99		
Kocuria rosea		11.51	0.39		1		
003210	**Very good identification**						**003210**
Staphylococcus vitulinus		99.26	0.91				
Macrococcus carouselicus		0.55	0.47				
Macrococcus bovicus		0.12	0.37				
003211	**Low discrimination**						**003211**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Staphylococcus vitulinus		57.15	0.50	MNS 1	99	99		
Staphylococcus capitis subsp. capitis		27.65	0.47		1	1		
Staphylococcus sciuri		14.80	0.46		99	99		
003212	**Good identification**							**003212**
Staphylococcus vitulinus		98.94	0.50	LAC 1				
Macrococcus carouselicus		0.54	0.06					
Staphylococcus sciuri		0.52	0.12					
003213	**Low discrimination**							**003213**
					OXI	NOV		
Staphylococcus sciuri		61.32	0.30	TRE 97	99	99		
Staphylococcus vitulinus		14.69	0.09	LAC 14	99	99		
Staphylococcus capitis subsp. capitis		14.36	0.13		1	1		
003220	**Very good identification**							**003220**
					NOV			
Staphylococcus vitulinus		95.53	0.59	MAN 99	99			
Kocuria rosea		3.93	0.32		1			
Macrococcus carouselicus		0.30	0.10		99			
003230	**Excellent identification**							**003230**
					NOV			
Staphylococcus vitulinus		99.98	1.00		99			
Staphylococcus sciuri		0.01	0.25		99			
Kocuria rosea		0.01	0.12		1			
003231	**Good identification**							**003231**
					OXI	NOV		
Staphylococcus vitulinus		93.11	0.59	MNS 1	99	99		
Staphylococcus sciuri		6.73	0.44		99	99		
Staphylococcus capitis subsp. capitis		0.17	0.06		1	1		
003232	**Very good identification**							**003232**
Staphylococcus vitulinus		99.86	0.59	LAC 1				
Staphylococcus sciuri		0.15	0.09					
003233	**Low discrimination**							**003233**
Staphylococcus sciuri		48.38	0.28	TRE 97				
Staphylococcus vitulinus		41.55	0.18	LAC 14				
Staphylococcus lentus		6.54	0.12					
003240	**Low discrimination**							**003240**
					OXI	NOV		
Staphylococcus auricularis		48.83	0.31	ESL 1	1	1		
Staphylococcus vitulinus		11.65	0.09	TRE 92	99	99		
Macrococcus carouselicus		9.92	0.10		99	99		
003241	**Low discrimination**							**003241**
					OXI	NOV	PYR	
Staphylococcus sciuri		49.36	0.27	TRE 97	99	99	1	
Kocuria kristinae		36.77	0.19	MAN 97	99	1	93	
Staphylococcus auricularis		7.71	0.11		1	1	40	
003250	**Low discrimination**							**003250**
Staphylococcus vitulinus		70.26	0.50	MLT 1				
Macrococcus bovicus		19.58	0.45					

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		8.46	0.40				
003251	**Very good identification**						**003251**
					OXI	NOV	
Staphylococcus sciuri		97.96	0.58	TRE 97	99	99	
Staphylococcus vitulinus		1.02	0.09		99	99	
Staphylococcus capitis subsp. capitis		0.49	0.06		1	1	
003253	**Low discrimination**						**003253**
					OXI	NOV	
Staphylococcus sciuri		89.80	0.42	TRE 97	99	99	
Aerococcus viridans		5.26	0.33	LAC 14	1	1	
Staphylococcus lentus		4.94	0.18		99	99	
003270	**Very good identification**						**003270**
Staphylococcus vitulinus		96.65	0.59	MLT 1			
Staphylococcus sciuri		3.25	0.37				
Macrococcus bovicus		0.10	0.04				
003271	**Very good identification**						**003271**
Staphylococcus sciuri		96.11	0.56	TRE 97			
Staphylococcus vitulinus		3.57	0.18				
Staphylococcus lentus		0.32	0.07				
003273	**Low discrimination**						**003273**
					OXI	NOV	
Staphylococcus sciuri		86.91	0.40	TRE 97	99	99	
Aerococcus viridans		6.75	0.33	LAC 14	1	1	
Staphylococcus lentus		6.34	0.18		99	99	
003300	**Low discrimination**						**003300**
					NOV	OXI	
Macrococcus carouselicus		81.40	0.43	NIT 6	99	99	
Kocuria varians		6.51	0.29	TRE 94	1	13	
Rothia mucilaginosa		5.30	0.15		1	1	
003310	**Low discrimination**						**003310**
Staphylococcus vitulinus		74.09	0.50	GAL 1			
Macrococcus carouselicus		17.30	0.39				
Staphylococcus sciuri		4.60	0.34				
003311	**Low discrimination**						**003311**
Staphylococcus sciuri		78.54	0.52	TRE 97			
Staphylococcus lentus		18.75	0.41				
Staphylococcus vitulinus		1.58	0.09				
003313	**Low discrimination**						**003313**
					OXI	NOV	
Staphylococcus lentus		83.49	0.53	TRE 99	99	99	
Staphylococcus sciuri		16.05	0.36		99	99	
Aerococcus viridans		0.46	0.20		1	1	
003330	**Very good identification**						**003330**
Staphylococcus vitulinus		98.21	0.59	GAL 1			
Staphylococcus sciuri		1.70	0.31				
Macrococcus carouselicus		0.09	-.02				
003331	**Low discrimination**						**003331**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		72.02	0.50	TRE 97				
Staphylococcus lentus		22.80	0.41					
Staphylococcus vitulinus		5.19	0.18					
003333	**Low discrimination**							**003333**
					OXI	NOV		
Staphylococcus lentus		86.92	0.53	TRE 99	99	99		
Staphylococcus sciuri		12.61	0.34		99	99		
Aerococcus viridans		0.47	0.20		1	1		
003341	**Low discrimination**							**003341**
					OXI	NOV	PYR	
Rothia mucilaginosa		69.02	0.33	VPT 95	1	1	75	
Staphylococcus sciuri		24.27	0.33	TRE 99	99	99	1	
Kocuria kristinae		3.10	0.09		99	1	93	
003350	**Low discrimination**							**003350**
Staphylococcus sciuri		62.14	0.46	TRE 97				
Macrococcus bovicus		31.76	0.37	MNS 89				
Staphylococcus vitulinus		2.69	0.09					
003351	**Low discrimination**							**003351**
					OXI	NOV		
Staphylococcus sciuri		88.51	0.64	TRE 97	99	99		
Staphylococcus lentus		11.41	0.48		99	99		
Rothia mucilaginosa		0.08	-.09		1	1		
003352	**Low discrimination**							**003352**
					OXI	NOV		
Staphylococcus sciuri		74.96	0.29	TRE 97	99	99		
Staphylococcus lentus		17.20	0.18	MNS 89	99	99		
Aerococcus viridans		7.85	0.25	LAC 14	1	1		
003353	**Low discrimination**							**003353**
					OXI	NOV		
Staphylococcus lentus		73.15	0.59	TRE 99	99	99		
Staphylococcus sciuri		26.06	0.48		99	99		
Aerococcus viridans		0.79	0.33		1	1		
003361	**Acceptable identification**							**003361**
					OXI	NOV		
Staphylococcus sciuri		91.42	0.31	TRE 97	99	99		
Staphylococcus lentus		5.10	0.07	MAN 97	99	99		
Rothia mucilaginosa		3.48	-.09		1	1		
003370	**Low discrimination**							**003370**
Staphylococcus sciuri		85.55	0.43	TRE 97				
Staphylococcus vitulinus		13.25	0.18	MNS 89				
Staphylococcus lentus		1.19	0.07					
003371	**Low discrimination**							**003371**
Staphylococcus sciuri		85.41	0.62	TRE 97				
Staphylococcus lentus		14.59	0.48					
003372	**Low discrimination**							**003372**
					OXI	NOV		
Staphylococcus sciuri		69.31	0.27	TRE 97	99	99		
Staphylococcus lentus		21.08	0.18	MNS 89	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		9.62	0.25	LAC 14	1	1	
003373	**Low discrimination**						**003373**
					OXI	NOV	
Staphylococcus lentus		78.15	0.59	TRE 99	99	99	
Staphylococcus sciuri		21.01	0.45		99	99	
Aerococcus viridans		0.84	0.33		1	1	
003400	**Good identification**						**003400**
					OXI	NOV	
Macrococcus carouselicus		94.61	0.69	NIT 6	99	99	
Staphylococcus auricularis		3.25	0.46		1	1	
Staphylococcus carnosus subsp. utilis		0.88	0.18		1	1	
003401	**Low discrimination**						**003401**
					OXI	NOV	
Macrococcus carouselicus		61.34	0.28	NIT 6	99	99	
Staphylococcus auricularis		23.21	0.26	MNS 1	1	1	
Staphylococcus cohnii subsp. cohnii		11.52	0.23		1	99	
003402	**Low discrimination**						**003402**
					OXI	NOV	
Macrococcus carouselicus		56.60	0.28	NIT 6	99	99	
Staphylococcus auricularis		21.42	0.26	LAC 1	1	1	
Staphylococcus piscifermentans		14.03	0.32		1	1	
003403	**Low discrimination**						**003403**
Aerococcus viridans		84.05	0.37	NIT 1			
Staphylococcus auricularis		15.95	0.07				
003410	**Low discrimination**						**003410**
Macrococcus carouselicus		75.20	0.65	NIT 6			
Macrococcus bovicus		12.83	0.53				
Staphylococcus vitulinus		10.07	0.45				
003411	**Low discrimination**						**003411**
					OXI		
Staphylococcus sciuri		50.94	0.36	SUC 99	99		
Macrococcus carouselicus		25.57	0.24		99		
Staphylococcus cohnii subsp. cohnii		9.56	0.26		1		
003413	**Low discrimination**						**003413**
					OXI	NOV	
Aerococcus viridans		70.95	0.47	NIT 1	1	1	
Staphylococcus sciuri		22.14	0.20		99	99	
Staphylococcus lentus		6.91	0.12		99	99	
003420	**Low discrimination**						**003420**
					NOV		
Macrococcus carouselicus		67.81	0.28	NIT 6	99		
Staphylococcus vitulinus		15.69	0.13	XYL 1	99		
Kocuria rosea		10.31	0.10		1		
003422	**Good identification**						**003422**
Aerococcus viridans		100.0	0.29	NIT 1			
003423	**Good identification**						**003423**
Aerococcus viridans		100.0	0.37	NIT 1			
003430	**Very good identification**						**003430**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus vitulinus		96.22	0.54	SUC 99				
Macrococcus carouselicus		2.69	0.24					
Staphylococcus sciuri		0.50	0.15					
003431	**Low discrimination**							**003431**
					OXI	NOV		
Staphylococcus sciuri		75.89	0.34	SUC 99	99	99		
Staphylococcus vitulinus		18.27	0.13		99	99		
Aerococcus viridans		5.83	0.27		1	1		
003432	**Low discrimination**							**003432**
					OXI	NOV		
Aerococcus viridans		55.18	0.39	NIT 1	1	1		
Staphylococcus vitulinus		44.82	0.13		99	99		
003433	**Low discrimination**							**003433**
					OXI	NOV		
Aerococcus viridans		75.03	0.47	NIT 1	1	1		
Staphylococcus sciuri		17.66	0.18		99	99		
Staphylococcus lentus		7.31	0.12		99	99		
003440	**Low discrimination**							**003440**
					OXI	NOV		
Staphylococcus auricularis		78.35	0.53	ESL 1	1	1		
Macrococcus carouselicus		9.86	0.28		99	99		
Staphylococcus cohnii subsp. cohnii		5.41	0.33		1	99		
003441	**Low discrimination**							**003441**
					OXI	NOV		
Staphylococcus auricularis		48.66	0.34	ESL 1	1	1		
Staphylococcus cohnii subsp. cohnii		32.78	0.34	MNS 10	1	99		
Staphylococcus sciuri		8.17	0.17		99	99		
003442	**Low discrimination**							**003442**
Staphylococcus auricularis		58.45	0.34	ESL 1				
Aerococcus viridans		25.14	0.42	LAC 10				
Staphylococcus piscifermentans		16.41	0.32					
003443	**Low discrimination**							**003443**
					OXI	NOV		
Aerococcus viridans		87.88	0.49	NIT 1	1	1		
Staphylococcus auricularis		9.73	0.14		1	1		
Staphylococcus sciuri		2.39	0.01		99	99		
003450	**Low discrimination**							**003450**
					OXI	YEP		
Macrococcus bovicus		83.06	0.61	ESL 10	99	99		
Macrococcus equipersicus		9.38	0.43	NIT 10	99	99		
Staphylococcus cohnii subsp. cohnii		2.30	0.35		1	12		
003451	**Low discrimination**							**003451**
					OXI	NOV	PYR	
Staphylococcus sciuri		77.73	0.48	SUC 99	99	99	1	
Staphylococcus cohnii subsp. cohnii		12.28	0.36		1	99	1	
Aerococcus viridans		4.79	0.39		1	1	95	
003452	**Low discrimination**							**003452**
					OXI	NOV		
Aerococcus viridans		70.11	0.51	NIT 1	1	1		
Macrococcus bovicus		15.66	0.20		99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		6.54	0.25		1	1	
003453	**Low discrimination**						**003453**
					OXI	NOV	
Aerococcus viridans		74.47	0.59	NIT 1	1	1	
Staphylococcus sciuri		21.85	0.32		99	99	
Staphylococcus lentus		3.68	0.18		99	99	
003462	**Good identification**						**003462**
Aerococcus viridans		100.0	0.42	NIT 1			
003463	**Good identification**						**003463**
Aerococcus viridans		100.0	0.49	NIT 1			
003470	**Low discrimination**						**003470**
Staphylococcus sciuri		41.07	0.27	SUC 99			
Macrococcus bovicus		23.39	0.20	MNS 89			
Staphylococcus vitulinus		21.27	0.13				
003471	**Acceptable identification**						**003471**
					OXI	NOV	
Staphylococcus sciuri		91.58	0.46	SUC 99	99	99	
Aerococcus viridans		7.48	0.39		1	1	
Staphylococcus lentus		0.94	0.07		99	99	
003472	**Acceptable identification**						**003472**
					OXI	NOV	
Aerococcus viridans		94.00	0.51	NIT 1	1	1	
Staphylococcus sciuri		6.00	0.11		99	99	
003473	**Low discrimination**						**003473**
					OXI	NOV	
Aerococcus viridans		78.69	0.59	NIT 1	1	1	
Staphylococcus sciuri		17.42	0.30		99	99	
Staphylococcus lentus		3.89	0.18		99	99	
003500	**Very good identification**						**003500**
					OXI	NOV	
Macrococcus carouselicus		96.96	0.62	NIT 6	99	99	
Staphylococcus auricularis		0.87	0.26		1	1	
Rothia mucilaginosa		0.82	0.15		1	1	
003503	**Low discrimination**						**003503**
					OXI	NOV	
Aerococcus viridans		77.39	0.37	NIT 1	1	1	
Staphylococcus lentus		22.61	0.12		99	99	
003510	**Low discrimination**						**003510**
Macrococcus carouselicus		82.81	0.58	NIT 6			
Macrococcus bovicus		14.12	0.46				
Macrococcus equipersicus		1.38	0.26				
003511	**Low discrimination**						**003511**
Staphylococcus sciuri		52.59	0.42	SUC 99			
Staphylococcus lentus		38.45	0.41				
Macrococcus carouselicus		5.83	0.16				
003512	**Low discrimination**						**003512**
					OXI	NOV	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		32.39	0.39	NIT 1	1	1	
Macrococcus carouselicus		31.16	0.16		99	99	
Staphylococcus piscifermentans		18.12	0.28		1	1	
003513	**Low discrimination**						**003513**
					OXI	NOV	
Staphylococcus lentus		85.73	0.53	SUC 99	99	99	
Aerococcus viridans		8.89	0.47		1	1	
Staphylococcus sciuri		5.38	0.26		99	99	
003522	**Low discrimination**						**003522**
					PYR		
Aerococcus viridans		79.80	0.29	NIT 1	95		
Kocuria varians		20.20	0.03		10		
003523	**Low discrimination**						**003523**
					OXI	NOV	
Aerococcus viridans		77.39	0.37	NIT 1	1	1	
Staphylococcus lentus		22.61	0.12		99	99	
003531	**Low discrimination**						**003531**
					OXI	NOV	
Staphylococcus sciuri		49.78	0.40	SUC 99	99	99	
Staphylococcus lentus		48.25	0.41		99	99	
Aerococcus viridans		1.97	0.27		1	1	
003532	**Low discrimination**						**003532**
					OXI	NOV	
Aerococcus viridans		73.58	0.39	NIT 1	1	1	
Staphylococcus lentus		16.72	0.12		99	99	
Staphylococcus sciuri		9.69	0.05		99	99	
003533	**Low discrimination**						**003533**
					OXI	NOV	
Staphylococcus lentus		86.88	0.53	SUC 99	99	99	
Aerococcus viridans		9.01	0.47		1	1	
Staphylococcus sciuri		4.12	0.24		99	99	
003540	**Low discrimination**						**003540**
					OXI	NOV	
Staphylococcus auricularis		28.75	0.34	ESL 1	1	1	
Rothia mucilaginosa		27.19	0.23	GAL 10	1	1	
Macrococcus carouselicus		13.96	0.20		99	99	
003541	**Low discrimination**						**003541**
					OXI	NOV	
Rothia mucilaginosa		80.44	0.33	VPT 95	1	1	
Staphylococcus sciuri		9.24	0.23	SUC 99	99	99	
Aerococcus viridans		3.16	0.30		1	1	
003543	**Low discrimination**						**003543**
					OXI	NOV	
Aerococcus viridans		81.32	0.49	NIT 1	1	1	
Staphylococcus lentus		12.06	0.18		99	99	
Staphylococcus sciuri		4.30	0.07		99	99	
003550	**Low discrimination**						**003550**
Macrococcus bovicus		78.02	0.53	ESL 10			
Macrococcus equipersicus		8.81	0.35	NIT 10			
Staphylococcus sciuri		8.31	0.36				
003551	**Low discrimination**						**003551**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Staphylococcus sciuri		68.97	0.54	SUC 99	99	99	
Staphylococcus lentus		27.22	0.48		99	99	
Aerococcus viridans		2.19	0.39		1	1	
003552	**Low discrimination**						**003552**
					OXI	NOV	
Aerococcus viridans		65.10	0.51	NIT 1	1	1	
Staphylococcus sciuri		10.69	0.19		99	99	
Staphylococcus piscifermentans		9.10	0.28		1	1	
003553	**Low discrimination**						**003553**
					OXI	NOV	
Staphylococcus lentus		75.72	0.59	SUC 99	99	99	
Aerococcus viridans		15.47	0.59		1	1	
Staphylococcus sciuri		8.81	0.38		99	99	
003562	**Good identification**						**003562**
Aerococcus viridans		100.0	0.42	NIT 1			
003563	**Low discrimination**						**003563**
					OXI	NOV	
Aerococcus viridans		84.16	0.49	NIT 1	1	1	
Staphylococcus lentus		12.48	0.18		99	99	
Staphylococcus sciuri		3.36	0.04		99	99	
003570	**Low discrimination**						**003570**
					OXI	NOV	
Staphylococcus sciuri		76.08	0.33	SUC 99	99	99	
Aerococcus viridans		11.11	0.32	MNS 89	1	1	
Macrococcus bovicus		9.57	0.12		99	99	
003571	**Low discrimination**						**003571**
					OXI	NOV	
Staphylococcus sciuri		63.89	0.52	SUC 99	99	99	
Staphylococcus lentus		33.42	0.48		99	99	
Aerococcus viridans		2.69	0.39		1	1	
003572	**Low discrimination**						**003572**
					OXI	NOV	
Aerococcus viridans		80.69	0.51	NIT 1	1	1	
Staphylococcus sciuri		10.00	0.17		99	99	
Staphylococcus lentus		9.31	0.18		99	99	
003573	**Low discrimination**						**003573**
					OXI	NOV	
Staphylococcus lentus		77.39	0.59	SUC 99	99	99	
Aerococcus viridans		15.81	0.59		1	1	
Staphylococcus sciuri		6.79	0.35		99	99	
003600	**Good identification**						**003600**
					OXI	NOV	
Macrococcus carouselicus		94.02	0.75	NIT 6	99	99	
Staphylococcus vitulinus		3.96	0.45		99	99	
Staphylococcus auricularis		1.47	0.45		1	1	
003601	**Low discrimination**						**003601**
					OXI	NOV	
Staphylococcus sciuri		47.70	0.46	MAN 97	99	99	
Macrococcus carouselicus		24.83	0.34		99	99	
Rothia mucilaginosa		13.13	0.25		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
003602	**Low discrimination**							**003602**
					OXI	NOV		
Macrococcus carouselicus		67.45	0.34	NIT 6	99	99		
Staphylococcus piscifermentans		12.35	0.36	LAC 1	1	1		
Staphylococcus auricularis		11.60	0.26		1	1		
003603	**Low discrimination**							**003603**
					OXI	NOV	PYR	
Staphylococcus sciuri		55.77	0.30	MAN 97	99	99	1	
Aerococcus viridans		19.46	0.37	LAC 14	1	1	95	
Kocuria kristinae		15.67	0.13		99	1	93	
003610	**Low discrimination**							**003610**
Staphylococcus vitulinus		85.08	0.86					
Macrococcus carouselicus		13.04	0.71					
Staphylococcus sciuri		1.58	0.59					
003611	**Good identification**							**003611**
Staphylococcus sciuri		91.41	0.78					
Staphylococcus vitulinus		6.14	0.45					
Staphylococcus lentus		1.31	0.41					
003612	**Low discrimination**							**003612**
Staphylococcus vitulinus		65.96	0.45	LAC 1				
Staphylococcus sciuri		19.75	0.43					
Macrococcus carouselicus		10.11	0.30					
003613	**Low discrimination**							**003613**
					OXI	NOV		
Staphylococcus sciuri		74.14	0.61	LAC 14	99	99		
Staphylococcus lentus		23.15	0.53		99	99		
Aerococcus viridans		2.40	0.47		1	1		
003620	**Low discrimination**							**003620**
					OXI	NOV		
Staphylococcus vitulinus		86.47	0.54	MAN 99	99	99		
Macrococcus carouselicus		7.67	0.34		99	99		
Staphylococcus muscae		4.37	0.18		1	1		
003621	**Acceptable identification**							**003621**
					OXI	NOV		
Staphylococcus sciuri		92.00	0.44	MAN 97	99	99		
Staphylococcus vitulinus		7.24	0.13		99	99		
Aerococcus viridans		0.77	0.17		1	1		
003623	**Low discrimination**							**003623**
					OXI	NOV		
Staphylococcus sciuri		62.59	0.28	MAN 97	99	99		
Aerococcus viridans		28.95	0.37	LAC 14	1	1		
Staphylococcus lentus		8.46	0.12		99	99		
003630	**Excellent identification**							**003630**
Staphylococcus vitulinus		99.43	0.95					
Staphylococcus sciuri		0.52	0.56					
Macrococcus carouselicus		0.06	0.30					
003631	**Low discrimination**							**003631**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		79.33	0.75					
Staphylococcus vitulinus		19.10	0.54					
Staphylococcus lentus		1.51	0.41					
003632	**Good identification**							**003632**
					OXI	NOV		
Staphylococcus vitulinus		91.01	0.54	LAC 1	99	99		
Staphylococcus sciuri		7.60	0.40		99	99		
Aerococcus viridans		1.13	0.39		1	1		
003633	**Low discrimination**							**003633**
					OXI	NOV		
Staphylococcus sciuri		67.95	0.59	LAC 14	99	99		
Staphylococcus lentus		28.12	0.53		99	99		
Aerococcus viridans		2.92	0.47		1	1		
003640	**Low discrimination**							**003640**
					OXI	NOV		
Staphylococcus auricularis		54.95	0.53	ESL 1	1	1		
Macrococcus carouselicus		15.21	0.34		99	99		
Staphylococcus sciuri		13.58	0.40		99	99		
003641	**Low discrimination**							**003641**
					OXI	NOV	PYR	
Staphylococcus sciuri		59.77	0.58	MAN 97	99	99	1	
Kocuria kristinae		26.16	0.45		99	1	93	
Rothia mucilaginosa		10.20	0.33		1	1	75	
003642	**Low discrimination**							**003642**
Staphylococcus auricularis		39.77	0.33	ESL 1				
Aerococcus viridans		18.53	0.42	LAC 10				
Staphylococcus piscifermentans		18.14	0.36					
003643	**Low discrimination**							**003643**
					OXI	NOV	PYR	
Staphylococcus sciuri		46.58	0.42	MAN 97	99	99	1	
Kocuria kristinae		31.31	0.32	LAC 14	99	1	93	
Aerococcus viridans		17.29	0.49		1	1	95	
003650	**Low discrimination**							**003650**
Staphylococcus sciuri		77.71	0.71	MNS 89				
Staphylococcus vitulinus		11.23	0.45					
Macrococcus bovicus		8.35	0.48					
003651	**Very good identification**							**003651**
					OXI	NOV		
Staphylococcus sciuri		99.08	0.89		99	99		
Staphylococcus lentus		0.77	0.48		99	99		
Aerococcus viridans		0.06	0.39		1	1		
003652	**Low discrimination**							**003652**
					OXI	NOV		
Staphylococcus sciuri		85.38	0.54	MNS 89	99	99		
Aerococcus viridans		10.19	0.51	LAC 14	1	1		
Staphylococcus piscifermentans		1.43	0.28		1	1		
003653	**Low discrimination**							**003653**
					OXI	NOV		
Staphylococcus sciuri		83.07	0.73	LAC 14	99	99		
Staphylococcus lentus		14.00	0.59		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		2.86	0.59		1	1		
003660	**Low discrimination**							**003660**
					OXI	NOV		
Staphylococcus sciuri		79.74	0.37	MAN 97	99	99		
Staphylococcus vitulinus		13.49	0.13	MNS 89	99	99		
Staphylococcus auricularis		4.32	0.12		1	1		
003661	**Very good identification**							**003661**
					OXI	NOV	PYR	
Staphylococcus sciuri		98.00	0.56	MAN 97	99	99	1	
Aerococcus viridans		0.87	0.30		1	1	95	
Kocuria kristinae		0.57	0.04		99	1	93	
003662	**Low discrimination**							**003662**
					OXI	NOV		
Aerococcus viridans		63.04	0.42	NIT 1	1	1		
Staphylococcus sciuri		36.96	0.21		99	99		
003663	**Low discrimination**							**003663**
					OXI	NOV		
Staphylococcus sciuri		63.91	0.40	MAN 97	99	99		
Aerococcus viridans		31.43	0.49	LAC 14	1	1		
Staphylococcus lentus		4.66	0.18		99	99		
003670	**Low discrimination**							**003670**
					OXI	NOV		
Staphylococcus sciuri		65.66	0.68	MNS 89	99	99		
Staphylococcus vitulinus		34.00	0.54		99	99		
Aerococcus viridans		0.19	0.32		1	1		
003671	**Very good identification**							**003671**
					OXI	NOV		
Staphylococcus sciuri		98.82	0.87		99	99		
Staphylococcus lentus		1.01	0.48		99	99		
Aerococcus viridans		0.08	0.39		1	1		
003672	**Low discrimination**							**003672**
					OXI	NOV		
Staphylococcus sciuri		82.09	0.52	MNS 89	99	99		
Aerococcus viridans		12.99	0.51	LAC 14	1	1		
Staphylococcus vitulinus		2.64	0.13		99	99		
003673	**Low discrimination**							**003673**
					OXI	NOV		
Staphylococcus sciuri		78.53	0.71	LAC 14	99	99		
Staphylococcus lentus		17.54	0.59		99	99		
Aerococcus viridans		3.58	0.59		1	1		
003700	**Low discrimination**							**003700**
					OXI	NOV		
Macrococcus carouselicus		69.84	0.68	NIT 6	99	99		
Rothia mucilaginosa		28.72	0.56		1	1		
Staphylococcus sciuri		0.76	0.34		99	99		
003701	**Good identification**							**003701**
					OXI	NOV		
Rothia mucilaginosa		90.97	0.66	VPT 95	1	1		
Staphylococcus sciuri		6.48	0.52		99	99		
Staphylococcus lentus		1.55	0.41		99	99		
003702	**Low discrimination**							**003702**
					OXI	NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus carouselicus		40.73	0.27	NIT 6	99	99	
Staphylococcus piscifermentans		26.10	0.39	LAC 1	1	1	
Rothia mucilaginosa		16.75	0.15		1	1	
003703	**Low discrimination**						**003703**
					OXI	NOV	
Staphylococcus lentus		71.05	0.53	MAN 99	99	99	
Staphylococcus sciuri		13.66	0.36		99	99	
Rothia mucilaginosa		11.90	0.25		1	1	
003710	**Low discrimination**						**003710**
Macrococcus carouselicus		56.57	0.64	NIT 6			
Staphylococcus sciuri		31.07	0.65				
Staphylococcus vitulinus		8.70	0.45				
003711	**Low discrimination**						**003711**
					OXI	NOV	
Staphylococcus sciuri		57.52	0.83		99	99	
Staphylococcus lentus		42.06	0.82		99	99	
Rothia mucilaginosa		0.25	0.25		1	1	
003712	**Low discrimination**						**003712**
Staphylococcus lentus		51.27	0.53	MNS 99			
Staphylococcus sciuri		39.40	0.49				
Macrococcus carouselicus		4.45	0.23				
003713	**Good identification**						**003713**
					OXI	NOV	
Staphylococcus lentus		94.00	0.94		99	99	
Staphylococcus sciuri		5.90	0.67		99	99	
Aerococcus viridans		0.10	0.47		1	1	
003720	**Low discrimination**						**003720**
					OXI	NOV	
Macrococcus carouselicus		40.19	0.27	NIT 6	99	99	
Staphylococcus sciuri		32.60	0.31	XYL 1	99	99	
Rothia mucilaginosa		16.53	0.15		1	1	
003721	**Low discrimination**						**003721**
					OXI	NOV	
Staphylococcus sciuri		66.28	0.50	MAN 97	99	99	
Staphylococcus lentus		20.98	0.41		99	99	
Rothia mucilaginosa		12.46	0.25		1	1	
003723	**Low discrimination**						**003723**
					OXI	NOV	
Staphylococcus lentus		84.77	0.53	MAN 99	99	99	
Staphylococcus sciuri		12.30	0.34		99	99	
Aerococcus viridans		2.93	0.37		1	1	
003730	**Low discrimination**						**003730**
Staphylococcus vitulinus		47.60	0.54	GAL 1			
Staphylococcus sciuri		47.43	0.62				
Staphylococcus lentus		3.76	0.41				
003731	**Low discrimination**						**003731**
Staphylococcus sciuri		50.73	0.81				
Staphylococcus lentus		49.17	0.82				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus vitulinus		0.06	0.13				
003732	**Low discrimination**						**003732**
					OXI	NOV	
Staphylococcus lentus		60.23	0.53	MNS 99	99	99	
Staphylococcus sciuri		34.92	0.46		99	99	
Aerococcus viridans		2.68	0.39		1	1	
003733	**Very good identification**						**003733**
					OXI	NOV	
Staphylococcus lentus		95.36	0.94		99	99	
Staphylococcus sciuri		4.52	0.65		99	99	
Aerococcus viridans		0.10	0.47		1	1	
003740	**Good identification**						**003740**
					OXI	NOV	
Rothia mucilaginosa		93.04	0.64	VPT 95	1	1	
Staphylococcus sciuri		3.96	0.46		99	99	
Macrococcus carouselicus		0.98	0.27		99	99	
003741	**Low discrimination**						**003741**
					OXI	NOV	
Rothia mucilaginosa		87.79	0.74	VPT 95	1	1	
Staphylococcus sciuri		10.08	0.64		99	99	
Staphylococcus lentus		1.30	0.48		99	99	
003743	**Low discrimination**						**003743**
					OXI	NOV	
Staphylococcus lentus		60.31	0.59	MAN 99	99	99	
Staphylococcus sciuri		21.49	0.48		99	99	
Rothia mucilaginosa		11.61	0.33		1	1	
003750	**Good identification**						**003750**
					YEP		
Staphylococcus sciuri		93.07	0.77	MNS 89	57		
Staphylococcus lentus		3.00	0.48		1		
Macrococcus bovicus		2.21	0.41		99		
003751	**Low discrimination**						**003751**
					OXI	NOV	
Staphylococcus sciuri		71.54	0.95		99	99	
Staphylococcus lentus		28.23	0.89		99	99	
Rothia mucilaginosa		0.20	0.33		1	1	
003752	**Low discrimination**						**003752**
					OXI	NOV	
Staphylococcus sciuri		56.08	0.60	MNS 89	99	99	
Staphylococcus lentus		39.39	0.59	LAC 14	99	99	
Aerococcus viridans		3.45	0.51		1	1	
003753	**Low discrimination**						**003753**
					OXI	NOV	
Staphylococcus lentus		89.41	1.00		99	99	
Staphylococcus sciuri		10.40	0.79		99	99	
Aerococcus viridans		0.19	0.59		1	1	
003760	**Low discrimination**						**003760**
					OXI	NOV	
Staphylococcus sciuri		74.39	0.43	MAN 97	99	99	
Rothia mucilaginosa		23.39	0.23	MNS 89	1	1	
Aerococcus viridans		1.18	0.22		1	1	
003761	**Low discrimination**						**003761**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Staphylococcus sciuri		77.19	0.62	MAN 97	99	99		
Staphylococcus lentus		13.19	0.48		99	99		
Rothia mucilaginosa		9.00	0.33		1	1		
003762	**Low discrimination**							**003762**
					OXI	NOV		
Staphylococcus sciuri		45.82	0.27	MAN 97	99	99		
Aerococcus viridans		40.25	0.42	MNS 89	1	1		
Staphylococcus lentus		13.93	0.18	LAC 14	99	99		
003763	**Low discrimination**							**003763**
					OXI	NOV		
Staphylococcus lentus		74.15	0.59	MAN 99	99	99		
Staphylococcus sciuri		19.93	0.45		99	99		
Aerococcus viridans		5.05	0.49		1	1		
003770	**Very good identification**							**003770**
Staphylococcus sciuri		95.49	0.74	MNS 89				
Staphylococcus lentus		4.08	0.48					
Staphylococcus vitulinus		0.26	0.13					
003771	**Low discrimination**							**003771**
					OXI			
Staphylococcus sciuri		65.56	0.93		99			
Staphylococcus lentus		34.29	0.89		99			
Staphylococcus equorum		0.06	0.27		1			
003772	**Low discrimination**							**003772**
					OXI	NOV		
Staphylococcus sciuri		49.01	0.58	MNS 89	99	99		
Staphylococcus lentus		45.63	0.59	LAC 14	99	99		
Aerococcus viridans		4.00	0.51		1	1		
003773	**Good identification**							**003773**
					OXI	NOV		
Staphylococcus lentus		91.53	1.00		99	99		
Staphylococcus sciuri		8.03	0.77		99	99		
Aerococcus viridans		0.19	0.59		1	1		
004000	**Low discrimination**							**004000**
Dermacoccus nishinomiyaensis		63.53	0.74	PHS 5				
Micrococcus sp.		34.33	0.59					
Kocuria rosea		1.56	0.39					
004002	**Low discrimination**							**004002**
					OXI			
Dermacoccus nishinomiyaensis		60.90	0.33	PHS 5	93			
Micrococcus sp.		32.90	0.18	LAC 1	89			
Kocuria varians		4.70	0.17		13			
004010	**Low discrimination**							**004010**
					NOV	YEP		
Macrococcus bovicus		60.44	0.48	PHS 1	99	99		
Dermacoccus nishinomiyaensis		16.49	0.33		80	1		
Micrococcus sp.		8.91	0.18		1	41		
004011	**Acceptable identification**							**004011**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		93.74	0.46	NIT 89	1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		4.96	0.28	PHS 11	1	99	12	
Macrococcus bovicus		0.94	0.07		99	99	99	
004020	**Low discrimination**							**004020**
Kocuria rosea		38.89	0.32	NIT 90				
Dermacoccus nishinomiyaensis		37.34	0.33	PHS 1				
Micrococcus sp.		20.18	0.18					
004040	**Low discrimination**							**004040**
					COA	OXI		
Staphylococcus aureus subsp. anaerobius		86.14	0.57	SUC 99	99	1		
Micrococcus sp.		9.73	0.36		1	89		
Dermacoccus nishinomiyaensis		2.42	0.33		1	93		
004041	**Low discrimination**							**004041**
					COA	NOV	OXI	
Staphylococcus cohnii subsp. cohnii		52.37	0.36	PHS 9	1	99	1	
Staphylococcus aureus subsp. anaerobius		38.78	0.16	TRE 99	99	1	1	
Micrococcus sp.		8.85	0.01		1	1	89	
004050	**Low discrimination**							**004050**
					OXI			
Macrococcus bovicus		77.33	0.55	PHS 1	99			
Macrococcus equipersicus		7.53	0.36		99			
Staphylococcus kloosii		6.35	0.52		1			
004051	**Low discrimination**							**004051**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		83.35	0.38	PHS 9	1	99	12	
Macrococcus bovicus		11.63	0.14	TRE 99	99	99	99	
Staphylococcus capitis subsp. capitis		5.02	0.04		1	1	1	
004052	**Low discrimination**							**004052**
					OXI			
Staphylococcus kloosii		89.80	0.53	TRE 99	1			
Macrococcus bovicus		10.21	0.14		99			
004070	**Low discrimination**							**004070**
					OXI			
Staphylococcus kloosii		67.00	0.40	TRE 99	1			
Macrococcus bovicus		33.00	0.14		99			
004072	**Good identification**							**004072**
Staphylococcus kloosii		100.0	0.40	TRE 99				
004100	**Good identification**							**004100**
					OXI			
Dermacoccus nishinomiyaensis		94.85	0.59	PHS 5	93			
Micrococcus sp.		2.72	0.18		89			
Kocuria varians		2.07	0.32		13			
004110	**Low discrimination**							**004110**
					YEP			
Macrococcus bovicus		79.13	0.40	PHS 1	99			
Dermacoccus nishinomiyaensis		9.60	0.18		1			
Macrococcus equipersicus		6.65	0.20		99			
004112	**Low discrimination**							**004112**
					OXI			
Staphylococcus kloosii		83.04	0.32	TRE 99	1			
Macrococcus bovicus		16.97	-.01	MLT 95	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
004150	**Low discrimination**						**004150**
					OXI		
Macrococcus bovicus		67.28	0.48	PHS 1	99		
Staphylococcus kloosii		25.00	0.58		1		
Macrococcus equipersicus		6.55	0.28		99		
004152	**Very good identification**						**004152**
					OXI		
Staphylococcus kloosii		97.55	0.59	TRE 99	1		
Macrococcus bovicus		2.45	0.07		99		
004170	**Acceptable identification**						**004170**
					OXI		
Staphylococcus kloosii		90.19	0.46	TRE 99	1		
Macrococcus bovicus		9.81	0.07		99		
004172	**Good identification**						**004172**
Staphylococcus kloosii		100.0	0.46	TRE 99			
004200	**Low discrimination**						**004200**
					COA	OXI	
Staphylococcus aureus subsp. anaerobius		74.79	0.57	MLT 99	99	1	
Dermacoccus nishinomiyaensis		13.26	0.49		1	93	
Micrococcus sp.		11.10	0.38		1	89	
004201	**Low discrimination**						**004201**
					COA	OXI	
Staphylococcus capitis subsp. capitis		48.59	0.28	NIT 89	1	1	
Staphylococcus aureus subsp. anaerobius		33.90	0.16	PHS 11	99	1	
Micrococcus sp.		10.16	0.03	MAN 93	1	89	
004210	**Low discrimination**						**004210**
					OXI	COA	
Macrococcus bovicus		47.77	0.35	PHS 1	99	1	
Staphylococcus aureus subsp. anaerobius		18.78	0.16		1	99	
Staphylococcus vitulinus		17.88	0.21		99	1	
004211	**Good identification**						**004211**
Staphylococcus capitis subsp. capitis		100.0	0.51	NIT 89			
004230	**Good identification**						**004230**
					NOV		
Staphylococcus vitulinus		98.63	0.29	NIT 91	99		
Kocuria rosea		1.37	-.08	PHS 1	1		
004240	**Excellent identification**						**004240**
					COA	OXI	
Staphylococcus aureus subsp. anaerobius		99.98	0.98		99	1	
Micrococcus sp.		0.01	0.15		1	89	
Staphylococcus auricularis		0.00	0.11		1	1	
004241	**Very good identification**						**004241**
					OXI	COA	
Staphylococcus aureus subsp. anaerobius		99.68	0.57	MNS 1	1	99	
Kocuria kristinae		0.32	0.13		99	1	
004242	**Very good identification**						**004242**
Staphylococcus aureus subsp. anaerobius		100.0	0.57	LAC 1			
004250	**Good identification**						**004250**
					OXI	COA	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus aureus subsp. anaerobius		93.97	0.57	MAN 1	1	99		
Macrococcus bovicus		5.63	0.43		99	1		
Macrococcus equipersicus		0.22	0.15		99	1		
004260	**Very good identification**							**004260**
Staphylococcus aureus subsp. anaerobius		100.0	0.57	XYL 1				
004300	**Low discrimination**							**004300**
					COA	OXI		
Dermacoccus nishinomiyaensis		70.48	0.34	PHS 5	1	93		
Staphylococcus aureus subsp. anaerobius		21.09	0.16	SUC 6	99	1		
Kocuria varians		3.28	0.15		1	13		
004310	**Acceptable identification**							**004310**
Macrococcus bovicus		91.78	0.28	PHS 1				
Macrococcus carouselicus		5.17	-.02					
Macrococcus equipersicus		3.05	-.01					
004340	**Very good identification**							**004340**
					COA			
Staphylococcus aureus subsp. anaerobius		99.96	0.57	GAL 1	99			
Staphylococcus epidermidis		0.05	0.01		1			
004350	**Low discrimination**							**004350**
					OXI	COA	NOV	
Macrococcus bovicus		63.35	0.35	PHS 1	99	1	99	
Staphylococcus aureus subsp. anaerobius		24.90	0.16		1	99	1	
Staphylococcus kloosii		9.31	0.37		1	1	99	
004352	**Good identification**							**004352**
Staphylococcus kloosii		100.0	0.38	SUC 9				
004372	**Acceptable identification**							**004372**
Staphylococcus arlettae		90.92	0.35	GLR 92				
Staphylococcus kloosii		9.08	0.26	TRE 99				
004373	**Good identification**							**004373**
Staphylococcus arlettae		100.0	0.37	GLR 92				
004400	**Low discrimination**							**004400**
					OXI			
Staphylococcus cohnii subsp. cohnii		84.07	0.66	PHS 9	1			
Dermacoccus nishinomiyaensis		5.99	0.33		93			
Macrococcus carouselicus		3.73	0.28		99			
004401	**Very good identification**							**004401**
					NOV			
Staphylococcus cohnii subsp. cohnii		98.71	0.67	PHS 9	99			
Staphylococcus schleiferi subsp. schleiferi		1.12	0.19		1			
Staphylococcus auricularis		0.17	0.07		1			
004410	**Low discrimination**							**004410**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		64.44	0.68	PHS 9	1	12		
Macrococcus bovicus		19.83	0.51		99	99		
Staphylococcus kloosii		13.19	0.67		1	99		
004411	**Very good identification**							**004411**
					OXI	NOV	YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		99.14	0.69	PHS 9	1	99	12	
Staphylococcus capitis subsp. capitis		0.39	0.11		1	1	1	
Macrococcus bovicus		0.28	0.10		99	99	99	
004412	**Good identification**							**004412**
					OXI	YEP		
Staphylococcus kloosii		93.94	0.67	MLT 95	1	99		
Staphylococcus cohnii subsp. cohnii		4.28	0.27		1	12		
Macrococcus bovicus		1.32	0.10		99	99		
004413	**Low discrimination**							**004413**
					NOV	YEP	PYR	
Staphylococcus cohnii subsp. cohnii		69.46	0.28	PHS 9	99	12	1	
Aerococcus viridans		16.33	0.27	LAC 1	1	1	95	
Staphylococcus kloosii		14.21	0.26		99	99	52	
004430	**Low discrimination**							**004430**
					OXI	YEP		
Staphylococcus kloosii		79.48	0.54	MLT 95	1	99		
Staphylococcus cohnii subsp. cohnii		15.69	0.27		1	12		
Macrococcus bovicus		4.83	0.10		99	99		
004431	**Good identification**							**004431**
Staphylococcus cohnii subsp. cohnii		100.0	0.28	PHS 9				
004432	**Very good identification**							**004432**
					NOV	YEP		
Staphylococcus kloosii		98.05	0.55	MLT 95	99	99		
Aerococcus viridans		1.95	0.19		1	1		
004433	**Good identification**							**004433**
Aerococcus viridans		100.0	0.27	ESL 90				
004440	**Very good identification**							**004440**
					NOV	YEP		
Staphylococcus cohnii subsp. cohnii		95.99	0.76	PHS 9	99	12		
Staphylococcus kloosii		1.52	0.52		99	99		
Staphylococcus auricularis		1.19	0.34		1	1		
004441	**Very good identification**							**004441**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		99.83	0.77	PHS 9	1	99		
Staphylococcus auricularis		0.13	0.14		1	1		
Kocuria kristinae		0.05	-.02		99	1		
004442	**Low discrimination**							**004442**
					NOV	YEP		
Staphylococcus kloosii		58.63	0.53	MAN 99	99	99		
Staphylococcus cohnii subsp. cohnii		34.64	0.35		99	12		
Staphylococcus auricularis		4.71	0.14		1	1		
004443	**Low discrimination**							**004443**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		88.81	0.36	PHS 9	99	1		
Aerococcus viridans		11.19	0.30	LAC 1	1	95		
004450	**Low discrimination**							**004450**
					OXI	YEP		
Staphylococcus kloosii		49.72	0.93		1	99		
Staphylococcus cohnii subsp. cohnii		40.50	0.79		1	12		
Macrococcus bovicus		9.18	0.59		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
004451	**Very good identification**							**004451**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		98.63	0.79	PHS 9	1	12		
Staphylococcus kloosii		1.13	0.52		1	99		
Macrococcus bovicus		0.21	0.18		99	99		
004452	**Very good identification**							**004452**
					OXI	YEP		
Staphylococcus kloosii		98.97	0.94		1	99		
Staphylococcus cohnii subsp. cohnii		0.75	0.38		1	12		
Macrococcus bovicus		0.17	0.18		99	99		
004453	**Low discrimination**							**004453**
					NOV	YEP	PYR	
Staphylococcus kloosii		47.45	0.53	MNS 1	99	99	52	
Staphylococcus cohnii subsp. cohnii		38.65	0.38		99	12	1	
Aerococcus viridans		11.48	0.39		1	1	95	
004460	**Low discrimination**							**004460**
					YEP			
Staphylococcus cohnii subsp. cohnii		71.91	0.35	PHS 9	12			
Staphylococcus kloosii		28.09	0.40	XYL 1	99			
004461	**Good identification**							**004461**
Staphylococcus cohnii subsp. cohnii		100.0	0.36	PHS 9				
004462	**Low discrimination**							**004462**
					NOV	YEP		
Staphylococcus kloosii		87.86	0.40	MAN 99	99	99		
Aerococcus viridans		12.14	0.22		1	1		
004463	**Good identification**							**004463**
Aerococcus viridans		100.0	0.30	ESL 90				
004470	**Very good identification**							**004470**
					OXI	YEP		
Staphylococcus kloosii		96.12	0.81		1	99		
Staphylococcus cohnii subsp. cohnii		3.16	0.38		1	12		
Macrococcus bovicus		0.72	0.18		99	99		
004471	**Low discrimination**							**004471**
					NOV	YEP	PYR	
Staphylococcus cohnii subsp. cohnii		75.97	0.38	PHS 9	99	12	1	
Staphylococcus kloosii		21.52	0.40	XYL 1	99	99	52	
Aerococcus viridans		2.51	0.20		1	1	95	
004472	**Very good identification**							**004472**
					NOV	YEP		
Staphylococcus kloosii		99.24	0.81		99	99		
Aerococcus viridans		0.42	0.32		1	1		
Staphylococcus arlettae		0.34	0.19		99	99		
004473	**Low discrimination**							**004473**
					NOV	YEP		
Staphylococcus kloosii		41.37	0.40	MNS 1	99	99		
Aerococcus viridans		40.03	0.39		1	1		
Staphylococcus arlettae		18.61	0.22		99	99		
004500	**Low discrimination**							**004500**
					OXI			
Staphylococcus cohnii subsp. cohnii		71.89	0.47	PHS 9	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Macrococcus carouselicus		12.30	0.20	GAL 10	99			
Dermacoccus nishinomiyaensis		8.78	0.18		93			
004501	**Acceptable identification**							**004501**
					NOV			
Staphylococcus cohnii subsp. cohnii		90.70	0.47	PHS 9	99			
Staphylococcus schleiferi subsp. schleiferi		9.30	0.19	GAL 10	1			
004502	**Low discrimination**							**004502**
					YEP			
Staphylococcus kloosii		83.13	0.32	MAN 99	99			
Staphylococcus cohnii subsp. cohnii		16.87	0.05	MLT 95	12			
004510	**Low discrimination**							**004510**
					OXI	YEP		
Staphylococcus kloosii		60.45	0.73	MLT 95	1	99		
Macrococcus bovicus		20.07	0.44		99	99		
Staphylococcus cohnii subsp. cohnii		16.91	0.49		1	12		
004511	**Good identification**							**004511**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		95.75	0.49	PHS 9	1	12		
Staphylococcus kloosii		3.19	0.32	GAL 10	1	99		
Macrococcus bovicus		1.06	0.03		99	99		
004512	**Very good identification**							**004512**
					OXI	YEP		
Staphylococcus kloosii		99.18	0.73	MLT 95	1	99		
Macrococcus bovicus		0.31	0.03		99	99		
Staphylococcus cohnii subsp. cohnii		0.26	0.08		1	12		
004513	**Low discrimination**							**004513**
					NOV	YEP	PYR	
Staphylococcus kloosii		53.43	0.32	MLT 95	99	99	52	
Aerococcus viridans		31.63	0.27	MNS 1	1	1	95	
Staphylococcus cohnii subsp. cohnii		14.95	0.08		99	12	1	
004530	**Very good identification**							**004530**
					OXI	YEP		
Staphylococcus kloosii		97.59	0.60	MLT 95	1	99		
Macrococcus bovicus		1.31	0.03		99	99		
Staphylococcus cohnii subsp. cohnii		1.10	0.08		1	12		
004532	**Very good identification**							**004532**
					NOV	YEP		
Staphylococcus kloosii		98.99	0.61	MLT 95	99	99		
Aerococcus viridans		1.02	0.19		1	1		
004533	**Low discrimination**							**004533**
					NOV	YEP		
Aerococcus viridans		70.31	0.27	ESL 90	1	1		
Staphylococcus kloosii		29.69	0.20	PHS 1	99	99		
004540	**Low discrimination**							**004540**
					NOV	YEP		
Staphylococcus cohnii subsp. cohnii		76.74	0.57	PHS 9	99	12		
Staphylococcus kloosii		21.16	0.58	GAL 10	99	99		
Staphylococcus auricularis		0.95	0.14		1	1		
004541	**Very good identification**							**004541**
					YEP			
Staphylococcus cohnii subsp. cohnii		99.74	0.58	PHS 9	12			
Staphylococcus kloosii		0.26	0.17	GAL 10	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
004542	**Very good identification**							**004542**
					NOV	YEP	PYR	
Staphylococcus kloosii		95.09	0.59	MAN 99	99	99	52	
Staphylococcus cohnii subsp. cohnii		3.22	0.16		99	12	1	
Aerococcus viridans		1.69	0.22		1	1	95	
004543	**Low discrimination**							**004543**
					NOV	YEP	PYR	
Aerococcus viridans		47.06	0.30	ESL 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		41.50	0.16	PHS 1	99	12	1	
Staphylococcus kloosii		11.44	0.18		99	99	52	
004550	**Good identification**							**004550**
					OXI	YEP		
Staphylococcus kloosii		91.74	0.99		1	99		
Staphylococcus cohnii subsp. cohnii		4.28	0.59		1	12		
Macrococcus bovicus		3.74	0.51		99	99		
004551	**Low discrimination**							**004551**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		82.57	0.60	PHS 9	1	12		
Staphylococcus kloosii		16.51	0.58	GAL 10	1	99		
Macrococcus bovicus		0.67	0.10		99	99		
004552	**Excellent identification**							**004552**
					NOV	YEP	PYR	
Staphylococcus kloosii		99.87	1.00		99	99	52	
Aerococcus viridans		0.05	0.32		1	1	95	
Staphylococcus cohnii subsp. cohnii		0.04	0.18		99	12	1	
004553	**Low discrimination**							**004553**
					NOV	YEP	PYR	
Staphylococcus kloosii		84.57	0.59	MNS 1	99	99	52	
Aerococcus viridans		10.54	0.39		1	1	95	
Staphylococcus cohnii subsp. cohnii		3.94	0.19		99	12	1	
004560	**Low discrimination**							**004560**
					YEP			
Staphylococcus kloosii		87.22	0.46	MAN 99	99			
Staphylococcus cohnii subsp. cohnii		12.78	0.16		12			
004562	**Acceptable identification**							**004562**
					NOV	YEP		
Staphylococcus kloosii		93.35	0.46	MAN 99	99	99		
Aerococcus viridans		6.65	0.22		1	1		
004563	**Good identification**							**004563**
Aerococcus viridans		100.0	0.30	ESL 90				
004570	**Very good identification**							**004570**
					OXI	YEP		
Staphylococcus kloosii		99.65	0.87		1	99		
Staphylococcus cohnii subsp. cohnii		0.19	0.18		1	12		
Macrococcus bovicus		0.16	0.10		99	99		
004571	**Low discrimination**							**004571**
					NOV	YEP	PYR	
Staphylococcus kloosii		79.24	0.46	MNS 1	99	99	52	
Staphylococcus cohnii subsp. cohnii		16.01	0.19		99	12	1	
Aerococcus viridans		4.75	0.20		1	1	95	
004572	**Very good identification**							**004572**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus kloosii		98.80	0.87		99	99		
Staphylococcus arlettae		0.99	0.35		99	99		
Aerococcus viridans		0.21	0.32		1	1		
004573	**Low discrimination**							**004573**
					NOV	YEP		
Staphylococcus arlettae		46.71	0.37	GLR 92	99	99		
Staphylococcus kloosii		35.56	0.46	SUC 99	99	99		
Aerococcus viridans		17.73	0.39		1	1		
004600	**Low discrimination**							**004600**
					OXI	COA	NOV	
Macrococcus carouselicus		44.07	0.34	ESL 94	99	1	99	
Staphylococcus muscae		25.12	0.18	PHS 1	1	1	1	
Staphylococcus aureus subsp. anaerobius		12.56	0.16		1	99	1	
004610	**Low discrimination**							**004610**
					OXI			
Macrococcus bovicus		46.63	0.39	PHS 1	99			
Macrococcus carouselicus		27.41	0.30		99			
Staphylococcus kloosii		12.27	0.46		1			
004611	**Low discrimination**							**004611**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		51.87	0.28	PHS 9	1	99		
Staphylococcus capitis subsp. capitis		37.14	0.16	SUC 1	1	1		
Staphylococcus sciuri		7.31	0.06		99	99		
004612	**Acceptable identification**							**004612**
					OXI	NOV	YEP	
Staphylococcus kloosii		92.10	0.47	SUC 9	1	99	99	
Aerococcus viridans		4.63	0.19	MLT 95	1	1	1	
Macrococcus bovicus		3.26	-.02		99	99	99	
004613	**Good identification**							**004613**
Aerococcus viridans		100.0	0.27	ESL 90				
004620	**Very good identification**							**004620**
Staphylococcus muscae		100.0	0.59	NIT 99				
004632	**Low discrimination**							**004632**
					NOV	YEP		
Staphylococcus kloosii		53.77	0.34	SUC 9	99	99		
Staphylococcus arlettae		35.41	0.19	MLT 95	99	99		
Aerococcus viridans		10.82	0.19		1	1		
004640	**Low discrimination**							**004640**
					COA	PYR		
Staphylococcus aureus subsp. anaerobius		89.12	0.57	TRE 1	99	1		
Staphylococcus petrasii subsp. pragensis		8.18	0.33		1	99		
Staphylococcus auricularis		1.27	0.33		1	40		
004641	**Low discrimination**							**004641**
					OXI	COA	NOV	
Kocuria kristinae		69.62	0.39	VPT 90	99	1	1	
Staphylococcus cohnii subsp. cohnii		15.79	0.36	PHS 1	1	1	99	
Staphylococcus aureus subsp. anaerobius		11.69	0.16		1	99	1	
004643	**Low discrimination**							**004643**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		89.75	0.27	VPT 90	99			
Aerococcus viridans		10.26	0.30	PHS 1	1			
004650	**Low discrimination**							**004650**
					OXI	YEP		
Staphylococcus kloosii		61.69	0.72	SUC 9	1	99		
Macrococcus bovicus		28.80	0.47		99	99		
Staphylococcus cohnii subsp. cohnii		5.13	0.38		1	12		
004651	**Low discrimination**							**004651**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		66.26	0.38	PHS 9	1	99	1	
Staphylococcus sciuri		11.09	0.18	SUC 1	99	99	1	
Kocuria kristinae		9.57	0.11		99	1	93	
004652	**Very good identification**							**004652**
					NOV	YEP		
Staphylococcus kloosii		97.69	0.73	SUC 9	99	99		
Aerococcus viridans		1.03	0.32		1	1		
Staphylococcus arlettae		0.85	0.19		99	99		
004653	**Low discrimination**							**004653**
					NOV	YEP		
Aerococcus viridans		47.91	0.39	ESL 90	1	1		
Staphylococcus arlettae		22.28	0.22	PHS 1	99	99		
Staphylococcus kloosii		19.59	0.32		99	99		
004670	**Good identification**							**004670**
					OXI			
Staphylococcus kloosii		94.66	0.60	SUC 9	1			
Staphylococcus arlettae		3.56	0.19		1			
Macrococcus bovicus		1.79	0.05		99			
004672	**Low discrimination**							**004672**
					NOV	YEP		
Staphylococcus arlettae		76.70	0.60	GLR 92	99	99		
Staphylococcus kloosii		22.35	0.61		99	99		
Aerococcus viridans		0.95	0.32		1	1		
004673	**Very good identification**							**004673**
					NOV	YEP		
Staphylococcus arlettae		97.66	0.63	GLR 92	99	99		
Aerococcus viridans		2.12	0.39		1	1		
Staphylococcus kloosii		0.22	0.20		99	99		
004700	**Acceptable identification**							**004700**
					OXI	NOV		
Macrococcus carouselicus		93.14	0.27	ESL 94	99	99		
Staphylococcus auricularis		4.15	0.06	PHS 1	1	1		
Staphylococcus cohnii subsp. cohnii		2.71	0.05		1	99		
004710	**Low discrimination**							**004710**
					OXI			
Staphylococcus kloosii		42.02	0.52	SUC 9	1			
Macrococcus bovicus		35.27	0.31	MLT 95	99			
Macrococcus carouselicus		20.73	0.23		99			
004712	**Good identification**							**004712**
					NOV	YEP		
Staphylococcus kloosii		97.47	0.53	SUC 9	99	99		
Aerococcus viridans		2.53	0.19	MLT 95	1	1		
004730	**Good identification**							**004730**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.40	SUC 9			
004732	**Low discrimination**						**004732**
					NOV	YEP	
Staphylococcus arlettae		63.53	0.35	GLR 92	99	99	
Staphylococcus kloosii		33.04	0.40	MLT 99	99	99	
Aerococcus viridans		3.43	0.19		1	1	
004733	**Acceptable identification**						**004733**
					NOV	YEP	
Staphylococcus arlettae		91.33	0.37	GLR 92	99	99	
Aerococcus viridans		8.67	0.27	MLT 99	1	1	
004741	**Acceptable identification**						**004741**
					OXI	NOV	
Kocuria kristinae		92.97	0.30	VPT 90	99	1	
Staphylococcus cohnii subsp. cohnii		7.03	0.16	PHS 1	1	99	
004742	**Low discrimination**						**004742**
					NOV	YEP	
Staphylococcus kloosii		84.75	0.38	SUC 9	99	99	
Aerococcus viridans		15.25	0.22	MAN 99	1	1	
004750	**Low discrimination**						**004750**
					OXI	YEP	
Staphylococcus kloosii		89.79	0.78	SUC 9	1	99	
Macrococcus bovicus		9.25	0.39		99	99	
Staphylococcus cohnii subsp. cohnii		0.43	0.18		1	12	
004752	**Very good identification**						**004752**
					NOV	YEP	
Staphylococcus kloosii		96.93	0.79	SUC 9	99	99	
Staphylococcus arlettae		2.45	0.35		99	99	
Aerococcus viridans		0.53	0.32		1	1	
004753	**Low discrimination**						**004753**
					NOV	YEP	
Staphylococcus arlettae		49.62	0.37	GLR 92	99	99	
Aerococcus viridans		18.83	0.39	XYL 99	1	1	
Staphylococcus kloosii		14.95	0.38		99	99	
004762	**Low discrimination**						**004762**
					NOV	YEP	
Staphylococcus arlettae		85.35	0.35	GLR 92	99	99	
Staphylococcus kloosii		8.52	0.26	MAN 99	99	99	
Aerococcus viridans		6.13	0.22		1	1	
004763	**Low discrimination**						**004763**
					NOV	YEP	
Staphylococcus arlettae		88.77	0.37	GLR 92	99	99	
Aerococcus viridans		11.23	0.30	MAN 99	1	1	
004770	**Low discrimination**						**004770**
					OXI		
Staphylococcus kloosii		89.33	0.66	SUC 9	1		
Staphylococcus arlettae		9.79	0.35		1		
Staphylococcus sciuri		0.50	0.03		99		
004771	**Low discrimination**						**004771**
					OXI		
Staphylococcus arlettae		70.18	0.37	GLR 92	1		
Staphylococcus sciuri		21.99	0.22	LAC 99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		4.88	0.25		1			
004772	**Good identification**							**004772**
					NOV	YEP		
Staphylococcus arlettae		90.74	0.76	GLR 92	99	99		
Staphylococcus kloosii		9.06	0.67		99	99		
Aerococcus viridans		0.20	0.32		1	1		
004773	**Very good identification**							**004773**
					NOV	YEP		
Staphylococcus arlettae		99.45	0.78	GLR 92	99	99		
Aerococcus viridans		0.38	0.39		1	1		
Staphylococcus kloosii		0.08	0.26		99	99		
005000	**Low discrimination**							**005000**
					NOV			
Dermacoccus nishinomiyaensis		45.01	0.33	ESL 1	80			
Macrococcus carouselicus		28.02	0.28	PHS 5	99			
Micrococcus sp.		24.32	0.18		1			
005010	**Low discrimination**							**005010**
Macrococcus equipersicus		43.74	0.34	URE 86				
Macrococcus bovicus		27.21	0.28	PHS 1				
Macrococcus carouselicus		26.59	0.24					
005050	**Low discrimination**							**005050**
					OXI			
Macrococcus equipersicus		51.72	0.43	URE 86	99			
Macrococcus bovicus		27.77	0.36	PHS 1	99			
Staphylococcus kloosii		20.51	0.52		1			
005052	**Acceptable identification**							**005052**
					OXI	NOV	YEP	
Staphylococcus kloosii		93.23	0.53	TRE 99	1	99	99	
Aerococcus viridans		4.58	0.25		1	1	1	
Macrococcus equipersicus		2.19	0.02		99	99	99	
005053	**Good identification**							**005053**
Aerococcus viridans		100.0	0.33	PHS 1				
005070	**Acceptable identification**							**005070**
					OXI			
Staphylococcus kloosii		90.75	0.40	TRE 99	1			
Macrococcus equipersicus		9.25	0.02		99			
005072	**Low discrimination**							**005072**
					NOV	YEP		
Staphylococcus kloosii		83.58	0.40	TRE 99	99	99		
Aerococcus viridans		16.42	0.25		1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
005073	**Good identification**						**005073**
Aerococcus viridans		100.0	0.33	PHS 1			
005112	**Good identification**						**005112**
Staphylococcus kloosii		100.0	0.32	TRE 99			
005150	**Low discrimination**						**005150**
					OXI		
Staphylococcus kloosii		53.88	0.58	TRE 99	1		
Macrococcus equipersicus		30.01	0.35		99		
Macrococcus bovicus		16.11	0.28		99		
005152	**Very good identification**						**005152**
					NOV	YEP	
Staphylococcus kloosii		97.53	0.59	TRE 99	99	99	
Aerococcus viridans		2.47	0.25		1	1	
005153	**Low discrimination**						**005153**
					NOV	YEP	
Aerococcus viridans		85.39	0.33	PHS 1	1	1	
Staphylococcus kloosii		14.61	0.18	TRE 95	99	99	
005170	**Good identification**						**005170**
Staphylococcus kloosii		100.0	0.46	TRE 99			
005172	**Acceptable identification**						**005172**
					NOV	YEP	
Staphylococcus kloosii		90.81	0.46	TRE 99	99	99	
Aerococcus viridans		9.19	0.25		1	1	
005173	**Good identification**						**005173**
Aerococcus viridans		100.0	0.33	PHS 1			
005200	**Low discrimination**						**005200**
					COA	OXI	
Macrococcus carouselicus		72.58	0.34	PHS 1	1	99	
Staphylococcus aureus subsp. anaerobius		20.69	0.16	TRE 94	99	1	
Dermacoccus nishinomiyaensis		3.67	0.08		1	93	
005210	**Low discrimination**						**005210**
Staphylococcus vitulinus		48.75	0.29	NIT 91			
Macrococcus carouselicus		42.49	0.30	PHS 1			
Macrococcus bovicus		5.35	0.16				
005230	**Good identification**						**005230**
Staphylococcus vitulinus		100.0	0.38	NIT 91			
005240	**Very good identification**						**005240**
Staphylococcus aureus subsp. anaerobius		100.0	0.57	ESL 1			
005251	**Good identification**						**005251**
Staphylococcus sciuri		100.0	0.28	NIT 98			
005252	**Low discrimination**						**005252**
					NOV	YEP	
Staphylococcus kloosii		66.82	0.32	SUC 9	99	99	
Aerococcus viridans		33.18	0.25	TRE 99	1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
005253	**Low discrimination**							**005253**
					OXI	NOV		
Aerococcus viridans		62.77	0.33	PHS 1	1	1		
Staphylococcus sciuri		37.23	0.12	TRE 95	99	99		
005273	**Low discrimination**							**005273**
					OXI	NOV	YEP	
Aerococcus viridans		41.17	0.33	PHS 1	1	1	1	
Staphylococcus arlettae		40.41	0.22	TRE 95	1	99	99	
Staphylococcus sciuri		18.42	0.10		99	99	57	
005300	**Good identification**							**005300**
Macrococcus carouselicus		100.0	0.27	PHS 1				
005350	**Low discrimination**							**005350**
					OXI			
Staphylococcus kloosii		34.75	0.37	SUC 9	1			
Macrococcus bovicus		26.26	0.16	TRE 99	99			
Staphylococcus sciuri		19.64	0.15		99			
005351	**Good identification**							**005351**
Staphylococcus sciuri		100.0	0.34	NIT 98				
005352	**Low discrimination**							**005352**
					NOV	YEP		
Staphylococcus kloosii		79.63	0.38	SUC 9	99	99		
Aerococcus viridans		20.37	0.25	TRE 99	1	1		
005371	**Good identification**							**005371**
Staphylococcus sciuri		100.0	0.32	NIT 98				
005372	**Low discrimination**							**005372**
					NOV	YEP		
Staphylococcus arlettae		83.20	0.35	GLR 92	99	99		
Aerococcus viridans		8.50	0.25	TRE 99	1	1		
Staphylococcus kloosii		8.31	0.26		99	99		
005373	**Low discrimination**							**005373**
					OXI	NOV	YEP	
Staphylococcus arlettae		72.56	0.37	GLR 92	1	99	99	
Aerococcus viridans		13.05	0.33	TRE 99	1	1	1	
Staphylococcus sciuri		11.33	0.15		99	99	57	
005400	**Good identification**							**005400**
					OXI	YEP		
Macrococcus carouselicus		97.86	0.53	PHS 1	99	67		
Staphylococcus cohnii subsp. cohnii		1.42	0.25		1	12		
Staphylococcus kloosii		0.37	0.26		1	99		
005403	**Good identification**							**005403**
Aerococcus viridans		100.0	0.37	PHS 1				
005410	**Low discrimination**							**005410**
					OXI			
Macrococcus carouselicus		54.90	0.49	PHS 1	99			
Staphylococcus kloosii		32.19	0.67		1			
Macrococcus equipersicus		5.76	0.34		99			
005411	**Low discrimination**							**005411**
					OXI	NOV	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		52.99	0.28	ESL 1	1	99	1	
Macrococcus carouselicus		17.07	0.08	PHS 9	99	99	1	
Aerococcus viridans		12.46	0.27		1	1	95	
005412	**Good identification**							**005412**
					OXI	NOV	YEP	
Staphylococcus kloosii		94.28	0.67	MLT 95	1	99	99	
Aerococcus viridans		4.22	0.39		1	1	1	
Macrococcus carouselicus		1.50	0.08		99	99	67	
005413	**Acceptable identification**							**005413**
					NOV	YEP		
Aerococcus viridans		91.18	0.47	PHS 1	1	1		
Staphylococcus kloosii		8.82	0.26		99	99		
005422	**Good identification**							**005422**
Aerococcus viridans		100.0	0.29	PHS 1				
005423	**Good identification**							**005423**
Aerococcus viridans		100.0	0.37	PHS 1				
005430	**Good identification**							**005430**
					OXI	NOV	YEP	
Staphylococcus kloosii		90.41	0.54	MLT 95	1	99	99	
Macrococcus carouselicus		6.23	0.08		99	99	67	
Aerococcus viridans		1.95	0.19		1	1	1	
005431	**Low discrimination**							**005431**
					OXI	NOV		
Aerococcus viridans		68.87	0.27	PHS 1	1	1		
Staphylococcus sciuri		31.14	0.04	LAC 90	99	99		
005432	**Low discrimination**							**005432**
					NOV	YEP		
Staphylococcus kloosii		84.81	0.55	MLT 95	99	99		
Aerococcus viridans		15.19	0.39		1	1		
005433	**Good identification**							**005433**
Aerococcus viridans		100.0	0.47	PHS 1				
005440	**Low discrimination**							**005440**
					OXI	YEP		
Staphylococcus kloosii		50.73	0.52	MAN 99	1	99		
Staphylococcus cohnii subsp. cohnii		32.47	0.35		1	12		
Macrococcus carouselicus		7.12	0.12		99	67		
005441	**Low discrimination**							**005441**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		88.81	0.36	ESL 1	99	1		
Aerococcus viridans		11.19	0.30	PHS 9	1	95		
005442	**Low discrimination**							**005442**
					NOV	YEP		
Staphylococcus kloosii		76.28	0.53	MAN 99	99	99		
Aerococcus viridans		23.72	0.42		1	1		
005443	**Good identification**							**005443**
Aerococcus viridans		100.0	0.49	PHS 1				
005450	**Good identification**							**005450**
					OXI			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		94.66	0.93		1			
Macrococcus equipersicus		2.41	0.43		99			
Macrococcus bovicus		1.94	0.39		99			
005451	**Low discrimination**							**005451**
					NOV	YEP	PYR	
Staphylococcus kloosii		42.77	0.52	MNS 1	99	99	52	
Staphylococcus cohnii subsp. cohnii		37.74	0.38		99	12	1	
Aerococcus viridans		11.21	0.39		1	1	95	
005452	**Excellent identification**							**005452**
					OXI	NOV	YEP	
Staphylococcus kloosii		99.02	0.94		1	99	99	
Aerococcus viridans		0.93	0.51		1	1	1	
Macrococcus equipersicus		0.02	0.02		99	99	99	
005453	**Low discrimination**							**005453**
					OXI	NOV	YEP	
Aerococcus viridans		67.63	0.59	PHS 1	1	1	1	
Staphylococcus kloosii		31.06	0.53		1	99	99	
Staphylococcus sciuri		0.69	0.02		99	99	57	
005460	**Low discrimination**							**005460**
					NOV	YEP		
Staphylococcus kloosii		86.98	0.40	MAN 99	99	99		
Aerococcus viridans		13.02	0.22		1	1		
005461	**Good identification**							**005461**
Aerococcus viridans		100.0	0.30	PHS 1				
005462	**Low discrimination**							**005462**
					NOV	YEP		
Aerococcus viridans		55.44	0.42	PHS 1	1	1		
Staphylococcus kloosii		44.56	0.40		99	99		
005463	**Good identification**							**005463**
Aerococcus viridans		100.0	0.49	PHS 1				
005470	**Very good identification**							**005470**
					OXI	NOV	YEP	
Staphylococcus kloosii		99.36	0.81		1	99	99	
Aerococcus viridans		0.45	0.32		1	1	1	
Macrococcus equipersicus		0.10	0.02		99	99	99	
005471	**Low discrimination**							**005471**
					OXI	NOV	YEP	
Aerococcus viridans		42.03	0.39	PHS 1	1	1	1	
Staphylococcus kloosii		40.10	0.40	LAC 90	1	99	99	
Staphylococcus sciuri		17.87	0.16		99	99	57	
005472	**Very good identification**							**005472**
					NOV	YEP		
Staphylococcus kloosii		96.05	0.81		99	99		
Aerococcus viridans		3.62	0.51		1	1		
Staphylococcus arlettae		0.33	0.19		99	99		
005473	**Low discrimination**							**005473**
					NOV	YEP		
Aerococcus viridans		85.73	0.59	PHS 1	1	1		
Staphylococcus kloosii		9.84	0.40		99	99		
Staphylococcus arlettae		4.43	0.22		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
005500	**Good identification**							**005500**
					OXI	YEP		
Macrococcus carouselicus		97.61	0.45	PHS 1	99	67		
Staphylococcus kloosii		1.67	0.32		1	99		
Staphylococcus cohnii subsp. cohnii		0.37	0.05		1	12		
005502	**Low discrimination**							**005502**
					OXI	NOV	YEP	
Staphylococcus kloosii		43.39	0.32	MAN 99	1	99	99	
Aerococcus viridans		33.02	0.29	MLT 95	1	1	1	
Macrococcus carouselicus		23.59	0.04		99	99	67	
005503	**Good identification**							**005503**
Aerococcus viridans		100.0	0.37	PHS 1				
005510	**Low discrimination**							**005510**
					OXI			
Staphylococcus kloosii		68.56	0.73	MLT 95	1			
Macrococcus carouselicus		25.82	0.41		99			
Macrococcus equipersicus		2.71	0.26		99			
005511	**Low discrimination**							**005511**
					OXI	NOV	YEP	
Staphylococcus kloosii		32.61	0.32	MLT 95	1	99	99	
Staphylococcus sciuri		24.33	0.12	MNS 1	99	99	57	
Aerococcus viridans		20.91	0.27		1	1	1	
005512	**Very good identification**							**005512**
					OXI	NOV	YEP	
Staphylococcus kloosii		97.41	0.73	MLT 95	1	99	99	
Aerococcus viridans		2.25	0.39		1	1	1	
Macrococcus carouselicus		0.34	0.00		99	99	67	
005513	**Low discrimination**							**005513**
					NOV	YEP		
Aerococcus viridans		84.20	0.47	PHS 1	1	1		
Staphylococcus kloosii		15.80	0.32		99	99		
005522	**Low discrimination**							**005522**
					NOV	YEP		
Aerococcus viridans		75.27	0.29	PHS 1	1	1		
Staphylococcus kloosii		24.73	0.20		99	99		
005523	**Good identification**							**005523**
Aerococcus viridans		100.0	0.37	PHS 1				
005530	**Very good identification**							**005530**
					OXI	NOV	YEP	
Staphylococcus kloosii		97.44	0.60	MLT 95	1	99	99	
Macrococcus carouselicus		1.48	0.00		99	99	67	
Aerococcus viridans		1.08	0.19		1	1	1	
005531	**Low discrimination**							**005531**
					OXI	NOV	YEP	
Aerococcus viridans		44.10	0.27	PHS 1	1	1	1	
Staphylococcus sciuri		38.71	0.10	LAC 90	99	99	57	
Staphylococcus kloosii		17.19	0.19		1	99	99	
005532	**Good identification**							**005532**
					NOV	YEP		
Staphylococcus kloosii		91.55	0.61	MLT 95	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		8.45	0.39		1	1		
005533	**Good identification**							**005533**
					NOV	YEP		
Aerococcus viridans		95.52	0.47	PHS 1	1	1		
Staphylococcus kloosii		4.48	0.20		99	99		
005540	**Low discrimination**							**005540**
					OXI	YEP		
Staphylococcus kloosii		89.59	0.58	MAN 99	1	99		
Staphylococcus cohnii subsp. cohnii		3.28	0.16		1	12		
Macrococcus carouselicus		2.78	0.04		99	67		
005541	**Low discrimination**							**005541**
					NOV	YEP	PYR	
Aerococcus viridans		47.48	0.30	PHS 1	1	1	95	
Staphylococcus cohnii subsp. cohnii		41.87	0.16	LAC 90	99	12	1	
Staphylococcus kloosii		10.66	0.17		99	99	52	
005542	**Low discrimination**							**005542**
					NOV	YEP		
Staphylococcus kloosii		86.19	0.59	MAN 99	99	99		
Aerococcus viridans		13.81	0.42		1	1		
005543	**Good identification**							**005543**
					NOV	YEP		
Aerococcus viridans		97.37	0.49	PHS 1	1	1		
Staphylococcus kloosii		2.63	0.18		99	99		
005550	**Very good identification**							**005550**
					OXI			
Staphylococcus kloosii		98.85	0.99		1			
Macrococcus equipersicus		0.56	0.35		99			
Macrococcus bovicus		0.45	0.32		99			
005551	**Low discrimination**							**005551**
					OXI	NOV	YEP	
Staphylococcus kloosii		75.01	0.58	MNS 1	1	99	99	
Staphylococcus sciuri		11.08	0.24		99	99	57	
Aerococcus viridans		10.13	0.39		1	1	1	
005552	**Excellent identification**							**005552**
					NOV	YEP		
Staphylococcus kloosii		99.52	1.00		99	99		
Aerococcus viridans		0.48	0.51		1	1		
005553	**Low discrimination**							**005553**
					OXI	NOV	YEP	
Aerococcus viridans		51.98	0.59	PHS 1	1	1	1	
Staphylococcus kloosii		46.35	0.59		1	99	99	
Staphylococcus sciuri		1.03	0.08		99	99	57	
005560	**Acceptable identification**							**005560**
					NOV	YEP		
Staphylococcus kloosii		92.84	0.46	MAN 99	99	99		
Aerococcus viridans		7.16	0.22		1	1		
005561	**Good identification**							**005561**
Aerococcus viridans		100.0	0.30	PHS 1				
005562	**Low discrimination**							**005562**
					NOV	YEP		
Staphylococcus kloosii		60.94	0.46	MAN 99	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		39.06	0.42		1	1		
005563	**Good identification**							**005563**
Aerococcus viridans		100.0	0.49	PHS 1				
005570	**Very good identification**							**005570**
					OXI	NOV	YEP	
Staphylococcus kloosii		99.71	0.87		1	99	99	
Aerococcus viridans		0.23	0.32		1	1	1	
Staphylococcus sciuri		0.06	0.03		99	99	57	
005571	**Low discrimination**							**005571**
					OXI	NOV	YEP	
Staphylococcus kloosii		50.36	0.46	MNS 1	1	99	99	
Aerococcus viridans		27.19	0.39		1	1	1	
Staphylococcus sciuri		22.45	0.22		99	99	57	
005572	**Very good identification**							**005572**
					NOV	YEP		
Staphylococcus kloosii		97.14	0.87		99	99		
Aerococcus viridans		1.89	0.51		1	1		
Staphylococcus arlettae		0.97	0.35		99	99		
005573	**Low discrimination**							**005573**
					NOV	YEP		
Aerococcus viridans		64.81	0.59	PHS 1	1	1		
Staphylococcus arlettae		18.97	0.37		99	99		
Staphylococcus kloosii		14.45	0.46		99	99		
005600	**Very good identification**							**005600**
Macrococcus carouselicus		100.0	0.59	PHS 1				
005603	**Low discrimination**							**005603**
					OXI	NOV	PYR	
Aerococcus viridans		82.81	0.37	PHS 1	1	1	95	
Kocuria kristinae		8.95	-.05		99	1	93	
Staphylococcus sciuri		8.25	0.00		99	99	1	
005610	**Low discrimination**							**005610**
					OXI			
Macrococcus carouselicus		89.68	0.55	PHS 1	99			
Staphylococcus vitulinus		3.70	0.24		99			
Staphylococcus kloosii		2.56	0.46		1			
005611	**Acceptable identification**							**005611**
					OXI	NOV		
Staphylococcus sciuri		94.01	0.47	NIT 98	99	99		
Macrococcus carouselicus		4.41	0.14		99	99		
Aerococcus viridans		1.58	0.27		1	1		
005612	**Low discrimination**							**005612**
					OXI	NOV	YEP	
Staphylococcus kloosii		52.11	0.47	SUC 9	1	99	99	
Aerococcus viridans		23.59	0.39	MLT 95	1	1	1	
Macrococcus carouselicus		17.01	0.14		99	99	67	
005613	**Low discrimination**							**005613**
					OXI	NOV		
Staphylococcus sciuri		51.77	0.31	NIT 98	99	99		
Aerococcus viridans		48.23	0.47	LAC 14	1	1		
005622	**Good identification**							**005622**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.29	PHS 1				
005623	**Good identification**							**005623**
Aerococcus viridans		100.0	0.37	PHS 1				
005630	**Low discrimination**							**005630**
Staphylococcus vitulinus		74.12	0.33	NIT 91				
Staphylococcus sciuri		13.37	0.26	PHS 1				
Macrococcus carouselicus		6.72	0.14					
005631	**Good identification**							**005631**
					OXI	NOV		
Staphylococcus sciuri		97.16	0.45	NIT 98	99	99		
Aerococcus viridans		2.17	0.27		1	1		
Staphylococcus vitulinus		0.67	-.08		99	99		
005632	**Low discrimination**							**005632**
					NOV	YEP		
Aerococcus viridans		44.85	0.39	PHS 1	1	1		
Staphylococcus kloosii		24.77	0.34		99	99		
Staphylococcus arlettae		16.31	0.19		99	99		
005633	**Low discrimination**							**005633**
					OXI	NOV	YEP	
Aerococcus viridans		49.60	0.47	PHS 1	1	1	1	
Staphylococcus sciuri		40.15	0.29		99	99	57	
Staphylococcus arlettae		10.25	0.22		1	99	99	
005641	**Low discrimination**							**005641**
					OXI	NOV	PYR	
Kocuria kristinae		58.62	0.27	VPT 90	99	1	93	
Staphylococcus sciuri		34.68	0.28	PHS 1	99	99	1	
Aerococcus viridans		6.70	0.30		1	1	95	
005642	**Low discrimination**							**005642**
					NOV	YEP		
Aerococcus viridans		75.87	0.42	PHS 1	1	1		
Staphylococcus kloosii		24.13	0.32		99	99		
005643	**Low discrimination**							**005643**
					OXI	NOV	PYR	
Aerococcus viridans		74.81	0.49	PHS 1	1	1	95	
Kocuria kristinae		18.19	0.15		99	1	93	
Staphylococcus sciuri		7.01	0.12		99	99	1	
005650	**Low discrimination**							**005650**
					OXI			
Staphylococcus kloosii		77.40	0.72	SUC 9	1			
Staphylococcus sciuri		14.29	0.41		99			
Macrococcus bovicus		4.02	0.27		99			
005651	**Very good identification**							**005651**
					OXI	NOV	YEP	
Staphylococcus sciuri		97.38	0.59	NIT 98	99	99	57	
Aerococcus viridans		1.75	0.39		1	1	1	
Staphylococcus kloosii		0.66	0.31		1	99	99	
005652	**Low discrimination**							**005652**
					OXI	NOV	YEP	
Staphylococcus kloosii		88.36	0.73	SUC 9	1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		8.42	0.51		1	1	1	
Staphylococcus sciuri		2.45	0.24		99	99	57	
005653	**Low discrimination**							**005653**
					OXI	NOV	YEP	
Staphylococcus sciuri		47.64	0.43	NIT 98	99	99	57	
Aerococcus viridans		47.20	0.59	LAC 14	1	1	1	
Staphylococcus arlettae		2.44	0.22		1	99	99	
005662	**Low discrimination**							**005662**
					NOV	YEP		
Aerococcus viridans		73.95	0.42	PHS 1	1	1		
Staphylococcus arlettae		20.17	0.19		99	99		
Staphylococcus kloosii		5.88	0.20		99	99		
005663	**Low discrimination**							**005663**
					OXI	NOV	YEP	
Aerococcus viridans		81.59	0.49	PHS 1	1	1	1	
Staphylococcus arlettae		12.65	0.22		1	99	99	
Staphylococcus sciuri		5.76	0.10		99	99	57	
005670	**Low discrimination**							**005670**
					OXI	NOV	YEP	
Staphylococcus kloosii		60.65	0.60	SUC 9	1	99	99	
Staphylococcus sciuri		33.78	0.38		99	99	57	
Aerococcus viridans		2.78	0.32		1	1	1	
005671	**Very good identification**							**005671**
					OXI	NOV	YEP	
Staphylococcus sciuri		96.43	0.57	NIT 98	99	99	57	
Aerococcus viridans		2.29	0.39		1	1	1	
Staphylococcus arlettae		1.07	0.22		1	99	99	
005672	**Low discrimination**							**005672**
					NOV	YEP		
Staphylococcus arlettae		70.08	0.60	GLR 92	99	99		
Staphylococcus kloosii		20.42	0.61		99	99		
Aerococcus viridans		7.79	0.51		1	1		
005673	**Low discrimination**							**005673**
					OXI	NOV	YEP	
Staphylococcus arlettae		74.04	0.63	GLR 92	1	99	99	
Aerococcus viridans		14.48	0.59		1	1	1	
Staphylococcus sciuri		11.02	0.41		99	99	57	
005700	**Good identification**							**005700**
Macrococcus carouselicus		99.76	0.51	PHS 1				
Staphylococcus sciuri		0.24	0.04					
005703	**Low discrimination**							**005703**
					OXI	NOV		
Aerococcus viridans		83.80	0.37	PHS 1	1	1		
Staphylococcus sciuri		16.20	0.06		99	99		
005710	**Low discrimination**							**005710**
					OXI			
Macrococcus carouselicus		78.69	0.47	PHS 1	99			
Staphylococcus kloosii		10.18	0.52		1			
Staphylococcus sciuri		9.49	0.35		99			
005711	**Good identification**							**005711**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		93.27	0.53	NIT 98				
Staphylococcus lentus		4.96	0.29					
Macrococcus carouselicus		0.97	0.06					
005712	**Low discrimination**							**005712**
					OXI	NOV	YEP	
Staphylococcus kloosii		69.19	0.53	SUC 9	1	99	99	
Aerococcus viridans		16.13	0.39	MLT 95	1	1	1	
Staphylococcus sciuri		9.69	0.18		99	99	57	
005713	**Low discrimination**							**005713**
					OXI	NOV		
Staphylococcus lentus		43.88	0.40	NIT 99	99	99		
Staphylococcus sciuri		37.92	0.37		99	99		
Aerococcus viridans		18.20	0.47		1	1		
005722	**Good identification**							**005722**
Aerococcus viridans		100.0	0.29	PHS 1				
005723	**Low discrimination**							**005723**
					OXI	NOV		
Aerococcus viridans		87.27	0.37	PHS 1	1	1		
Staphylococcus sciuri		12.73	0.04		99	99		
005730	**Low discrimination**							**005730**
					OXI			
Staphylococcus sciuri		65.14	0.32	NIT 98	99			
Staphylococcus kloosii		23.15	0.40	MNS 89	1			
Macrococcus carouselicus		7.23	0.06		99			
005731	**Acceptable identification**							**005731**
					OXI	NOV		
Staphylococcus sciuri		92.43	0.51	NIT 98	99	99		
Staphylococcus lentus		6.51	0.29		99	99		
Aerococcus viridans		1.06	0.27		1	1		
005732	**Low discrimination**							**005732**
					NOV	YEP		
Staphylococcus arlettae		44.94	0.35	GLR 92	99	99		
Staphylococcus kloosii		23.38	0.40	MLT 99	99	99		
Aerococcus viridans		21.80	0.39		1	1		
005733	**Low discrimination**							**005733**
					OXI	YEP		
Staphylococcus lentus		38.98	0.40	NIT 99	99	1		
Staphylococcus sciuri		25.41	0.35		99	57		
Staphylococcus arlettae		18.93	0.37		1	99		
005740	**Low discrimination**							**005740**
					OXI			
Staphylococcus kloosii		41.73	0.37	SUC 9	1			
Macrococcus carouselicus		26.55	0.10	MAN 99	99			
Staphylococcus sciuri		23.58	0.15		99			
005741	**Low discrimination**							**005741**
					OXI	NOV	PYR	
Staphylococcus sciuri		70.66	0.34	NIT 98	99	99	1	
Kocuria kristinae		20.51	0.17	MAN 97	99	1	93	
Aerococcus viridans		7.03	0.30		1	1	95	
005742	**Low discrimination**							**005742**
					NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		61.83	0.42	PHS 1	1	1		
Staphylococcus kloosii		38.17	0.38		99	99		
005743	**Low discrimination**							**005743**
					OXI	NOV	PYR	
Aerococcus viridans		76.93	0.49	PHS 1	1	1	95	
Staphylococcus sciuri		13.99	0.18		99	99	1	
Kocuria kristinae		6.23	0.05		99	1	93	
005750	**Low discrimination**							**005750**
					OXI			
Staphylococcus kloosii		82.53	0.78	SUC 9	1			
Staphylococcus sciuri		15.23	0.47		99			
Macrococcus bovicus		0.95	0.19		99			
005751	**Very good identification**							**005751**
					OXI	NOV		
Staphylococcus sciuri		95.72	0.65	NIT 98	99	99		
Staphylococcus lentus		2.75	0.35		99	99		
Aerococcus viridans		0.88	0.39		1	1		
005752	**Good identification**							**005752**
					OXI	NOV	YEP	
Staphylococcus kloosii		90.62	0.79	SUC 9	1	99	99	
Aerococcus viridans		4.45	0.51		1	1	1	
Staphylococcus sciuri		2.51	0.30		99	99	57	
005753	**Low discrimination**							**005753**
					OXI	NOV		
Staphylococcus sciuri		42.93	0.49	NIT 98	99	99		
Staphylococcus lentus		26.81	0.47	LAC 14	99	99		
Aerococcus viridans		21.91	0.59		1	1		
005761	**Low discrimination**							**005761**
					OXI	NOV	PYR	
Staphylococcus sciuri		83.22	0.32	NIT 98	99	99	1	
Aerococcus viridans		10.98	0.30	MAN 97	1	1	95	
Staphylococcus gallinarum		5.80	0.02		1	99	12	
005762	**Low discrimination**							**005762**
					NOV	YEP		
Staphylococcus arlettae		57.25	0.35	GLR 92	99	99		
Aerococcus viridans		37.03	0.42	MAN 99	1	1		
Staphylococcus kloosii		5.72	0.26		99	99		
005763	**Low discrimination**							**005763**
					NOV	PYR		
Aerococcus viridans		45.53	0.49	PHS 1	1	95		
Staphylococcus arlettae		39.99	0.37		99	1		
Staphylococcus gallinarum		6.55	0.10		99	12		
005770	**Low discrimination**							**005770**
					OXI			
Staphylococcus kloosii		59.16	0.66	SUC 9	1			
Staphylococcus sciuri		32.95	0.44		99			
Staphylococcus arlettae		6.49	0.35		1			
005771	**Good identification**							**005771**
					OXI	YEP		
Staphylococcus sciuri		90.09	0.63	NIT 98	99	57		
Staphylococcus lentus		3.43	0.35		99	1		
Staphylococcus arlettae		2.91	0.37		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
005772	**Low discrimination**						**005772**
					NOV	YEP	
Staphylococcus arlettae		88.63	0.76	GLR 92	99	99	
Staphylococcus kloosii		8.85	0.67		99	99	
Aerococcus viridans		1.74	0.51		1	1	
005773	**Low discrimination**						**005773**
					OXI	YEP	
Staphylococcus arlettae		87.11	0.78	GLR 92	1	99	
Staphylococcus sciuri		4.44	0.47		99	57	
Staphylococcus lentus		3.68	0.47		99	1	
006000	**Low discrimination**						**006000**
Dermacoccus nishinomiyaensis		67.89	0.69	PHS 5			
Kocuria rosea		26.66	0.59				
Micrococcus sp.		3.43	0.33				
006001	**Low discrimination**						**006001**
					OXI		
Staphylococcus capitis subsp. capitis		47.40	0.41	PHS 11	1		
Kocuria rosea		24.26	0.33	MAN 93	35		
Dermacoccus nishinomiyaensis		11.86	0.28		93		
006002	**Low discrimination**						**006002**
					OXI		
Dermacoccus nishinomiyaensis		46.51	0.28	PHS 5	93		
Kocuria varians		31.18	0.32	LAC 1	13		
Kocuria rosea		18.26	0.18		35		
006010	**Low discrimination**						**006010**
					YEP	NOV	
Kocuria rosea		64.08	0.39	PHS 1	1	1	
Dermacoccus nishinomiyaensis		14.83	0.28	MAN 10	1	80	
Macrococcus bovicus		10.74	0.28		99	99	
006011	**Very good identification**						**006011**
Staphylococcus capitis subsp. capitis		98.63	0.64	PHS 11			
Staphylococcus carnosus subsp. carnosus		0.90	0.23				
Kocuria rosea		0.42	0.13				
006013	**Low discrimination**						**006013**
Staphylococcus carnosus subsp. carnosus		64.04	0.36	ARG 99			
Staphylococcus capitis subsp. capitis		35.96	0.29	BGA 90			
006020	**Low discrimination**						**006020**
					OXI		
Kocuria rosea		84.03	0.51	PHS 1	35		
Staphylococcus muscae		6.46	0.18		1		
Dermacoccus nishinomiyaensis		5.04	0.28		93		
006030	**Acceptable identification**						**006030**
					NOV		
Kocuria rosea		91.67	0.32	PHS 1	1		
Staphylococcus vitulinus		8.34	0.09	MAN 10	99		
006040	**Low discrimination**						**006040**
					COA	OXI	
Dermacoccus nishinomiyaensis		26.62	0.28	PHS 5	1	93	
Staphylococcus auricularis		20.36	0.32	MLT 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus aureus subsp. anaerobius		17.05	0.16		99	1		
006050	**Acceptable identification**							**006050**
					NOV	YEP	OXI	
Macrococcus bovicus		93.36	0.36	NIT 10	99	99	99	
Staphylococcus cohnii subsp. cohnii		4.23	0.14	PHS 1	99	12	1	
Kocuria rosea		2.41	-.02		1	1	35	
006100	**Low discrimination**							**006100**
					OXI			
Dermacoccus nishinomiyaensis		82.34	0.54	PHS 5	93			
Kocuria varians		15.57	0.47		13			
Kocuria rosea		1.72	0.18		35			
006101	**Low discrimination**							**006101**
					OXI			
Staphylococcus schleiferi subsp. schleiferi		61.80	0.29	VPT 99	1			
Kocuria varians		20.74	0.25	ARG 86	13			
Dermacoccus nishinomiyaensis		12.74	0.13		93			
006102	**Low discrimination**							**006102**
					OXI			
Kocuria varians		84.08	0.35	PHS 1	13			
Dermacoccus nishinomiyaensis		15.92	0.13		93			
006120	**Low discrimination**							**006120**
					OXI			
Kocuria varians		77.74	0.37	PHS 1	13			
Dermacoccus nishinomiyaensis		11.82	0.13		93			
Kocuria rosea		10.44	0.10		35			
006140	**Low discrimination**							**006140**
					OXI			
Kocuria varians		73.97	0.33	PHS 1	13			
Dermacoccus nishinomiyaensis		18.00	0.13		93			
Staphylococcus auricularis		8.03	0.12		1			
006150	**Good identification**							**006150**
					OXI			
Macrococcus bovicus		96.73	0.28	NIT 10	99			
Staphylococcus kloosii		3.27	0.17	PHS 1	1			
006200	**Low discrimination**							**006200**
					OXI			
Dermacoccus nishinomiyaensis		45.73	0.44	PHS 5	93			
Kocuria rosea		31.26	0.39	SUC 6	35			
Staphylococcus muscae		9.27	0.18		1			
006201	**Good identification**							**006201**
					OXI			
Staphylococcus capitis subsp. capitis		95.46	0.47	PHS 11	1			
Kocuria rosea		2.92	0.13	MAN 93	35			
Dermacoccus nishinomiyaensis		0.82	0.03		93			
006210	**Low discrimination**							**006210**
					NOV	OXI		
Staphylococcus vitulinus		78.17	0.41	PHS 1	99	99		
Staphylococcus capitis subsp. capitis		12.64	0.29		1	1		
Kocuria rosea		6.09	0.20		1	35		
006211	**Very good identification**							**006211**
					OXI	NOV		
Staphylococcus capitis subsp. capitis		99.84	0.70	PHS 11	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		0.08	0.10		99	99	
Staphylococcus vitulinus		0.06	0.00		99	99	
006213	**Good identification**						**006213**
Staphylococcus capitis subsp. capitis		100.0	0.35	PHS 11			
006220	**Very good identification**						**006220**
					NOV	OXI	
Staphylococcus muscae		98.30	0.59	TRE 99	1	1	
Kocuria rosea		1.44	0.32		1	35	
Staphylococcus vitulinus		0.13	0.09		99	99	
006230	**Acceptable identification**						**006230**
					NOV	OXI	
Staphylococcus vitulinus		91.81	0.50	PHS 1	99	99	
Staphylococcus muscae		7.06	0.18		1	1	
Kocuria rosea		1.13	0.12		1	35	
006231	**Low discrimination**						**006231**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		81.62	0.29	PHS 11	1	1	
Staphylococcus vitulinus		13.79	0.09	XYL 1	99	99	
Staphylococcus sciuri		4.59	0.08		99	99	
006240	**Very good identification**						**006240**
					COA		
Staphylococcus aureus subsp. anaerobius		98.18	0.57	NIT 1	99		
Staphylococcus auricularis		1.09	0.31		1		
Staphylococcus epidermidis		0.26	0.17		1		
006243	**Good identification**						**006243**
					YEP		
Staphylococcus epidermidis		97.05	0.33	VPT 96	1		
Staphylococcus chromogenes		2.95	-.02	URE 92	95		
006251	**Low discrimination**						**006251**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		78.11	0.29	PHS 11	1	1	
Staphylococcus sciuri		21.89	0.22	MLT 1	99	99	
006300	**Low discrimination**						**006300**
					OXI		
Dermacoccus nishinomiyaensis		68.11	0.29	PHS 5	93		
Kocuria varians		27.52	0.29	SUC 6	13		
Kocuria rosea		2.47	-.02		35		
006303	**Low discrimination**						**006303**
					YEP		
Staphylococcus hyicus		88.08	0.34	ARG 96	1		
Staphylococcus chromogenes		8.58	0.17	TRE 92	95		
Staphylococcus epidermidis		1.79	0.07		1		
006311	**Low discrimination**						**006311**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		87.37	0.29	PHS 11	1	1	
Staphylococcus sciuri		12.63	0.16	GAL 1	99	99	
006340	**Low discrimination**						**006340**
					COA		
Staphylococcus epidermidis		49.05	0.33	VPT 96	1		
Staphylococcus aureus subsp. anaerobius		33.72	0.16	URE 92	99		
Kocuria varians		5.62	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
006341	**Good identification**							**006341**
					YEP			
Staphylococcus epidermidis		98.33	0.38	VPT 96	1			
Staphylococcus chromogenes		1.67	-.02	URE 92	95			
006342	**Acceptable identification**							**006342**
Staphylococcus epidermidis		94.67	0.43	VPT 96				
Staphylococcus hominis subsp. hominis		3.53	0.17	URE 92				
Kocuria varians		0.97	0.04					
006343	**Acceptable identification**							**006343**
					COA	YEP	PYR	
Staphylococcus epidermidis		93.77	0.48	VPT 96	1	1	1	
Staphylococcus chromogenes		4.53	0.17	URE 92	1	95	45	
Staphylococcus intermedius		1.23	0.07		99	1	90	
006351	**Acceptable identification**							**006351**
					OXI	COA		
Staphylococcus sciuri		94.50	0.28	ESL 99	99	1		
Staphylococcus intermedius		5.50	0.00	TRE 97	1	99		
006400	**Low discrimination**							**006400**
					NOV			
Staphylococcus auricularis		39.00	0.46	PHS 1	1			
Staphylococcus cohnii subsp. cohnii		19.44	0.43		99			
Staphylococcus muscae		13.25	0.18		1			
006401	**Low discrimination**							**006401**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		51.72	0.39	VPT 99	1			
Staphylococcus cohnii subsp. cohnii		37.97	0.44	ARG 86	99			
Staphylococcus auricularis		7.81	0.26		1			
006410	**Low discrimination**							**006410**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		64.54	0.45	NIT 7	1	12		
Macrococcus bovicus		29.32	0.32	PHS 9	99	99		
Staphylococcus kloosii		1.77	0.26		1	99		
006411	**Low discrimination**							**006411**
					NOV			
Staphylococcus cohnii subsp. cohnii		65.85	0.46	NIT 7	99			
Staphylococcus capitis subsp. capitis		27.56	0.29	PHS 9	1			
Staphylococcus carnosus subsp. carnosus		5.26	0.16		1			
006413	**Low discrimination**							**006413**
					NOV			
Staphylococcus carnosus subsp. carnosus		85.32	0.28	ARG 99	1			
Staphylococcus felis		11.98	0.04	BGA 90	1			
Staphylococcus cohnii subsp. cohnii		2.70	0.05		99			
006420	**Very good identification**							**006420**
Staphylococcus muscae		99.79	0.59	SUC 99				
Kocuria rosea		0.13	0.10					
Kocuria varians		0.05	0.10					
006440	**Low discrimination**							**006440**
					NOV			
Staphylococcus auricularis		59.39	0.53	PHS 1	1			
Staphylococcus cohnii subsp. cohnii		40.17	0.53		99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		0.27	0.06		1			
006441	**Low discrimination**							**006441**
					NOV			
Staphylococcus cohnii subsp. cohnii		86.51	0.54	NIT 7	99			
Staphylococcus auricularis		13.12	0.34	PHS 9	1			
Staphylococcus schleiferi subsp. schleiferi		0.38	-.02		1			
006442	**Acceptable identification**							**006442**
					NOV			
Staphylococcus auricularis		94.21	0.34	PHS 1	1			
Staphylococcus cohnii subsp. cohnii		5.79	0.12	LAC 10	99			
006450	**Low discrimination**							**006450**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		65.05	0.56	NIT 7	1	12		
Macrococcus bovicus		21.77	0.39	PHS 9	99	99		
Staphylococcus kloosii		10.72	0.52		1	99		
006451	**Very good identification**							**006451**
					OXI	PYR		
Staphylococcus cohnii subsp. cohnii		98.01	0.56	NIT 7	1	1		
Staphylococcus sciuri		1.09	0.12	PHS 9	99	1		
Staphylococcus saprophyticus subsp. bovis		0.59	0.06		1	99		
006452	**Low discrimination**							**006452**
					PYR	YEP		
Staphylococcus kloosii		89.91	0.53	NIT 1	52	99		
Staphylococcus cohnii subsp. cohnii		5.09	0.14		1	12		
Staphylococcus saprophyticus subsp. bovis		3.30	0.06		99	80		
006470	**Low discrimination**							**006470**
					OXI	YEP		
Staphylococcus kloosii		75.34	0.40	NIT 1	1	99		
Staphylococcus cohnii subsp. cohnii		18.48	0.14		1	12		
Macrococcus bovicus		6.18	-.02		99	99		
006472	**Good identification**							**006472**
Staphylococcus kloosii		100.0	0.40	NIT 1				
006501	**Low discrimination**							**006501**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		89.55	0.39	VPT 99	1			
Staphylococcus cohnii subsp. cohnii		7.31	0.24	ARG 86	99			
Staphylococcus auricularis		1.50	0.07		1			
006512	**Good identification**							**006512**
Staphylococcus kloosii		100.0	0.32	NIT 1				
006540	**Low discrimination**							**006540**
					OXI	NOV	YEP	
Staphylococcus auricularis		47.29	0.34	PHS 1	1	1	1	
Staphylococcus cohnii subsp. cohnii		31.99	0.34	GAL 10	1	99	12	
Macrococcus caseolyticus		16.70	0.14		99	99	99	
006541	**Low discrimination**							**006541**
					NOV			
Staphylococcus cohnii subsp. cohnii		83.98	0.34	NIT 7	99			
Staphylococcus auricularis		12.73	0.14	PHS 9	1			
Staphylococcus schleiferi subsp. schleiferi		3.28	-.02	GAL 10	1			
006550	**Low discrimination**							**006550**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	YEP	
Staphylococcus kloosii		47.10	0.58	NIT 1	1	99	
Macrococcus bovicus		21.12	0.32		99	99	
Staphylococcus cohnii subsp. cohnii		16.36	0.36		1	12	
006551	**Low discrimination**						**006551**
					OXI	PYR	
Staphylococcus cohnii subsp. cohnii		69.52	0.37	NIT 7	1	1	
Staphylococcus saprophyticus subsp. bovis		15.11	0.18	PHS 9	1	99	
Staphylococcus sciuri		13.50	0.18	GAL 10	99	1	
006552	**Good identification**						**006552**
Staphylococcus kloosii		92.98	0.59	NIT 1			
Staphylococcus saprophyticus subsp. bovis		7.02	0.18				
006570	**Good identification**						**006570**
Staphylococcus kloosii		100.0	0.46	NIT 1			
006572	**Good identification**						**006572**
Staphylococcus kloosii		100.0	0.46	NIT 1			
006600	**Very good identification**						**006600**
					OXI	NOV	
Staphylococcus muscae		97.10	0.59	XYL 99	1	1	
Staphylococcus auricularis		2.66	0.45		1	1	
Macrococcus carouselicus		0.11	0.10		99	99	
006603	**Low discrimination**						**006603**
					YEP		
Staphylococcus hyicus		86.72	0.34	ARG 96	1		
Staphylococcus chromogenes		10.79	0.20	GAL 92	95		
Staphylococcus auricularis		1.37	0.06		1		
006610	**Low discrimination**						**006610**
					OXI	NOV	
Staphylococcus vitulinus		63.42	0.36	PHS 1	99	99	
Staphylococcus muscae		23.43	0.18		1	1	
Staphylococcus sciuri		5.43	0.23		99	99	
006611	**Low discrimination**						**006611**
					OXI	NOV	
Staphylococcus sciuri		53.64	0.41	ESL 99	99	99	
Staphylococcus capitis subsp. capitis		45.00	0.35		1	1	
Staphylococcus vitulinus		0.78	-.05		99	99	
006620	**Excellent identification**						**006620**
					OXI	NOV	
Staphylococcus muscae		100.0	1.00		1	1	
Staphylococcus vitulinus		0.00	0.04		99	99	
Staphylococcus auricularis		0.00	0.04		1	1	
006621	**Very good identification**						**006621**
					OXI	NOV	
Staphylococcus muscae		99.96	0.59	MNS 1	1	1	
Staphylococcus sciuri		0.04	0.08		99	99	
006622	**Very good identification**						**006622**
Staphylococcus muscae		100.0	0.59	LAC 1			
006630	**Good identification**						**006630**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Staphylococcus muscae		92.96	0.59	MAN 1	1	1		
Staphylococcus vitulinus		6.87	0.45		99	99		
Staphylococcus sciuri		0.16	0.20		99	99		
006631	**Low discrimination**							**006631**
					OXI	NOV		
Staphylococcus sciuri		56.83	0.39	ESL 99	99	99		
Staphylococcus muscae		40.20	0.18		1	1		
Staphylococcus vitulinus		2.97	0.04		99	99		
006640	**Low discrimination**							**006640**
					PYR			
Staphylococcus auricularis		72.24	0.53	PHS 1	40			
Staphylococcus muscae		11.40	0.18		10			
Staphylococcus petrasii subsp. pragensis		5.76	0.13		99			
006641	**Low discrimination**							**006641**
					OXI	NOV	PYR	
Staphylococcus auricularis		57.41	0.33	PHS 1	1	1	40	
Kocuria kristinae		18.28	0.16	MNS 10	99	1	93	
Staphylococcus sciuri		17.78	0.22		99	99	1	
006642	**Low discrimination**							**006642**
					PYR			
Staphylococcus auricularis		66.48	0.33	PHS 1	40			
Staphylococcus hominis subsp. hominis		23.71	0.26	LAC 10	1			
Staphylococcus haemolyticus		7.56	0.15		98			
006650	**Low discrimination**							**006650**
					OXI	PYR		
Staphylococcus sciuri		37.11	0.34	ESL 99	99	1		
Staphylococcus saprophyticus subsp. bovis		26.36	0.31	MNS 89	1	99		
Macrococcus bovicus		21.07	0.27		99	1		
006651	**Very good identification**							**006651**
					OXI	PYR		
Staphylococcus sciuri		96.98	0.53	ESL 99	99	1		
Staphylococcus saprophyticus subsp. bovis		2.13	0.18		1	99		
Staphylococcus cohnii subsp. cohnii		0.89	0.15		1	1		
006653	**Low discrimination**							**006653**
					OXI	NOV	PYR	
Staphylococcus sciuri		85.74	0.37	ESL 99	99	99	1	
Staphylococcus chromogenes		5.57	0.08	LAC 14	1	1	45	
Staphylococcus saprophyticus subsp. bovis		2.89	0.06		1	99	99	
006660	**Very good identification**							**006660**
					OXI	NOV		
Staphylococcus muscae		99.92	0.59	MLT 1	1	1		
Staphylococcus auricularis		0.07	0.12		1	1		
Staphylococcus sciuri		0.02	0.01		99	99		
006671	**Good identification**							**006671**
Staphylococcus sciuri		100.0	0.51	ESL 99				
006673	**Low discrimination**							**006673**
					OXI	YEP		
Staphylococcus sciuri		85.48	0.34	ESL 99	99	57		
Staphylococcus arlettae		11.72	0.22	LAC 14	1	99		
Staphylococcus lentus		2.80	0.05		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
006701	**Low discrimination**							**006701**
					OXI	YEP	NOV	
Staphylococcus hyicus		77.51	0.34	ARG 96	1	1	1	
Staphylococcus chromogenes		9.65	0.20	LAC 92	1	95	1	
Staphylococcus sciuri		4.11	0.16		99	57	99	
006703	**Low discrimination**							**006703**
					YEP	PYR	COA	
Staphylococcus hyicus		89.11	0.56	ARG 96	1	1	24	
Staphylococcus chromogenes		8.68	0.39		95	45	1	
Staphylococcus intermedius		1.81	0.27		1	90	99	
006710	**Low discrimination**							**006710**
Staphylococcus sciuri		72.11	0.28	ESL 99				
Macrococcus bovicus		14.57	0.12	MNS 89				
Macrococcus carouselicus		4.92	-.02					
006711	**Low discrimination**							**006711**
					OXI	COA		
Staphylococcus sciuri		83.77	0.47	ESL 99	99	1		
Staphylococcus lentus		8.99	0.29		99	1		
Staphylococcus intermedius		5.02	0.19		1	99		
006713	**Low discrimination**							**006713**
					OXI	COA	YEP	
Staphylococcus lentus		28.54	0.40	ESL 99	99	1	1	
Staphylococcus intermedius		27.63	0.36		1	99	1	
Staphylococcus chromogenes		13.82	0.28		1	1	95	
006720	**Very good identification**							**006720**
Staphylococcus muscae		100.0	0.59	GAL 1				
006731	**Low discrimination**							**006731**
Staphylococcus sciuri		87.54	0.45	ESL 99				
Staphylococcus lentus		12.46	0.29					
006733	**Low discrimination**							**006733**
Staphylococcus lentus		75.61	0.40	ESL 99				
Staphylococcus sciuri		24.40	0.28					
006740	**Low discrimination**							**006740**
					OXI	NOV		
Staphylococcus auricularis		48.32	0.33	PHS 1	1	1		
Staphylococcus hominis subsp. hominis		15.90	0.26	GAL 10	1	1		
Macrococcus caseolyticus		12.32	0.11		99	99		
006741	**Low discrimination**							**006741**
					OXI	YEP	NOV	
Staphylococcus sciuri		34.60	0.28	ESL 99	99	57	99	
Staphylococcus chromogenes		21.59	0.20	MAN 97	1	95	1	
Rothia mucilaginosa		19.66	0.13		1	1	1	
006742	**Low discrimination**							**006742**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		34.31	0.27	URE 99	1	1	1	
Macrococcus caseolyticus		33.97	0.14	PHS 3	99	80	99	
Staphylococcus haemolyticus		11.39	0.16		1	98	2	
006743	**Low discrimination**							**006743**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					YEP	PYR	COA	
Staphylococcus chromogenes		67.41	0.39	URE 99	95	45	1	
Staphylococcus intermedius		21.08	0.30	ARG 90	1	90	99	
Staphylococcus hyicus		6.99	0.15		1	1	24	
006750	**Low discrimination**							**006750**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		52.72	0.43	URE 99	1	99		
Staphylococcus sciuri		36.03	0.40		99	1		
Macrococcus bovicus		4.52	0.19		99	1		
006751	**Low discrimination**							**006751**
					OXI	PYR		
Staphylococcus sciuri		88.30	0.59	ESL 99	99	1		
Staphylococcus lentus		5.12	0.35		99	6		
Staphylococcus saprophyticus subsp. bovis		3.99	0.31		1	99		
006752	**Low discrimination**							**006752**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		49.94	0.31	URE 99	1	99		
Staphylococcus sciuri		22.22	0.24		99	1		
Staphylococcus kloosii		16.35	0.38		1	52		
006753	**Low discrimination**							**006753**
					OXI	COA		
Staphylococcus lentus		35.03	0.47	ESL 99	99	1		
Staphylococcus sciuri		27.76	0.43		99	1		
Staphylococcus intermedius		25.05	0.39		1	99		
006770	**Acceptable identification**							**006770**
					OXI	PYR		
Staphylococcus sciuri		94.68	0.38	ESL 99	99	1		
Staphylococcus kloosii		3.47	0.25	MNS 89	1	52		
Staphylococcus saprophyticus subsp. bovis		1.86	0.02		1	99		
006771	**Low discrimination**							**006771**
					OXI			
Staphylococcus sciuri		89.00	0.56	ESL 99	99			
Staphylococcus lentus		6.84	0.35		99			
Staphylococcus gallinarum		3.92	0.23		1			
006772	**Low discrimination**							**006772**
					OXI			
Staphylococcus arlettae		63.89	0.35	GLR 92	1			
Staphylococcus sciuri		26.16	0.22	NIT 1	99			
Staphylococcus kloosii		6.38	0.26		1			
006773	**Low discrimination**							**006773**
					OXI			
Staphylococcus lentus		41.51	0.47	ESL 99	99			
Staphylococcus sciuri		24.82	0.40		99			
Staphylococcus gallinarum		16.44	0.31		1			
007000	**Low discrimination**							**007000**
					NOV			
Dermacoccus nishinomiyaensis		65.02	0.28	ESL 1	80			
Kocuria rosea		25.54	0.18	PHS 5	1			
Macrococcus carouselicus		4.59	0.03		99			
007151	**Good identification**							**007151**
Staphylococcus sciuri		100.0	0.28	SUC 99				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
007210	**Acceptable identification**						**007210**
Staphylococcus vitulinus		94.34	0.50	PHS 1			
Staphylococcus sciuri		5.14	0.33				
Macrococcus carouselicus		0.52	0.06				
007211	**Low discrimination**						**007211**
					OXI	NOV	
Staphylococcus sciuri		86.32	0.51	TRE 97	99	99	
Staphylococcus capitis subsp. capitis		11.71	0.29		1	1	
Staphylococcus vitulinus		1.98	0.09		99	99	
007213	**Good identification**						**007213**
Staphylococcus sciuri		100.0	0.35	TRE 97			
007230	**Very good identification**						**007230**
Staphylococcus vitulinus		98.50	0.59	PHS 1			
Staphylococcus sciuri		1.50	0.30				
007231	**Acceptable identification**						**007231**
Staphylococcus sciuri		92.41	0.49	TRE 97			
Staphylococcus vitulinus		7.59	0.18				
007233	**Good identification**						**007233**
Staphylococcus sciuri		100.0	0.32	TRE 97			
007241	**Good identification**						**007241**
Staphylococcus sciuri		100.0	0.32	TRE 97			
007250	**Acceptable identification**						**007250**
Staphylococcus sciuri		94.07	0.44	TRE 97			
Staphylococcus vitulinus		4.64	0.09	MNS 89			
Macrococcus bovicus		1.29	0.04				
007251	**Very good identification**						**007251**
Staphylococcus sciuri		100.0	0.63	TRE 97			
007252	**Good identification**						**007252**
Staphylococcus sciuri		100.0	0.28	TRE 97			
007253	**Good identification**						**007253**
Staphylococcus sciuri		99.20	0.47	TRE 97			
Staphylococcus lentus		0.80	0.05	LAC 14			
007261	**Good identification**						**007261**
Staphylococcus sciuri		100.0	0.29	TRE 97			
007270	**Low discrimination**						**007270**
Staphylococcus sciuri		84.99	0.42	TRE 97			
Staphylococcus vitulinus		15.01	0.18	MNS 89			
007271	**Very good identification**						**007271**
Staphylococcus sciuri		100.0	0.61	TRE 97			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
007273	**Good identification**						**007273**
Staphylococcus sciuri		98.94	0.44	TRE 97			
Staphylococcus lentus		1.06	0.05	LAC 14			
007310	**Low discrimination**						**007310**
Staphylococcus sciuri		89.46	0.38	TRE 97			
Staphylococcus vitulinus		8.54	0.09	MNS 89			
Macrococcus carouselicus		1.99	-.02				
007311	**Very good identification**						**007311**
Staphylococcus sciuri		96.61	0.57	TRE 97			
Staphylococcus lentus		3.39	0.29				
007313	**Low discrimination**						**007313**
Staphylococcus sciuri		56.70	0.41	TRE 97			
Staphylococcus lentus		43.30	0.40	LAC 14			
007330	**Low discrimination**						**007330**
Staphylococcus sciuri		74.50	0.36	TRE 97			
Staphylococcus vitulinus		25.50	0.18	MNS 89			
007331	**Very good identification**						**007331**
Staphylococcus sciuri		95.56	0.55	TRE 97			
Staphylococcus lentus		4.44	0.29				
007333	**Low discrimination**						**007333**
Staphylococcus lentus		50.30	0.40	TRE 99			
Staphylococcus sciuri		49.70	0.38				
007341	**Good identification**						**007341**
					OXI	NOV	
Staphylococcus sciuri		98.34	0.38	TRE 97	99	99	
Rothia mucilaginosa		1.66	-.09	MAN 97	1	1	
007350	**Good identification**						**007350**
Staphylococcus sciuri		100.0	0.50	TRE 97			
007351	**Very good identification**						**007351**
Staphylococcus sciuri		98.14	0.69	TRE 97			
Staphylococcus lentus		1.86	0.35				
007352	**Good identification**						**007352**
Staphylococcus sciuri		96.74	0.34	TRE 97			
Staphylococcus lentus		3.26	0.05	MNS 89			
007353	**Low discrimination**						**007353**
Staphylococcus sciuri		70.82	0.53	TRE 97			
Staphylococcus lentus		29.18	0.47	LAC 14			
007361	**Good identification**						**007361**
					OXI		
Staphylococcus sciuri		95.60	0.35	TRE 97	99		
Staphylococcus gallinarum		4.40	0.02	MAN 97	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
007370	**Good identification**							**007370**
Staphylococcus sciuri		100.0	0.48	TRE 97				
007371	**Very good identification**							**007371**
					OXI			
Staphylococcus sciuri		96.20	0.66	TRE 97	99			
Staphylococcus lentus		2.41	0.35		99			
Staphylococcus gallinarum		1.38	0.23		1			
007372	**Good identification**							**007372**
Staphylococcus sciuri		95.73	0.32	TRE 97				
Staphylococcus lentus		4.27	0.05	MNS 89				
007373	**Low discrimination**							**007373**
					OXI			
Staphylococcus sciuri		56.60	0.50	TRE 97	99			
Staphylococcus lentus		30.92	0.47	LAC 14	99			
Staphylococcus gallinarum		12.25	0.31		1			
007400	**Low discrimination**							**007400**
					OXI	NOV		
Macrococcus carouselicus		87.37	0.28	NIT 6	99	99		
Staphylococcus piscifermentans		9.62	0.25	PHS 1	1	1		
Staphylococcus auricularis		3.01	0.05		1	1		
007402	**Good identification**							**007402**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
007411	**Good identification**							**007411**
					OXI	NOV		
Staphylococcus sciuri		97.99	0.41	SUC 99	99	99		
Staphylococcus cohnii subsp. cohnii		1.07	0.05		1	99		
Staphylococcus carnosus subsp. carnosus		0.94	-.04		1	1		
007431	**Good identification**							**007431**
Staphylococcus sciuri		100.0	0.39	SUC 99				
007442	**Good identification**							**007442**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
007450	**Low discrimination**							**007450**
					OXI			
Staphylococcus kloosii		44.47	0.52	NIT 1	1			
Staphylococcus sciuri		39.78	0.34		99			
Macrococcus bovicus		10.04	0.20		99			
007451	**Very good identification**							**007451**
					OXI			
Staphylococcus sciuri		99.09	0.53	SUC 99	99			
Staphylococcus cohnii subsp. cohnii		0.91	0.15		1			
007452	**Low discrimination**							**007452**
					OXI	NOV	YEP	
Staphylococcus kloosii		83.86	0.53	NIT 1	1	99	99	
Staphylococcus sciuri		11.27	0.18		99	99	57	
Staphylococcus piscifermentans		4.86	0.21		1	1	1	
007453	**Good identification**							**007453**
					OXI	NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		95.70	0.37	SUC 99	99	99		
Staphylococcus lentus		2.37	0.05	LAC 14	99	99		
Aerococcus viridans		1.94	0.18		1	1		
007470	**Low discrimination**							**007470**
					OXI			
Staphylococcus sciuri		72.97	0.32	SUC 99	99			
Staphylococcus kloosii		27.03	0.40	MNS 89	1			
007471	**Good identification**							**007471**
Staphylococcus sciuri		100.0	0.51	SUC 99				
007472	**Low discrimination**							**007472**
					OXI			
Staphylococcus kloosii		71.14	0.40	NIT 1	1			
Staphylococcus sciuri		28.86	0.16		99			
007473	**Acceptable identification**							**007473**
					OXI	NOV		
Staphylococcus sciuri		94.38	0.34	SUC 99	99	99		
Staphylococcus lentus		3.10	0.05	LAC 14	99	99		
Aerococcus viridans		2.53	0.18		1	1		
007502	**Good identification**							**007502**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
007510	**Low discrimination**							**007510**
					OXI			
Staphylococcus sciuri		50.66	0.28	SUC 99	99			
Macrococcus carouselicus		26.68	0.16	MNS 89	99			
Staphylococcus kloosii		11.21	0.32		1			
007511	**Acceptable identification**							**007511**
Staphylococcus sciuri		90.31	0.47	SUC 99				
Staphylococcus lentus		9.69	0.29					
007512	**Low discrimination**							**007512**
					OXI	NOV	YEP	
Staphylococcus kloosii		39.52	0.32	NIT 1	1	99	99	
Staphylococcus piscifermentans		33.65	0.25	MLT 95	1	1	1	
Staphylococcus sciuri		26.83	0.12		99	99	57	
007513	**Low discrimination**							**007513**
					OXI	NOV		
Staphylococcus lentus		69.41	0.40	SUC 99	99	99		
Staphylococcus sciuri		29.69	0.31		99	99		
Staphylococcus carnosus subsp. carnosus		0.90	-.04		1	1		
007531	**Low discrimination**							**007531**
Staphylococcus sciuri		87.54	0.45	SUC 99				
Staphylococcus lentus		12.46	0.29					
007533	**Low discrimination**							**007533**
Staphylococcus lentus		75.61	0.40	SUC 99				
Staphylococcus sciuri		24.40	0.28					
007540	**Low discrimination**							**007540**
					OXI	NOV	YEP	
Staphylococcus piscifermentans		66.66	0.28	URE 90	1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		24.43	0.09	ARG 99	99	99	57	
Staphylococcus kloosii		8.92	0.17		1	99	99	
007541	**Good identification**							**007541**
					OXI	NOV		
Staphylococcus sciuri		95.09	0.28	SUC 99	99	99		
Rothia mucilaginosa		4.91	-.09	MAN 97	1	1		
007542	**Acceptable identification**							**007542**
					NOV	YEP		
Staphylococcus piscifermentans		91.19	0.32	URE 90	1	1		
Staphylococcus kloosii		8.81	0.18	ARG 99	99	99		
007550	**Low discrimination**							**007550**
					OXI			
Staphylococcus kloosii		50.59	0.58	NIT 1	1			
Staphylococcus sciuri		45.25	0.40		99			
Macrococcus bovicus		2.52	0.12		99			
007551	**Good identification**							**007551**
					OXI	YEP		
Staphylococcus sciuri		94.40	0.59	SUC 99	99	57		
Staphylococcus lentus		5.47	0.35		99	1		
Staphylococcus kloosii		0.13	0.17		1	99		
007552	**Low discrimination**							**007552**
					OXI	NOV	YEP	
Staphylococcus kloosii		83.82	0.59	NIT 1	1	99	99	
Staphylococcus sciuri		11.27	0.24		99	99	57	
Staphylococcus piscifermentans		3.76	0.25		1	1	1	
007553	**Low discrimination**							**007553**
					OXI	NOV		
Staphylococcus lentus		55.31	0.47	SUC 99	99	99		
Staphylococcus sciuri		43.83	0.43		99	99		
Aerococcus viridans		0.46	0.18		1	1		
007570	**Low discrimination**							**007570**
					OXI			
Staphylococcus sciuri		72.97	0.38	SUC 99	99			
Staphylococcus kloosii		27.03	0.46	MNS 89	1			
007571	**Low discrimination**							**007571**
					OXI			
Staphylococcus sciuri		89.22	0.56	SUC 99	99			
Staphylococcus lentus		6.85	0.35		99			
Staphylococcus gallinarum		3.93	0.23		1			
007572	**Low discrimination**							**007572**
					OXI	YEP		
Staphylococcus kloosii		68.44	0.46	NIT 1	1	99		
Staphylococcus sciuri		27.76	0.22		99	57		
Staphylococcus lentus		3.80	0.05		99	1		
007573	**Low discrimination**							**007573**
					OXI			
Staphylococcus lentus		49.76	0.47	SUC 99	99			
Staphylococcus sciuri		29.75	0.40		99			
Staphylococcus gallinarum		19.71	0.31		1			
007600	**Low discrimination**							**007600**
					OXI	NOV		
Macrococcus carouselicus		47.32	0.34	NIT 6	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus muscae		26.97	0.18	PHS 1	1	1	
Staphylococcus sciuri		19.13	0.33		99	99	
007601	**Good identification**						**007601**
Staphylococcus sciuri		100.0	0.51	MAN 97			
007602	**Low discrimination**						**007602**
					OXI	NOV	
Staphylococcus piscifermentans		64.97	0.32	URE 90	1	1	
Staphylococcus sciuri		35.03	0.16	ARG 99	99	99	
007603	**Good identification**						**007603**
					OXI	NOV	
Staphylococcus sciuri		97.76	0.35	MAN 97	99	99	
Staphylococcus hyicus		2.24	-.07	LAC 14	1	1	
007610	**Low discrimination**						**007610**
Staphylococcus sciuri		72.89	0.64	MNS 89			
Staphylococcus vitulinus		23.27	0.45				
Micrococcus carouselicus		3.57	0.30				
007611	**Very good identification**						**007611**
Staphylococcus sciuri		99.75	0.82				
Staphylococcus lentus		0.21	0.29				
Staphylococcus vitulinus		0.04	0.04				
007612	**Good identification**						**007612**
					OXI	NOV	
Staphylococcus sciuri		95.75	0.47	MNS 89	99	99	
Staphylococcus piscifermentans		2.35	0.25	LAC 14	1	1	
Staphylococcus vitulinus		1.90	0.04		99	99	
007613	**Very good identification**						**007613**
Staphylococcus sciuri		95.62	0.66	LAC 14			
Staphylococcus lentus		4.38	0.40				
007620	**Very good identification**						**007620**
					OXI	NOV	
Staphylococcus muscae		99.26	0.59	ESL 1	1	1	
Staphylococcus sciuri		0.54	0.30		99	99	
Staphylococcus vitulinus		0.20	0.13		99	99	
007621	**Low discrimination**						**007621**
					OXI	NOV	
Staphylococcus sciuri		81.23	0.49	MAN 97	99	99	
Staphylococcus muscae		18.77	0.18		1	1	
007623	**Good identification**						**007623**
Staphylococcus sciuri		100.0	0.32	MAN 97			
007630	**Low discrimination**						**007630**
					OXI	NOV	
Staphylococcus vitulinus		51.96	0.54	PHS 1	99	99	
Staphylococcus sciuri		45.41	0.61		99	99	
Staphylococcus muscae		2.63	0.18		1	1	
007631	**Very good identification**						**007631**
Staphylococcus sciuri		99.58	0.80				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		0.28	0.29					
Staphylococcus vitulinus		0.14	0.13					
007632	**Acceptable identification**							**007632**
Staphylococcus sciuri		93.37	0.45	MNS 89				
Staphylococcus vitulinus		6.63	0.13	LAC 14				
007633	**Good identification**							**007633**
					OXI			
Staphylococcus sciuri		94.19	0.64	LAC 14	99			
Staphylococcus lentus		5.72	0.40		99			
Staphylococcus equorum		0.09	-.02		1			
007640	**Acceptable identification**							**007640**
					OXI	NOV		
Staphylococcus sciuri		92.81	0.44	MAN 97	99	99		
Staphylococcus piscifermentans		4.97	0.28	MNS 89	1	1		
Staphylococcus auricularis		2.23	0.12		1	1		
007641	**Very good identification**							**007641**
					OXI	NOV	PYR	
Staphylococcus sciuri		99.64	0.63	MAN 97	99	99	1	
Kocuria kristinae		0.26	0.04		99	1	93	
Rothia mucilaginosa		0.10	-.09		1	1	75	
007642	**Low discrimination**							**007642**
					OXI	NOV		
Staphylococcus sciuri		66.98	0.28	MAN 97	99	99		
Staphylococcus piscifermentans		33.02	0.32	MNS 89	1	1		
007643	**Good identification**							**007643**
Staphylococcus sciuri		99.20	0.47	MAN 97				
Staphylococcus lentus		0.80	0.05	LAC 14				
007650	**Very good identification**							**007650**
					OXI			
Staphylococcus sciuri		99.67	0.75	MNS 89	99			
Staphylococcus kloosii		0.11	0.31		1			
Staphylococcus vitulinus		0.09	0.04		99			
007651	**Excellent identification**							**007651**
Staphylococcus sciuri		99.89	0.94					
Staphylococcus lentus		0.11	0.35					
007652	**Very good identification**							**007652**
					OXI	NOV	YEP	
Staphylococcus sciuri		98.43	0.59	MNS 89	99	99	57	
Staphylococcus kloosii		0.73	0.32	LAC 14	1	99	99	
Staphylococcus piscifermentans		0.64	0.25		1	1	1	
007653	**Very good identification**							**007653**
					OXI	NOV		
Staphylococcus sciuri		97.57	0.78	LAC 14	99	99		
Staphylococcus lentus		2.41	0.47		99	99		
Aerococcus viridans		0.02	0.18		1	1		
007660	**Low discrimination**							**007660**
					OXI	NOV		
Staphylococcus sciuri		66.81	0.42	MAN 97	99	99		
Staphylococcus muscae		33.20	0.18	MNS 89	1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
007661	**Very good identification**						**007661**
					OXI		
Staphylococcus sciuri		99.73	0.61	MAN 97	99		
Staphylococcus gallinarum		0.28	0.02		1		
007663	**Good identification**						**007663**
					OXI		
Staphylococcus sciuri		95.03	0.44	MAN 97	99		
Staphylococcus gallinarum		3.95	0.10	LAC 14	1		
Staphylococcus lentus		1.02	0.05		99		
007670	**Very good identification**						**007670**
					OXI		
Staphylococcus sciuri		99.66	0.73	MNS 89	99		
Staphylococcus vitulinus		0.31	0.13		99		
Staphylococcus kloosii		0.04	0.19		1		
007671	**Very good identification**						**007671**
					OXI		
Staphylococcus sciuri		99.75	0.92		99		
Staphylococcus lentus		0.15	0.35		99		
Staphylococcus gallinarum		0.09	0.23		1		
007672	**Very good identification**						**007672**
					OXI	YEP	
Staphylococcus sciuri		98.66	0.57	MNS 89	99	57	
Staphylococcus arlettae		0.83	0.19	LAC 14	1	99	
Staphylococcus lentus		0.26	0.05		99	1	
007673	**Very good identification**						**007673**
					OXI		
Staphylococcus sciuri		95.25	0.75	LAC 14	99		
Staphylococcus lentus		3.12	0.47		99		
Staphylococcus gallinarum		1.24	0.31		1		
007700	**Low discrimination**						**007700**
					OXI	NOV	
Staphylococcus sciuri		51.91	0.38	MAN 97	99	99	
Macrococcus carouselicus		28.35	0.27	MNS 89	99	99	
Rothia mucilaginosa		11.66	0.15		1	1	
007701	**Low discrimination**						**007701**
					OXI	NOV	
Staphylococcus sciuri		89.27	0.57	MAN 97	99	99	
Rothia mucilaginosa		7.43	0.25		1	1	
Staphylococcus lentus		3.13	0.29		99	99	
007702	**Low discrimination**						**007702**
					OXI	NOV	
Staphylococcus piscifermentans		58.90	0.36	URE 90	1	1	
Staphylococcus sciuri		41.10	0.22	ARG 99	99	99	
007703	**Low discrimination**						**007703**
					OXI	NOV	
Staphylococcus sciuri		52.28	0.41	MAN 97	99	99	
Staphylococcus lentus		39.92	0.40	LAC 14	99	99	
Staphylococcus hyicus		7.11	0.15		1	1	
007710	**Very good identification**						**007710**
Staphylococcus sciuri		97.72	0.70	MNS 89			
Macrococcus carouselicus		1.06	0.23				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lentus		0.86	0.29				
007711	**Good identification**						**007711**
Staphylococcus sciuri		90.31	0.88				
Staphylococcus lentus		9.69	0.70				
007712	**Low discrimination**						**007712**
					OXI	NOV	
Staphylococcus sciuri		82.64	0.53	MNS 89	99	99	
Staphylococcus lentus		15.79	0.40	LAC 14	99	99	
Staphylococcus piscifermentans		1.57	0.28		1	1	
007713	**Low discrimination**						**007713**
Staphylococcus lentus		70.04	0.81				
Staphylococcus sciuri		29.96	0.72				
007720	**Low discrimination**						**007720**
					OXI	NOV	
Staphylococcus sciuri		50.94	0.36	MAN 97	99	99	
Staphylococcus muscae		49.06	0.18	MNS 89	1	1	
007721	**Very good identification**						**007721**
					OXI		
Staphylococcus sciuri		95.07	0.55	MAN 97	99		
Staphylococcus lentus		4.42	0.29		99		
Staphylococcus gallinarum		0.51	0.02		1		
007723	**Low discrimination**						**007723**
					OXI		
Staphylococcus lentus		48.37	0.40	MAN 99	99		
Staphylococcus sciuri		47.79	0.38		99		
Staphylococcus gallinarum		3.85	0.10		1		
007730	**Very good identification**						**007730**
Staphylococcus sciuri		98.27	0.67	MNS 89			
Staphylococcus lentus		1.14	0.29				
Staphylococcus vitulinus		0.59	0.13				
007731	**Low discrimination**						**007731**
					OXI		
Staphylococcus sciuri		87.39	0.86		99		
Staphylococcus lentus		12.44	0.70		99		
Staphylococcus gallinarum		0.15	0.23		1		
007732	**Low discrimination**						**007732**
Staphylococcus sciuri		79.79	0.51	MNS 89			
Staphylococcus lentus		20.21	0.40	LAC 14			
007733	**Low discrimination**						**007733**
					OXI		
Staphylococcus lentus		75.06	0.81		99		
Staphylococcus sciuri		24.22	0.69		99		
Staphylococcus gallinarum		0.61	0.31		1		
007740	**Low discrimination**						**007740**
					OXI	NOV	
Staphylococcus sciuri		84.70	0.50	MAN 97	99	99	
Rothia mucilaginosa		11.80	0.23	MNS 89	1	1	
Staphylococcus piscifermentans		3.50	0.32		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
007741	**Good identification**						**007741**
					OXI	NOV	
Staphylococcus sciuri		93.28	0.69	MAN 97	99	99	
Rothia mucilaginosa		4.82	0.33		1	1	
Staphylococcus lentus		1.77	0.35		99	99	
007742	**Low discrimination**						**007742**
					OXI	NOV	
Staphylococcus sciuri		70.69	0.34	MAN 97	99	99	
Staphylococcus piscifermentans		26.93	0.36	MNS 89	1	1	
Staphylococcus lentus		2.38	0.05	LAC 14	99	99	
007743	**Low discrimination**						**007743**
					OXI		
Staphylococcus sciuri		69.69	0.53	MAN 97	99		
Staphylococcus lentus		28.72	0.47	LAC 14	99		
Staphylococcus gallinarum		1.13	0.10		1		
007750	**Very good identification**						**007750**
					OXI	YEP	
Staphylococcus sciuri		99.34	0.81	MNS 89	99	57	
Staphylococcus lentus		0.47	0.35		99	1	
Staphylococcus kloosii		0.11	0.37		1	99	
007751	**Good identification**						**007751**
					OXI		
Staphylococcus sciuri		94.49	1.00		99		
Staphylococcus lentus		5.48	0.76		99		
Staphylococcus gallinarum		0.03	0.23		1		
007752	**Low discrimination**						**007752**
					OXI	YEP	
Staphylococcus sciuri		89.64	0.65	MNS 89	99	57	
Staphylococcus lentus		9.24	0.47	LAC 14	99	1	
Staphylococcus kloosii		0.67	0.38		1	99	
007753	**Low discrimination**						**007753**
					OXI		
Staphylococcus lentus		55.66	0.88		99		
Staphylococcus sciuri		44.11	0.84		99		
Staphylococcus gallinarum		0.22	0.31		1		
007760	**Good identification**						**007760**
					OXI		
Staphylococcus sciuri		98.86	0.48	MAN 97	99		
Staphylococcus gallinarum		1.14	0.02	MNS 89	1		
007761	**Low discrimination**						**007761**
					OXI		
Staphylococcus sciuri		85.72	0.66	MAN 97	99		
Staphylococcus gallinarum		12.08	0.43		1		
Staphylococcus lentus		2.15	0.35		99		
007762	**Low discrimination**						**007762**
					OXI		
Staphylococcus sciuri		82.12	0.32	MAN 97	99		
Staphylococcus gallinarum		14.22	0.10	MNS 89	1		
Staphylococcus lentus		3.67	0.05	LAC 14	99		
007763	**Low discrimination**						**007763**
					OXI		
Staphylococcus gallinarum		57.74	0.51	URE 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		27.26	0.50	MAN 91	99		
Staphylococcus lentus		14.89	0.47		99		
007770	**Very good identification**						**007770**
					OXI		
Staphylococcus sciuri		98.99	0.79	MNS 89	99		
Staphylococcus lentus		0.62	0.35		99		
Staphylococcus gallinarum		0.36	0.23		1		
007771	**Low discrimination**						**007771**
					OXI		
Staphylococcus sciuri		89.15	0.98		99		
Staphylococcus lentus		6.85	0.76		99		
Staphylococcus gallinarum		3.93	0.64		1		
007772	**Low discrimination**						**007772**
					OXI		
Staphylococcus sciuri		82.03	0.63	MNS 89	99		
Staphylococcus lentus		11.21	0.47	LAC 14	99		
Staphylococcus gallinarum		4.44	0.31		1		
007773	**Low discrimination**						**007773**
					OXI		
Staphylococcus lentus		49.90	0.88		99		
Staphylococcus sciuri		29.83	0.81		99		
Staphylococcus gallinarum		19.76	0.72		1		
010000	**Low discrimination**						**010000**
Micrococcus sp.		71.00	0.59	ORN 1			
Dermacoccus nishinomiyaensis		25.22	0.59				
Kocuria rosea		3.23	0.39				
010010	**Low discrimination**						**010010**
					NOV		
Macrococcus bovicus		73.02	0.48	ORN 1	99		
Micrococcus sp.		10.77	0.18		1		
Macrococcus equipersicus		6.14	0.27		99		
010020	**Low discrimination**						**010020**
Kocuria rosea		55.69	0.32	ORN 1			
Micrococcus sp.		28.90	0.18	NIT 90			
Dermacoccus nishinomiyaensis		10.26	0.18				
010040	**Low discrimination**						**010040**
					NOV	YEP	
Micrococcus sp.		88.94	0.36	ORN 1	1	41	
Dermacoccus nishinomiyaensis		4.24	0.18	MLT 7	80	1	
Macrococcus bovicus		1.91	0.14		99	99	
010050	**Good identification**						**010050**
					OXI	YEP	
Macrococcus bovicus		90.17	0.55	ORN 1	99	99	
Macrococcus equipersicus		8.78	0.36		99	99	
Staphylococcus cohnii subsp. cohnii		0.62	0.17		1	12	
010100	**Low discrimination**						**010100**
					OXI		
Dermacoccus nishinomiyaensis		78.21	0.44	ORN 1	93		
Micrococcus sp.		11.68	0.18		89		
Kocuria varians		8.88	0.32		13		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
010110	**Acceptable identification**						**010110**
					YEP		
Macrococcus bovicus		90.31	0.40	ORN 1	99		
Macrococcus equipersicus		7.59	0.20		99		
Dermacoccus nishinomiyaensis		2.10	0.03		1		
010150	**Acceptable identification**						**010150**
Macrococcus bovicus		91.13	0.48	ORN 1			
Macrococcus equipersicus		8.88	0.28				
010200	**Low discrimination**						**010200**
Micrococcus sp.		76.56	0.38	ORN 1			
Dermacoccus nishinomiyaensis		17.55	0.34	SUC 9			
Kocuria rosea		3.91	0.20				
010210	**Low discrimination**						**010210**
					NOV		
Macrococcus bovicus		64.31	0.35	ORN 1	99		
Staphylococcus vitulinus		24.07	0.21		99		
Micrococcus sp.		3.75	-.03		1		
010211	**Good identification**						**010211**
Staphylococcus capitis subsp. capitis		100.0	0.29	ORN 1			
010230	**Good identification**						**010230**
					NOV		
Staphylococcus vitulinus		98.63	0.29	ORN 1	99		
Kocuria rosea		1.37	-.08	NIT 91	1		
010250	**Good identification**						**010250**
Macrococcus bovicus		96.29	0.43	ORN 1			
Macrococcus equipersicus		3.71	0.15				
010310	**Acceptable identification**						**010310**
Macrococcus bovicus		91.78	0.28	ORN 1			
Macrococcus carouselicus		5.17	-.02				
Macrococcus equipersicus		3.05	-.01				
010350	**Good identification**						**010350**
Macrococcus bovicus		96.29	0.35	ORN 1			
Macrococcus equipersicus		3.71	0.08				
010400	**Low discrimination**						**010400**
					NOV	OXI	
Staphylococcus cohnii subsp. cohnii		44.12	0.46	ORN 1	99	1	
Macrococcus carouselicus		19.16	0.28		99	99	
Micrococcus sp.		16.63	0.18		1	89	
010401	**Good identification**						**010401**
					NOV		
Staphylococcus cohnii subsp. cohnii		98.34	0.47	ORN 1	99		
Staphylococcus auricularis		1.66	0.07		1		
010410	**Low discrimination**						**010410**
					OXI	YEP	
Macrococcus bovicus		68.48	0.51	ORN 1	99	99	
Staphylococcus cohnii subsp. cohnii		22.73	0.48		1	12	

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Micrococcus carouselicus		4.96	0.24		99	67			
010411	**Good identification**								**010411**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		97.27	0.49	ORN 1	1	12			
Micrococcus bovicus		2.73	0.10		99	99			
010413	**Low discrimination**								**010413**
					NOV	PYR			
Aerococcus viridans		69.72	0.27	ORN 1	1	95			
Staphylococcus cohnii subsp. cohnii		30.28	0.08	ESL 90	99	1			
010433	**Good identification**								**010433**
Aerococcus viridans		100.0	0.27	ORN 1					
010440	**Low discrimination**								**010440**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		85.07	0.56	ORN 1	1	99	12		
Staphylococcus auricularis		10.30	0.34		1	1	1		
Micrococcus bovicus		2.43	0.18		99	99	99		
010441	**Very good identification**								**010441**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		98.35	0.57	ORN 1	1	99			
Staphylococcus auricularis		1.22	0.14		1	1			
Kocuria kristinae		0.43	-.02		99	1			
010443	**Low discrimination**								**010443**
					NOV	PYR			
Aerococcus viridans		55.23	0.30	ORN 1	1	95			
Staphylococcus cohnii subsp. cohnii		44.77	0.16	ESL 90	99	1			
010450	**Low discrimination**								**010450**
					OXI	YEP			
Micrococcus bovicus		65.99	0.59	ORN 1	99	99			
Staphylococcus cohnii subsp. cohnii		29.73	0.58		1	12			
Micrococcus equipersicus		4.29	0.36		99	99			
010451	**Very good identification**								**010451**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		97.66	0.59	ORN 1	1	99	1	12	
Micrococcus bovicus		2.02	0.18		99	99	1	99	
Aerococcus viridans		0.32	0.20		1	1	95	1	
010453	**Low discrimination**								**010453**
					NOV	PYR			
Aerococcus viridans		74.41	0.39	ORN 1	1	95			
Staphylococcus cohnii subsp. cohnii		25.59	0.18	ESL 90	99	1			
010463	**Good identification**								**010463**
Aerococcus viridans		100.0	0.30	ORN 1					
010472	**Good identification**								**010472**
Aerococcus viridans		100.0	0.32	ORN 1					
010473	**Good identification**								**010473**
Aerococcus viridans		100.0	0.39	ORN 1					
010501	**Good identification**								**010501**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	ORN 1					

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
010510	**Low discrimination**								**010510**
					OXI	YEP			
Macrococcus bovicus		82.34	0.44	ORN 1	99	99			
Staphylococcus cohnii subsp. cohnii		7.09	0.28		1	12			
Macrococcus carouselicus		5.96	0.16		99	67			
010511	**Acceptable identification**								**010511**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		90.22	0.29	ORN 1	1	12			
Macrococcus bovicus		9.78	0.03	GAL 10	99	99			
010513	**Good identification**								**010513**
Aerococcus viridans		100.0	0.27	ORN 1					
010533	**Good identification**								**010533**
Aerococcus viridans		100.0	0.27	ORN 1					
010540	**Low discrimination**								**010540**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		78.84	0.36	ORN 1	1	99	12		
Staphylococcus auricularis		9.54	0.14	GAL 10	1	1	1		
Macrococcus bovicus		8.68	0.10		99	99	99		
010541	**Good identification**								**010541**
Staphylococcus cohnii subsp. cohnii		100.0	0.37	ORN 1					
010543	**Good identification**								**010543**
Aerococcus viridans		100.0	0.30	ORN 1					
010550	**Low discrimination**								**010550**
					OXI	YEP			
Macrococcus bovicus		84.44	0.51	ORN 1	99	99			
Staphylococcus cohnii subsp. cohnii		9.86	0.39		1	12			
Macrococcus equipersicus		5.48	0.28		99	99			
010551	**Acceptable identification**								**010551**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		90.15	0.39	ORN 1	1	99	1	12	
Macrococcus bovicus		7.20	0.10	GAL 10	99	99	1	99	
Aerococcus viridans		2.65	0.20		1	1	95	1	
010552	**Low discrimination**								**010552**
					OXI	NOV	YEP		
Aerococcus viridans		52.80	0.32	ORN 1	1	1	1		
Macrococcus bovicus		37.21	0.10	ESL 90	99	99	99		
Staphylococcus kloosii		9.99	0.18		1	99	99		
010553	**Good identification**								**010553**
Aerococcus viridans		100.0	0.39	ORN 1					
010563	**Good identification**								**010563**
Aerococcus viridans		100.0	0.30	ORN 1					
010572	**Good identification**								**010572**
Aerococcus viridans		100.0	0.32	ORN 1					
010573	**Good identification**								**010573**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.39	ORN 1			
010600	**Low discrimination**						**010600**
					NOV	OXI	
Macrococcus carouselicus		79.18	0.34	ORN 1	99	99	
Staphylococcus auricularis		13.62	0.26	ESL 94	1	1	
Micrococcus sp.		3.35	-.03		1	89	
010610	**Low discrimination**						**010610**
Macrococcus bovicus		56.70	0.39	ORN 1			
Macrococcus carouselicus		33.33	0.30				
Staphylococcus vitulinus		7.96	0.15				
010613	**Good identification**						**010613**
Aerococcus viridans		100.0	0.27	ORN 1			
010633	**Good identification**						**010633**
Aerococcus viridans		100.0	0.27	ORN 1			
010640	**Low discrimination**						**010640**
					NOV	PYR	
Staphylococcus petrasii subsp. pragensis		84.36	0.33	VPT 95	1	99	
Staphylococcus auricularis		13.07	0.33	ORN 1	1	40	
Staphylococcus cohnii subsp. cohnii		1.18	0.15		99	1	
010641	**Acceptable identification**						**010641**
					OXI	NOV	
Kocuria kristinae		93.91	0.39	VPT 90	99	1	
Staphylococcus auricularis		2.47	0.13	ORN 1	1	1	
Staphylococcus cohnii subsp. cohnii		2.18	0.16		1	99	
010643	**Low discrimination**						**010643**
					OXI		
Kocuria kristinae		75.49	0.27	VPT 90	99		
Staphylococcus lugdunensis		15.88	0.14	ORN 1	1		
Aerococcus viridans		8.63	0.30		1		
010650	**Acceptable identification**						**010650**
					OXI	YEP	
Macrococcus bovicus		94.86	0.47	ORN 1	99	99	
Macrococcus equipersicus		2.44	0.15		99	99	
Staphylococcus cohnii subsp. cohnii		1.73	0.17		1	12	
010652	**Low discrimination**						**010652**
					OXI	NOV	
Aerococcus viridans		70.87	0.32	ORN 1	1	1	
Macrococcus bovicus		29.13	0.05	ESL 90	99	99	
010653	**Low discrimination**						**010653**
					OXI		
Aerococcus viridans		89.16	0.39	ORN 1	1		
Kocuria kristinae		10.84	-.02	ESL 90	99		
010663	**Good identification**						**010663**
Aerococcus viridans		100.0	0.30	ORN 1			
010672	**Low discrimination**						**010672**
					NOV	YEP	
Aerococcus viridans		64.70	0.32	ORN 1	1	1	
Staphylococcus arlettae		35.30	0.15	ESL 90	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
010673	**Low discrimination**						**010673**
					NOV	YEP	
Aerococcus viridans		76.34	0.39	ORN 1	1	1	
Staphylococcus arlettae		23.66	0.18	ESL 90	99	99	
010700	**Good identification**						**010700**
					OXI	NOV	
Macrococcus carouselicus		95.73	0.27	ORN 1	99	99	
Staphylococcus auricularis		4.27	0.06	ESL 94	1	1	
010710	**Low discrimination**						**010710**
Macrococcus bovicus		62.11	0.31	ORN 1			
Macrococcus carouselicus		36.51	0.23				
Macrococcus equipersicus		1.38	-.01				
010713	**Good identification**						**010713**
Aerococcus viridans		100.0	0.27	ORN 1			
010733	**Good identification**						**010733**
Aerococcus viridans		100.0	0.27	ORN 1			
010741	**Good identification**						**010741**
Kocuria kristinae		100.0	0.30	VPT 90			
010750	**Good identification**						**010750**
Macrococcus bovicus		97.50	0.39	ORN 1			
Macrococcus equipersicus		2.50	0.08				
010752	**Low discrimination**						**010752**
					OXI	NOV	
Aerococcus viridans		85.02	0.32	ORN 1	1	1	
Macrococcus bovicus		14.98	-.02	ESL 90	99	99	
010753	**Good identification**						**010753**
Aerococcus viridans		100.0	0.39	ORN 1			
010763	**Good identification**						**010763**
Aerococcus viridans		100.0	0.30	ORN 1			
010772	**Low discrimination**						**010772**
					NOV	YEP	
Staphylococcus arlettae		75.56	0.31	ORN 1	99	99	
Aerococcus viridans		24.44	0.32	GLR 92	1	1	
010773	**Low discrimination**						**010773**
					NOV	YEP	
Staphylococcus arlettae		63.72	0.34	ORN 1	99	99	
Aerococcus viridans		36.28	0.39	GLR 92	1	1	
011000	**Low discrimination**						**011000**
					NOV		
Macrococcus carouselicus		44.03	0.28	ORN 1	99		
Micrococcus sp.		38.22	0.18	TRE 94	1		
Dermacoccus nishinomiyaensis		13.58	0.18		80		
011010	**Low discrimination**						**011010**
Macrococcus equipersicus		44.84	0.34	URE 86			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Micrococcus bovicus		27.90	0.28	ORN 1		
Micrococcus carouselicus		27.26	0.24			
011050	**Low discrimination**					**011050**
Micrococcus equipersicus		65.07	0.43	URE 86		
Micrococcus bovicus		34.93	0.36	ORN 1		
011053	**Good identification**					**011053**
Aerococcus viridans		100.0	0.33	ORN 1		
011073	**Good identification**					**011073**
Aerococcus viridans		100.0	0.33	ORN 1		
011150	**Low discrimination**					**011150**
Micrococcus equipersicus		65.07	0.35	URE 86		
Micrococcus bovicus		34.93	0.28	ORN 1		
011153	**Good identification**					**011153**
Aerococcus viridans		100.0	0.33	ORN 1		
011173	**Good identification**					**011173**
Aerococcus viridans		100.0	0.33	ORN 1		
011200	**Good identification**					**011200**
					NOV	
Micrococcus carouselicus		95.94	0.34	ORN 1	99	
Micrococcus sp.		4.06	-.03	TRE 94	1	
011210	**Low discrimination**					**011210**
Staphylococcus vitulinus		48.75	0.29	ORN 1		
Micrococcus carouselicus		42.49	0.30	NIT 91		
Micrococcus bovicus		5.35	0.16			
011230	**Good identification**					**011230**
Staphylococcus vitulinus		100.0	0.38	ORN 1		
011253	**Good identification**					**011253**
Aerococcus viridans		100.0	0.33	ORN 1		
011273	**Good identification**					**011273**
Aerococcus viridans		100.0	0.33	ORN 1		
011300	**Good identification**					**011300**
Micrococcus carouselicus		100.0	0.27	ORN 1		
011353	**Good identification**					**011353**
Aerococcus viridans		100.0	0.33	ORN 1		
011373	**Good identification**					**011373**
Aerococcus viridans		100.0	0.33	ORN 1		
011400	**Good identification**					**011400**
					OXI YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Macrococcus carouselicus		99.50	0.53	ORN 1	99	67			
Macrococcus equipersicus		0.35	0.08		99	99			
Staphylococcus cohnii subsp. cohnii		0.15	0.05		1	12			
011403	**Good identification**								**011403**
Aerococcus viridans		100.0	0.37	ORN 1					
011410	**Low discrimination**								**011410**
Macrococcus carouselicus		82.71	0.49	ORN 1					
Macrococcus equipersicus		8.68	0.34						
Macrococcus bovicus		8.10	0.32						
011412	**Low discrimination**								**011412**
					OXI	NOV			
Aerococcus viridans		73.79	0.39	ORN 1	1	1			
Macrococcus carouselicus		26.21	0.08		99	99			
011413	**Good identification**								**011413**
Aerococcus viridans		100.0	0.47	ORN 1					
011422	**Good identification**								**011422**
Aerococcus viridans		100.0	0.29	ORN 1					
011423	**Good identification**								**011423**
Aerococcus viridans		100.0	0.37	ORN 1					
011431	**Good identification**								**011431**
Aerococcus viridans		100.0	0.27	ORN 1					
011432	**Good identification**								**011432**
Aerococcus viridans		100.0	0.39	ORN 1					
011433	**Good identification**								**011433**
Aerococcus viridans		100.0	0.47	ORN 1					
011441	**Low discrimination**								**011441**
					NOV	PYR			
Aerococcus viridans		55.23	0.30	ORN 1	1	95			
Staphylococcus cohnii subsp. cohnii		44.77	0.16	LAC 90	99	1			
011442	**Good identification**								**011442**
Aerococcus viridans		100.0	0.42	ORN 1					
011443	**Good identification**								**011443**
Aerococcus viridans		100.0	0.49	ORN 1					
011450	**Low discrimination**								**011450**
					OXI	NOV			
Macrococcus equipersicus		52.13	0.43	URE 86	99	99			
Macrococcus bovicus		41.98	0.39	ORN 1	99	99			
Aerococcus viridans		2.32	0.32		1	1			
011451	**Low discrimination**								**011451**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		65.82	0.39	ORN 1	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		22.63	0.18	LAC 90	1	99	1	12	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Micrococcus equipersicus		6.40	0.02		99	99	1	99	
011452	**Good identification**								**011452**
					OXI	NOV			
Aerococcus viridans		95.65	0.51	ORN 1	1	1			
Micrococcus equipersicus		2.41	0.02		99	99			
Micrococcus bovicus		1.94	-.02		99	99			
011453	**Very good identification**								**011453**
Aerococcus viridans		100.0	0.59	ORN 1					
011461	**Good identification**								**011461**
Aerococcus viridans		100.0	0.30	ORN 1					
011462	**Good identification**								**011462**
Aerococcus viridans		100.0	0.42	ORN 1					
011463	**Good identification**								**011463**
Aerococcus viridans		100.0	0.49	ORN 1					
011470	**Low discrimination**								**011470**
					OXI	NOV			
Aerococcus viridans		70.94	0.32	ORN 1	1	1			
Micrococcus equipersicus		16.09	0.02	LAC 90	99	99			
Micrococcus bovicus		12.96	-.02		99	99			
011471	**Good identification**								**011471**
Aerococcus viridans		100.0	0.39	ORN 1					
011472	**Good identification**								**011472**
Aerococcus viridans		100.0	0.51	ORN 1					
011473	**Very good identification**								**011473**
Aerococcus viridans		100.0	0.59	ORN 1					
011500	**Good identification**								**011500**
Micrococcus carouselicus		99.65	0.45	ORN 1					
Micrococcus equipersicus		0.35	0.00						
011502	**Low discrimination**								**011502**
					OXI	NOV			
Aerococcus viridans		58.33	0.29	ORN 1	1	1			
Micrococcus carouselicus		41.67	0.04		99	99			
011503	**Good identification**								**011503**
Aerococcus viridans		100.0	0.37	ORN 1					
011510	**Low discrimination**								**011510**
Micrococcus carouselicus		82.62	0.41	ORN 1					
Micrococcus equipersicus		8.67	0.26						
Micrococcus bovicus		8.10	0.24						
011511	**Low discrimination**								**011511**
					OXI	NOV			
Aerococcus viridans		63.01	0.27	ORN 1	1	1			
Micrococcus carouselicus		37.00	0.00	LAC 90	99	99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
011512	**Low discrimination**						**011512**
					OXI	NOV	
Aerococcus viridans		86.79	0.39	ORN 1	1	1	
Macrococcus carouselicus		13.21	0.00		99	99	
011513	**Good identification**						**011513**
Aerococcus viridans		100.0	0.47	ORN 1			
011522	**Good identification**						**011522**
Aerococcus viridans		100.0	0.29	ORN 1			
011523	**Good identification**						**011523**
Aerococcus viridans		100.0	0.37	ORN 1			
011531	**Good identification**						**011531**
Aerococcus viridans		100.0	0.27	ORN 1			
011532	**Good identification**						**011532**
Aerococcus viridans		100.0	0.39	ORN 1			
011533	**Good identification**						**011533**
Aerococcus viridans		100.0	0.47	ORN 1			
011541	**Good identification**						**011541**
Aerococcus viridans		100.0	0.30	ORN 1			
011542	**Good identification**						**011542**
Aerococcus viridans		100.0	0.42	ORN 1			
011543	**Good identification**						**011543**
Aerococcus viridans		100.0	0.49	ORN 1			
011550	**Low discrimination**						**011550**
					OXI	NOV	
Macrococcus equipersicus		50.94	0.35	URE 86	99	99	
Macrococcus bovicus		41.03	0.32	ORN 1	99	99	
Aerococcus viridans		5.29	0.32		1	1	
011551	**Good identification**						**011551**
Aerococcus viridans		100.0	0.39	ORN 1			
011552	**Good identification**						**011552**
					NOV	YEP	
Aerococcus viridans		97.94	0.51	ORN 1	1	1	
Staphylococcus kloosii		2.06	0.18		99	99	
011553	**Very good identification**						**011553**
Aerococcus viridans		100.0	0.59	ORN 1			
011561	**Good identification**						**011561**
Aerococcus viridans		100.0	0.30	ORN 1			
011562	**Good identification**						**011562**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.42	ORN 1			
011563	**Good identification**						**011563**
Aerococcus viridans		100.0	0.49	ORN 1			
011570	**Good identification**						**011570**
Aerococcus viridans		100.0	0.32	ORN 1			
011571	**Good identification**						**011571**
Aerococcus viridans		100.0	0.39	ORN 1			
011572	**Good identification**						**011572**
Aerococcus viridans		100.0	0.51	ORN 1			
011573	**Very good identification**						**011573**
Aerococcus viridans		100.0	0.59	ORN 1			
011600	**Very good identification**						**011600**
Macrococcus carouselicus		100.0	0.59	ORN 1			
011603	**Acceptable identification**						**011603**
					OXI		
Aerococcus viridans		90.25	0.37	ORN 1	1		
Kocuria kristinae		9.75	-.05		99		
011610	**Good identification**						**011610**
Macrococcus carouselicus		94.35	0.55	ORN 1			
Staphylococcus vitulinus		3.89	0.24				
Macrococcus bovicus		1.14	0.19				
011612	**Low discrimination**						**011612**
					OXI	NOV	
Aerococcus viridans		58.10	0.39	ORN 1	1	1	
Macrococcus carouselicus		41.90	0.14		99	99	
011613	**Good identification**						**011613**
Aerococcus viridans		100.0	0.47	ORN 1			
011622	**Good identification**						**011622**
Aerococcus viridans		100.0	0.29	ORN 1			
011623	**Good identification**						**011623**
Aerococcus viridans		100.0	0.37	ORN 1			
011630	**Acceptable identification**						**011630**
					OXI	NOV	
Staphylococcus vitulinus		90.53	0.33	ORN 1	99	99	
Macrococcus carouselicus		8.21	0.14	NIT 91	99	99	
Aerococcus viridans		1.27	0.19		1	1	
011631	**Low discrimination**						**011631**
					OXI	NOV	
Aerococcus viridans		76.34	0.27	ORN 1	1	1	
Staphylococcus vitulinus		23.66	-.08	LAC 90	99	99	
011632	**Acceptable identification**						**011632**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Aerococcus viridans		92.56	0.39	ORN 1	1	1		
Staphylococcus vitulinus		7.44	-.08		99	99		
011633	**Good identification**							**011633**
Aerococcus viridans		100.0	0.47	ORN 1				
011641	**Low discrimination**							**011641**
					OXI			
Kocuria kristinae		89.75	0.27	VPT 90	99			
Aerococcus viridans		10.26	0.30	ORN 1	1			
011642	**Good identification**							**011642**
Aerococcus viridans		100.0	0.42	ORN 1				
011643	**Low discrimination**							**011643**
					OXI			
Aerococcus viridans		80.45	0.49	ORN 1	1			
Kocuria kristinae		19.56	0.15		99			
011650	**Low discrimination**							**011650**
Micrococcus bovicus		48.28	0.27	ORN 1				
Micrococcus equipersicus		23.72	0.22	ESL 10				
Micrococcus carouselicus		17.33	0.14					
011651	**Low discrimination**							**011651**
					OXI	NOV	PYR	
Aerococcus viridans		68.84	0.39	ORN 1	1	1	95	
Staphylococcus sciuri		22.79	0.13	LAC 90	99	99	1	
Kocuria kristinae		8.37	-.02		99	1	93	
011652	**Good identification**							**011652**
Aerococcus viridans		100.0	0.51	ORN 1				
011653	**Very good identification**							**011653**
Aerococcus viridans		100.0	0.59	ORN 1				
011661	**Good identification**							**011661**
Aerococcus viridans		100.0	0.30	ORN 1				
011662	**Good identification**							**011662**
Aerococcus viridans		100.0	0.42	ORN 1				
011663	**Good identification**							**011663**
Aerococcus viridans		100.0	0.49	ORN 1				
011670	**Low discrimination**							**011670**
					OXI	NOV		
Aerococcus viridans		84.69	0.32	ORN 1	1	1		
Staphylococcus vitulinus		15.31	-.08	LAC 90	99	99		
011671	**Low discrimination**							**011671**
					OXI	NOV		
Aerococcus viridans		80.02	0.39	ORN 1	1	1		
Staphylococcus sciuri		19.98	0.11	LAC 90	99	99		
011672	**Acceptable identification**							**011672**
					NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		94.28	0.51	ORN 1	1	1	
Staphylococcus arlettae		5.72	0.15		99	99	
011673	**Very good identification**						**011673**
					NOV	YEP	
Aerococcus viridans		96.67	0.59	ORN 1	1	1	
Staphylococcus arlettae		3.33	0.18		99	99	
011700	**Good identification**						**011700**
Macrococcus carouselicus		100.0	0.51	ORN 1			
011703	**Good identification**						**011703**
Aerococcus viridans		100.0	0.37	ORN 1			
011710	**Good identification**						**011710**
Macrococcus carouselicus		97.96	0.47	ORN 1			
Macrococcus bovicus		1.18	0.12				
Macrococcus equipersicus		0.50	0.06				
011712	**Low discrimination**						**011712**
					OXI	NOV	
Aerococcus viridans		76.39	0.39	ORN 1	1	1	
Macrococcus carouselicus		23.61	0.06		99	99	
011713	**Acceptable identification**						**011713**
					OXI	NOV	
Aerococcus viridans		91.12	0.47	ORN 1	1	1	
Staphylococcus lentus		8.88	0.12		99	99	
011722	**Good identification**						**011722**
Aerococcus viridans		100.0	0.29	ORN 1			
011723	**Good identification**						**011723**
Aerococcus viridans		100.0	0.37	ORN 1			
011731	**Low discrimination**						**011731**
					OXI	NOV	
Aerococcus viridans		65.99	0.27	ORN 1	1	1	
Staphylococcus sciuri		34.01	0.05	LAC 90	99	99	
011732	**Good identification**						**011732**
Aerococcus viridans		100.0	0.39	ORN 1			
011733	**Acceptable identification**						**011733**
					OXI	NOV	
Aerococcus viridans		91.12	0.47	ORN 1	1	1	
Staphylococcus lentus		8.88	0.12		99	99	
011742	**Good identification**						**011742**
Aerococcus viridans		100.0	0.42	ORN 1			
011743	**Acceptable identification**						**011743**
					OXI		
Aerococcus viridans		92.51	0.49	ORN 1	1		
Kocuria kristinae		7.50	0.05		99		
011751	**Low discrimination**						**011751**
					OXI	NOV	
Aerococcus viridans		56.56	0.39	ORN 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		36.34	0.19	LAC 90	99	99		
Staphylococcus lentus		7.10	0.07		99	99		
011752	**Good identification**							**011752**
Aerococcus viridans		100.0	0.51	ORN 1				
011753	**Good identification**							**011753**
					OXI	NOV		
Aerococcus viridans		94.25	0.59	ORN 1	1	1		
Staphylococcus lentus		4.66	0.18		99	99		
Staphylococcus sciuri		1.10	0.03		99	99		
011761	**Good identification**							**011761**
Aerococcus viridans		100.0	0.30	ORN 1				
011762	**Good identification**							**011762**
Aerococcus viridans		100.0	0.42	ORN 1				
011763	**Good identification**							**011763**
Aerococcus viridans		100.0	0.49	ORN 1				
011770	**Good identification**							**011770**
Aerococcus viridans		100.0	0.32	ORN 1				
011771	**Low discrimination**							**011771**
					OXI	NOV		
Aerococcus viridans		62.10	0.39	ORN 1	1	1		
Staphylococcus sciuri		30.11	0.17	LAC 90	99	99		
Staphylococcus lentus		7.79	0.07		99	99		
011772	**Low discrimination**							**011772**
					NOV	YEP		
Aerococcus viridans		74.43	0.51	ORN 1	1	1		
Staphylococcus arlettae		25.57	0.31		99	99		
011773	**Low discrimination**							**011773**
					OXI	NOV	YEP	
Aerococcus viridans		79.78	0.59	ORN 1	1	1	1	
Staphylococcus arlettae		15.57	0.34		1	99	99	
Staphylococcus lentus		3.94	0.18		99	99	1	
012000	**Low discrimination**							**012000**
Kocuria rosea		59.10	0.59	ORN 1				
Dermacoccus nishinomiyaensis		28.88	0.54					
Micrococcus sp.		7.61	0.33					
012001	**Low discrimination**							**012001**
Kocuria rosea		70.31	0.33	ORN 1				
Staphylococcus capitis subsp. capitis		11.23	0.19	MNS 5				
Kocuria varians		7.10	0.21					
012002	**Low discrimination**							**012002**
					OXI			
Kocuria varians		49.96	0.32	ORN 1	13			
Kocuria rosea		29.26	0.18		35			
Dermacoccus nishinomiyaensis		14.30	0.13		93			
012010	**Low discrimination**							**012010**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					YEP	NOV	
Kocuria rosea		79.76	0.39	ORN 1	1	1	
Macrococcus bovicus		13.37	0.28	MAN 10	99	99	
Dermacoccus nishinomiyaensis		3.54	0.13		1	80	
012011	**Good identification**						**012011**
Staphylococcus capitis subsp. capitis		95.02	0.42	ORN 1			
Kocuria rosea		4.98	0.13				
012020	**Acceptable identification**						**012020**
					OXI		
Kocuria rosea		93.93	0.51	ORN 1	35		
Kocuria varians		4.71	0.34		13		
Dermacoccus nishinomiyaensis		1.08	0.13		93		
012030	**Acceptable identification**						**012030**
					NOV		
Kocuria rosea		91.67	0.32	ORN 1	1		
Staphylococcus vitulinus		8.34	0.09	MAN 10	99		
012040	**Low discrimination**						**012040**
Staphylococcus auricularis		34.02	0.32	ORN 1			
Kocuria varians		23.22	0.29	TRE 92			
Kocuria rosea		17.47	0.18				
012050	**Good identification**						**012050**
					NOV	YEP	
Macrococcus bovicus		97.48	0.36	ORN 1	99	99	
Kocuria rosea		2.52	-.02	NIT 10	1	1	
012100	**Low discrimination**						**012100**
					OXI		
Dermacoccus nishinomiyaensis		47.22	0.39	ORN 1	93		
Kocuria varians		46.52	0.47		13		
Kocuria rosea		5.12	0.18		35		
012102	**Good identification**						**012102**
					OXI		
Kocuria varians		96.49	0.35	ORN 1	13		
Dermacoccus nishinomiyaensis		3.51	-.02		93		
012120	**Low discrimination**						**012120**
					OXI		
Kocuria varians		85.95	0.37	ORN 1	13		
Kocuria rosea		11.55	0.10		35		
Dermacoccus nishinomiyaensis		2.51	-.02		93		
012140	**Low discrimination**						**012140**
					OXI		
Kocuria varians		86.56	0.33	ORN 1	13		
Staphylococcus auricularis		9.40	0.12		1		
Dermacoccus nishinomiyaensis		4.04	-.02		93		
012150	**Good identification**						**012150**
Macrococcus bovicus		100.0	0.28	ORN 1			
012200	**Low discrimination**						**012200**
Kocuria rosea		64.32	0.39	ORN 1			
Dermacoccus nishinomiyaensis		18.06	0.29	SUC 10			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Micrococcus sp.		7.37	0.12				
012210	**Low discrimination**						**012210**
					NOV	YEP	
Staphylococcus vitulinus		89.25	0.41	ORN 1	99	75	
Kocuria rosea		6.95	0.20		1	1	
Macrococcus bovicus		2.62	0.16		99	99	
012211	**Good identification**						**012211**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		98.93	0.47	ORN 1	1	1	
Staphylococcus vitulinus		0.76	0.00		99	99	
Kocuria rosea		0.31	-.07		1	35	
012220	**Low discrimination**						**012220**
					NOV	OXI	
Kocuria rosea		86.77	0.32	ORN 1	1	35	
Staphylococcus vitulinus		7.89	0.09	SUC 10	99	99	
Kocuria varians		5.34	0.16		1	13	
012230	**Good identification**						**012230**
					NOV		
Staphylococcus vitulinus		98.78	0.50	ORN 1	99		
Kocuria rosea		1.22	0.12		1		
012240	**Low discrimination**						**012240**
Staphylococcus auricularis		79.65	0.31	ORN 1			
Kocuria varians		8.03	0.12	TRE 92			
Kocuria rosea		4.92	-.02				
012300	**Low discrimination**						**012300**
					OXI		
Kocuria varians		61.20	0.29	ORN 1	13		
Dermacoccus nishinomiyaensis		29.08	0.14	SUC 12	93		
Kocuria rosea		5.49	-.02		35		
012400	**Low discrimination**						**012400**
Staphylococcus auricularis		64.32	0.46	ORN 1			
Staphylococcus carnosus subsp. utilis		17.31	0.18				
Kocuria rosea		6.70	0.18				
012410	**Low discrimination**						**012410**
					OXI	YEP	
Macrococcus bovicus		72.79	0.32	ORN 1	99	99	
Staphylococcus cohnii subsp. cohnii		16.37	0.25	NIT 10	1	12	
Macrococcus carouselicus		3.03	-.01		99	67	
012440	**Good identification**						**012440**
					NOV		
Staphylococcus auricularis		92.90	0.53	ORN 1	1		
Staphylococcus cohnii subsp. cohnii		6.42	0.33		99		
Kocuria varians		0.42	0.06		1		
012441	**Low discrimination**						**012441**
					NOV		
Staphylococcus auricularis		59.75	0.34	ORN 1	1		
Staphylococcus cohnii subsp. cohnii		40.25	0.34	MNS 10	99		
012442	**Good identification**						**012442**
Staphylococcus auricularis		100.0	0.34	ORN 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
012450	**Low discrimination**							**012450**
					OXI	NOV	YEP	
Macrococcus bovicus		74.62	0.39	ORN 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		22.77	0.35	NIT 10	1	99	12	
Staphylococcus auricularis		2.62	0.12		1	1	1	
012451	**Good identification**							**012451**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		97.04	0.36	ORN 1	1	12		
Macrococcus bovicus		2.97	-.02	NIT 7	99	99		
012540	**Low discrimination**							**012540**
					OXI	NOV	YEP	
Staphylococcus auricularis		67.46	0.34	ORN 1	1	1	1	
Macrococcus caseolyticus		23.82	0.14	GAL 10	99	99	99	
Staphylococcus cohnii subsp. cohnii		4.66	0.13		1	99	12	
012550	**Acceptable identification**							**012550**
					OXI	PYR	YEP	
Macrococcus bovicus		90.20	0.32	ORN 1	99	1	99	
Staphylococcus cohnii subsp. cohnii		7.14	0.15	NIT 10	1	1	12	
Staphylococcus saprophyticus subsp. bovis		2.66	0.02		1	99	80	
012600	**Acceptable identification**							**012600**
					NOV	OXI		
Staphylococcus auricularis		93.38	0.45	ORN 1	1	1		
Macrococcus carouselicus		3.85	0.10		99	99		
Kocuria rosea		1.17	-.02		1	35		
012610	**Low discrimination**							**012610**
Staphylococcus vitulinus		89.70	0.36	ORN 1				
Macrococcus bovicus		7.02	0.19					
Macrococcus carouselicus		2.37	0.06					
012630	**Good identification**							**012630**
Staphylococcus vitulinus		100.0	0.45	ORN 1				
012640	**Low discrimination**							**012640**
Staphylococcus auricularis		89.83	0.53	ORN 1				
Staphylococcus petrasii subsp. pragensis		7.16	0.13					
Staphylococcus haemolyticus		2.02	0.20					
012641	**Low discrimination**							**012641**
					OXI			
Staphylococcus lugdunensis		64.99	0.36	VPT 99	1			
Staphylococcus auricularis		26.56	0.33	LAC 90	1			
Kocuria kristinae		8.45	0.16		99			
012642	**Low discrimination**							**012642**
					PYR			
Staphylococcus auricularis		68.84	0.33	ORN 1	40			
Staphylococcus lugdunensis		15.31	0.14	LAC 10	96			
Staphylococcus hominis subsp. hominis		8.02	0.16		1			
012643	**Very good identification**							**012643**
					OXI			
Staphylococcus lugdunensis		99.14	0.55	VPT 99	1			
Staphylococcus auricularis		0.50	0.13		1			
Kocuria kristinae		0.36	0.04		99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
012650	**Low discrimination**						**012650**
					OXI	NOV	
Micrococcus bovicus		71.21	0.27	ORN 1	99	99	
Staphylococcus haemolyticus		12.04	0.15	NIT 10	1	2	
Staphylococcus auricularis		9.22	0.12		1	1	
012740	**Low discrimination**						**012740**
					OXI	NOV	
Staphylococcus auricularis		58.67	0.33	ORN 1	1	1	
Micrococcus caseolyticus		14.96	0.11	GAL 10	99	99	
Staphylococcus haemolyticus		13.38	0.21		1	2	
012741	**Low discrimination**						**012741**
Staphylococcus lugdunensis		81.41	0.36	VPT 99			
Rothia mucilaginosa		11.36	0.13	LAC 90			
Staphylococcus auricularis		3.70	0.13				
012743	**Very good identification**						**012743**
					OXI		
Staphylococcus lugdunensis		99.88	0.55	VPT 99	1		
Kocuria kristinae		0.12	-.06		99		
013210	**Good identification**						**013210**
Staphylococcus vitulinus		99.45	0.50	ORN 1			
Micrococcus carouselicus		0.55	0.06				
013230	**Very good identification**						**013230**
Staphylococcus vitulinus		100.0	0.59	ORN 1			
013400	**Good identification**						**013400**
					OXI	NOV	
Micrococcus carouselicus		96.68	0.28	ORN 1	99	99	
Staphylococcus auricularis		3.33	0.05	NIT 6	1	1	
013600	**Acceptable identification**						**013600**
					OXI	NOV	
Micrococcus carouselicus		94.54	0.34	ORN 1	99	99	
Staphylococcus vitulinus		3.98	0.04	NIT 6	99	99	
Staphylococcus auricularis		1.48	0.04		1	1	
013610	**Low discrimination**						**013610**
Staphylococcus vitulinus		85.10	0.45	ORN 1			
Micrococcus carouselicus		13.04	0.30				
Staphylococcus sciuri		1.58	0.18				
013611	**Acceptable identification**						**013611**
Staphylococcus sciuri		93.70	0.36	ORN 1			
Staphylococcus vitulinus		6.30	0.04				
013630	**Very good identification**						**013630**
Staphylococcus vitulinus		99.48	0.54	ORN 1			
Staphylococcus sciuri		0.52	0.15				
013631	**Low discrimination**						**013631**
Staphylococcus sciuri		80.59	0.34	ORN 1			
Staphylococcus vitulinus		19.41	0.13				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
013650	**Low discrimination**						**013650**
Staphylococcus sciuri		79.88	0.30	ORN 1			
Staphylococcus vitulinus		11.54	0.04	MNS 89			
Macrococcus bovicus		8.58	0.07				
013651	**Good identification**						**013651**
Staphylococcus sciuri		99.23	0.48	ORN 1			
Staphylococcus lentus		0.77	0.07				
013653	**Low discrimination**						**013653**
					OXI	NOV	
Staphylococcus sciuri		83.13	0.32	ORN 1	99	99	
Staphylococcus lentus		14.01	0.18	LAC 14	99	99	
Aerococcus viridans		2.86	0.18		1	1	
013670	**Low discrimination**						**013670**
Staphylococcus sciuri		65.88	0.27	ORN 1			
Staphylococcus vitulinus		34.12	0.13	MNS 89			
013671	**Good identification**						**013671**
Staphylococcus sciuri		98.99	0.46	ORN 1			
Staphylococcus lentus		1.02	0.07				
013673	**Low discrimination**						**013673**
					OXI	NOV	
Staphylococcus sciuri		78.80	0.30	ORN 1	99	99	
Staphylococcus lentus		17.60	0.18	LAC 14	99	99	
Aerococcus viridans		3.60	0.18		1	1	
013700	**Low discrimination**						**013700**
					OXI	NOV	
Macrococcus carouselicus		70.86	0.27	ORN 1	99	99	
Rothia mucilaginosa		29.14	0.15	NIT 6	1	1	
013711	**Low discrimination**						**013711**
Staphylococcus sciuri		57.77	0.42	ORN 1			
Staphylococcus lentus		42.23	0.41				
013713	**Acceptable identification**						**013713**
Staphylococcus lentus		94.09	0.53	ORN 1			
Staphylococcus sciuri		5.91	0.26				
013731	**Low discrimination**						**013731**
Staphylococcus sciuri		50.78	0.40	ORN 1			
Staphylococcus lentus		49.22	0.41				
013733	**Good identification**						**013733**
Staphylococcus lentus		95.48	0.53	ORN 1			
Staphylococcus sciuri		4.52	0.24				
013741	**Low discrimination**						**013741**
					OXI	NOV	
Rothia mucilaginosa		87.85	0.33	VPT 95	1	1	
Staphylococcus sciuri		10.09	0.23	ORN 1	99	99	
Staphylococcus lentus		1.30	0.07		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
013750	**Acceptable identification**						**013750**
					YEP		
Staphylococcus sciuri		94.70	0.36	ORN 1	57		
Staphylococcus lentus		3.05	0.07	MNS 89	1		
Macrococcus bovicus		2.25	0.00		99		
013751	**Low discrimination**						**013751**
					OXI	NOV	
Staphylococcus sciuri		71.57	0.54	ORN 1	99	99	
Staphylococcus lentus		28.24	0.48		99	99	
Rothia mucilaginosa		0.20	-.09		1	1	
013753	**Low discrimination**						**013753**
					OXI	NOV	
Staphylococcus lentus		89.41	0.59	ORN 1	99	99	
Staphylococcus sciuri		10.40	0.38		99	99	
Aerococcus viridans		0.19	0.18		1	1	
013770	**Good identification**						**013770**
Staphylococcus sciuri		95.90	0.33	ORN 1			
Staphylococcus lentus		4.10	0.07	MNS 89			
013771	**Low discrimination**						**013771**
Staphylococcus sciuri		65.66	0.52	ORN 1			
Staphylococcus lentus		34.34	0.48				
013773	**Good identification**						**013773**
					OXI	NOV	
Staphylococcus lentus		91.76	0.59	ORN 1	99	99	
Staphylococcus sciuri		8.06	0.35		99	99	
Aerococcus viridans		0.19	0.18		1	1	
014000	**Low discrimination**						**014000**
Dermacoccus nishinomiyaensis		63.90	0.33	ORN 1			
Micrococcus sp.		34.53	0.18	PHS 5			
Kocuria rosea		1.57	-.02				
014240	**Very good identification**						**014240**
Staphylococcus aureus subsp. anaerobius		100.0	0.57	ORN 1			
014410	**Low discrimination**						**014410**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		66.12	0.27	ORN 1	1	12	
Macrococcus bovicus		20.35	0.10	PHS 9	99	99	
Staphylococcus kloosii		13.53	0.26		1	99	
014411	**Good identification**						**014411**
Staphylococcus cohnii subsp. cohnii		100.0	0.28	ORN 1			
014440	**Good identification**						**014440**
Staphylococcus cohnii subsp. cohnii		100.0	0.35	ORN 1			
014441	**Good identification**						**014441**
Staphylococcus cohnii subsp. cohnii		100.0	0.36	ORN 1			
014450	**Low discrimination**						**014450**
					OXI	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		50.02	0.52	ORN 1	1	99	
Staphylococcus cohnii subsp. cohnii		40.74	0.38		1	12	
Macrococcus bovicus		9.24	0.18		99	99	
014451	**Good identification**						**014451**
Staphylococcus cohnii subsp. cohnii		100.0	0.38	ORN 1			
014452	**Good identification**						**014452**
Staphylococcus kloosii		100.0	0.53	ORN 1			
014470	**Good identification**						**014470**
Staphylococcus kloosii		100.0	0.40	ORN 1			
014472	**Good identification**						**014472**
Staphylococcus kloosii		100.0	0.40	ORN 1			
014510	**Low discrimination**						**014510**
					OXI	YEP	
Staphylococcus kloosii		62.05	0.32	ORN 1	1	99	
Macrococcus bovicus		20.60	0.03	MLT 95	99	99	
Staphylococcus cohnii subsp. cohnii		17.36	0.08		1	12	
014512	**Good identification**						**014512**
Staphylococcus kloosii		100.0	0.32	ORN 1			
014550	**Good identification**						**014550**
					OXI	YEP	
Staphylococcus kloosii		91.96	0.58	ORN 1	1	99	
Staphylococcus cohnii subsp. cohnii		4.29	0.18		1	12	
Macrococcus bovicus		3.75	0.10		99	99	
014552	**Very good identification**						**014552**
Staphylococcus kloosii		100.0	0.59	ORN 1			
014570	**Good identification**						**014570**
Staphylococcus kloosii		100.0	0.46	ORN 1			
014572	**Good identification**						**014572**
Staphylococcus kloosii		100.0	0.46	ORN 1			
014650	**Low discrimination**						**014650**
					OXI		
Staphylococcus kloosii		68.17	0.31	ORN 1	1		
Macrococcus bovicus		31.83	0.05	SUC 9	99		
014652	**Good identification**						**014652**
Staphylococcus kloosii		100.0	0.32	ORN 1			
014750	**Acceptable identification**						**014750**
					OXI		
Staphylococcus kloosii		90.66	0.37	ORN 1	1		
Macrococcus bovicus		9.34	-.02	SUC 9	99		
014752	**Good identification**						**014752**
Staphylococcus kloosii		100.0	0.38	ORN 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
014772	**Acceptable identification**					**014772**
Staphylococcus arlettae		90.92	0.35	ORN 1		
Staphylococcus kloosii		9.08	0.26	GLR 92		
014773	**Good identification**					**014773**
Staphylococcus arlettae		100.0	0.37	ORN 1		
015450	**Good identification**					**015450**
					OXI	
Staphylococcus kloosii		95.60	0.52	ORN 1	1	
Macrococcus equipersicus		2.44	0.02		99	
Macrococcus bovicus		1.96	-.02		99	
015452	**Good identification**					**015452**
Staphylococcus kloosii		100.0	0.53	ORN 1		
015470	**Good identification**					**015470**
Staphylococcus kloosii		100.0	0.40	ORN 1		
015472	**Good identification**					**015472**
Staphylococcus kloosii		100.0	0.40	ORN 1		
015510	**Low discrimination**					**015510**
					OXI	
Staphylococcus kloosii		72.65	0.32	ORN 1	1	
Macrococcus carouselicus		27.35	0.00	MLT 95	99	
015512	**Good identification**					**015512**
Staphylococcus kloosii		100.0	0.32	ORN 1		
015550	**Very good identification**					**015550**
Staphylococcus kloosii		100.0	0.58	ORN 1		
015552	**Very good identification**					**015552**
Staphylococcus kloosii		100.0	0.59	ORN 1		
015570	**Good identification**					**015570**
Staphylococcus kloosii		100.0	0.46	ORN 1		
015572	**Good identification**					**015572**
Staphylococcus kloosii		100.0	0.46	ORN 1		
015650	**Low discrimination**					**015650**
					OXI	
Staphylococcus kloosii		84.42	0.31	ORN 1	1	
Staphylococcus sciuri		15.58	-.01	SUC 9	99	
015652	**Good identification**					**015652**
Staphylococcus kloosii		100.0	0.32	ORN 1		
015750	**Low discrimination**					**015750**
					OXI	
Staphylococcus kloosii		84.42	0.37	ORN 1	1	
Staphylococcus sciuri		15.58	0.05	SUC 9	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
015752	**Good identification**						**015752**
Staphylococcus kloosii		100.0	0.38	ORN 1			
015772	**Acceptable identification**						**015772**
Staphylococcus arlettae		90.92	0.35	ORN 1			
Staphylococcus kloosii		9.08	0.26	GLR 92			
015773	**Low discrimination**						**015773**
					OXI	YEP	
Staphylococcus arlettae		88.68	0.37	ORN 1	1	99	
Staphylococcus sciuri		4.52	0.05	GLR 92	99	57	
Staphylococcus lentus		3.74	0.05		99	1	
016000	**Low discrimination**						**016000**
Dermacoccus nishinomiyaensis		68.15	0.28	ORN 1			
Kocuria rosea		26.76	0.18	PHS 5			
Micrococcus sp.		3.45	-.09				
016211	**Good identification**						**016211**
Staphylococcus capitis subsp. capitis		100.0	0.29	ORN 1			
016343	**Good identification**						**016343**
Staphylococcus epidermidis		100.0	0.31	VPT 96			
016620	**Very good identification**						**016620**
Staphylococcus muscae		100.0	0.59	ORN 1			
017351	**Good identification**						**017351**
Staphylococcus sciuri		100.0	0.28	ORN 1			
017611	**Good identification**						**017611**
Staphylococcus sciuri		100.0	0.41	ORN 1			
017631	**Good identification**						**017631**
Staphylococcus sciuri		100.0	0.39	ORN 1			
017650	**Good identification**						**017650**
Staphylococcus sciuri		100.0	0.34	ORN 1			
017651	**Very good identification**						**017651**
Staphylococcus sciuri		100.0	0.53	ORN 1			
017653	**Good identification**						**017653**
Staphylococcus sciuri		97.59	0.37	ORN 1			
Staphylococcus lentus		2.41	0.05	LAC 14			
017670	**Good identification**						**017670**
Staphylococcus sciuri		100.0	0.32	ORN 1			
017671	**Good identification**						**017671**
Staphylococcus sciuri		100.0	0.51	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
017673	**Good identification**						**017673**
Staphylococcus sciuri		96.83	0.34	ORN 1			
Staphylococcus lentus		3.18	0.05	LAC 14			
017710	**Good identification**						**017710**
Staphylococcus sciuri		100.0	0.28	ORN 1			
017711	**Acceptable identification**						**017711**
Staphylococcus sciuri		90.31	0.47	ORN 1			
Staphylococcus lentus		9.69	0.29				
017713	**Low discrimination**						**017713**
Staphylococcus lentus		70.04	0.40	ORN 1			
Staphylococcus sciuri		29.96	0.31				
017731	**Low discrimination**						**017731**
Staphylococcus sciuri		87.54	0.45	ORN 1			
Staphylococcus lentus		12.46	0.29				
017733	**Low discrimination**						**017733**
Staphylococcus lentus		75.61	0.40	ORN 1			
Staphylococcus sciuri		24.40	0.28				
017741	**Good identification**						**017741**
					OXI	NOV	
Staphylococcus sciuri		95.09	0.28	ORN 1	99	99	
Rothia mucilaginosa		4.91	-.09	MAN 97	1	1	
017750	**Good identification**						**017750**
Staphylococcus sciuri		100.0	0.40	ORN 1			
017751	**Good identification**						**017751**
Staphylococcus sciuri		94.52	0.59	ORN 1			
Staphylococcus lentus		5.48	0.35				
017753	**Low discrimination**						**017753**
Staphylococcus lentus		55.79	0.47	ORN 1			
Staphylococcus sciuri		44.21	0.43				
017770	**Good identification**						**017770**
Staphylococcus sciuri		100.0	0.38	ORN 1			
017771	**Low discrimination**						**017771**
					OXI		
Staphylococcus sciuri		89.22	0.56	ORN 1	99		
Staphylococcus lentus		6.85	0.35		99		
Staphylococcus gallinarum		3.93	0.23		1		
017773	**Low discrimination**						**017773**
					OXI		
Staphylococcus lentus		49.96	0.47	ORN 1	99		
Staphylococcus sciuri		29.87	0.40		99		
Staphylococcus gallinarum		19.79	0.31		1		
020000	**Low discrimination**						**020000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Micrococcus sp.		63.37	0.59	BGA 1	89			
Dermacoccus nishinomiyaensis		22.51	0.59		93			
Kocuria varians		10.71	0.60		13			
020002	**Low discrimination**							**020002**
					OXI			
Kocuria varians		75.63	0.49		13			
Micrococcus sp.		16.03	0.18		89			
Dermacoccus nishinomiyaensis		5.69	0.18		93			
020003	**Low discrimination**							**020003**
					PYR			
Kocuria varians		77.24	0.27	MNS 8	10			
Aerococcus viridans		22.76	0.30		95			
020010	**Low discrimination**							**020010**
					NOV			
Macrococcus bovicus		71.72	0.48	BGA 1	99			
Micrococcus sp.		10.57	0.18		1			
Macrococcus equipersicus		6.03	0.27		99			
020012	**Low discrimination**							**020012**
					NOV	PYR	OXI	
Aerococcus viridans		57.23	0.32	ESL 90	1	95	1	
Macrococcus bovicus		25.23	0.07	TRE 95	99	1	99	
Kocuria varians		17.55	0.08		1	10	13	
020013	**Good identification**							**020013**
Aerococcus viridans		100.0	0.40	ESL 90				
020020	**Low discrimination**							**020020**
					OXI			
Kocuria varians		64.16	0.51		13			
Kocuria rosea		21.04	0.32		35			
Micrococcus sp.		10.92	0.18		89			
020021	**Low discrimination**							**020021**
Kocuria varians		83.44	0.29	MNS 8				
Kocuria rosea		16.56	0.05					
020022	**Good identification**							**020022**
					PYR			
Kocuria varians		96.97	0.39		10			
Aerococcus viridans		3.03	0.23		95			
020032	**Good identification**							**020032**
Aerococcus viridans		100.0	0.32	ESL 90				
020033	**Good identification**							**020033**
Aerococcus viridans		100.0	0.40	ESL 90				
020040	**Low discrimination**							**020040**
					OXI			
Micrococcus sp.		56.13	0.36	BGA 1	89			
Kocuria varians		27.66	0.46	MLT 7	13			
Staphylococcus auricularis		10.74	0.37		1			
020042	**Low discrimination**							**020042**
					PYR			

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Kocuria varians		70.14	0.35		10				
Aerococcus viridans		14.04	0.35		95				
Staphylococcus auricularis		10.73	0.17		40				
020043	**Low discrimination**								**020043**
					PYR				
Aerococcus viridans		84.30	0.42	ESL 90	95				
Kocuria varians		15.70	0.13	TRE 95	10				
020050	**Low discrimination**								**020050**
					OXI	YEP			
Macrococcus bovicus		89.63	0.55	BGA 1	99	99			
Macrococcus equipersicus		8.73	0.36		99	99			
Staphylococcus cohnii subsp. cohnii		0.61	0.17		1	12			
020051	**Low discrimination**								**020051**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		36.79	0.33	ESL 90	1	1	95	1	
Macrococcus bovicus		36.49	0.14	TRE 95	99	99	1	99	
Staphylococcus cohnii subsp. cohnii		26.71	0.18	LAC 90	1	99	1	12	
020052	**Low discrimination**								**020052**
					OXI	NOV			
Aerococcus viridans		79.55	0.45	ESL 90	1	1			
Macrococcus bovicus		20.45	0.14	TRE 95	99	99			
020053	**Good identification**								**020053**
Aerococcus viridans		100.0	0.52	ESL 90					
020060	**Acceptable identification**								**020060**
					OXI				
Kocuria varians		94.49	0.37		13				
Micrococcus sp.		5.51	-.05		89				
020063	**Acceptable identification**								**020063**
					PYR				
Aerococcus viridans		93.86	0.42	ESL 90	95				
Kocuria varians		6.14	0.04	TRE 95	10				
020071	**Good identification**								**020071**
Aerococcus viridans		100.0	0.33	ESL 90					
020072	**Good identification**								**020072**
Aerococcus viridans		100.0	0.45	ESL 90					
020073	**Good identification**								**020073**
Aerococcus viridans		100.0	0.52	ESL 90					
020100	**Low discrimination**								**020100**
					OXI				
Kocuria varians		76.12	0.64		13				
Dermacoccus nishinomiyaensis		20.32	0.44		93				
Micrococcus sp.		3.03	0.18		89				
020101	**Good identification**								**020101**
					OXI				
Kocuria varians		96.99	0.42	MNS 8	13				
Dermacoccus nishinomiyaensis		3.01	0.03		93				
020102	**Good identification**								**020102**
					OXI	PYR			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		98.34	0.52		13	10		
Dermacoccus nishinomiyaensis		0.94	0.03		93	45		
Aerococcus viridans		0.72	0.23		1	95		
020103	**Low discrimination**							**020103**
					PYR			
Kocuria varians		83.58	0.31	MNS 8	10			
Aerococcus viridans		16.42	0.30		95			
020110	**Low discrimination**							**020110**
					NOV	OXI		
Macrococcus bovicus		83.72	0.40	BGA 1	99	99		
Kocuria varians		7.30	0.23		1	13		
Macrococcus equipersicus		7.04	0.20		99	99		
020112	**Low discrimination**							**020112**
					NOV	PYR	OXI	
Aerococcus viridans		60.65	0.32	ESL 90	1	95	1	
Kocuria varians		27.89	0.11	TRE 95	1	10	13	
Macrococcus bovicus		11.46	-.01		99	1	99	
020113	**Good identification**							**020113**
Aerococcus viridans		100.0	0.40	ESL 90				
020120	**Very good identification**							**020120**
					OXI			
Kocuria varians		99.24	0.54		13			
Dermacoccus nishinomiyaensis		0.76	0.03		93			
020121	**Good identification**							**020121**
Kocuria varians		100.0	0.32	MNS 8				
020122	**Good identification**							**020122**
					PYR			
Kocuria varians		97.96	0.43		10			
Aerococcus viridans		2.04	0.23		95			
020132	**Low discrimination**							**020132**
					PYR			
Aerococcus viridans		86.09	0.32	ESL 90	95			
Kocuria varians		13.91	0.02	TRE 95	10			
020133	**Good identification**							**020133**
Aerococcus viridans		100.0	0.40	ESL 90				
020140	**Acceptable identification**							**020140**
					OXI			
Kocuria varians		93.16	0.50		13			
Staphylococcus auricularis		2.68	0.17		1			
Micrococcus sp.		1.27	-.05		89			
020141	**Low discrimination**							**020141**
					PYR			
Kocuria varians		89.91	0.28	MNS 8	10			
Aerococcus viridans		10.09	0.23		95			
020142	**Low discrimination**							**020142**
					PYR			
Kocuria varians		88.23	0.39		10			
Aerococcus viridans		11.77	0.35		95			
020143	**Low discrimination**							**020143**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR			
Aerococcus viridans		78.16	0.42	ESL 90	95			
Kocuria varians		21.84	0.17	TRE 95	10			
020150	**Low discrimination**							**020150**
					OXI	NOV		
Macrococcus bovicus		89.32	0.48	BGA 1	99	99		
Macrococcus equipersicus		8.70	0.28		99	99		
Aerococcus viridans		0.91	0.25		1	1		
020151	**Low discrimination**							**020151**
					OXI	NOV		
Aerococcus viridans		70.17	0.33	ESL 90	1	1		
Macrococcus bovicus		29.83	0.07	TRE 95	99	99		
020152	**Low discrimination**							**020152**
					OXI	NOV	YEP	
Aerococcus viridans		86.61	0.45	ESL 90	1	1	1	
Macrococcus bovicus		9.55	0.07	TRE 95	99	99	99	
Staphylococcus kloosii		3.84	0.18		1	99	99	
020153	**Good identification**							**020153**
Aerococcus viridans		100.0	0.52	ESL 90				
020160	**Good identification**							**020160**
Kocuria varians		100.0	0.41					
020162	**Low discrimination**							**020162**
					PYR			
Kocuria varians		72.48	0.29		10			
Aerococcus viridans		27.52	0.35		95			
020163	**Acceptable identification**							**020163**
					PYR			
Aerococcus viridans		91.06	0.42	ESL 90	95			
Kocuria varians		8.94	0.08	TRE 95	10			
020171	**Good identification**							**020171**
Aerococcus viridans		100.0	0.33	ESL 90				
020172	**Good identification**							**020172**
Aerococcus viridans		100.0	0.45	ESL 90				
020173	**Good identification**							**020173**
Aerococcus viridans		100.0	0.52	ESL 90				
020200	**Low discrimination**							**020200**
					OXI			
Micrococcus sp.		63.05	0.38	BGA 1	89			
Kocuria varians		14.69	0.42	SUC 9	13			
Dermacoccus nishinomiyaensis		14.45	0.34		93			
020202	**Low discrimination**							**020202**
					OXI			
Kocuria varians		75.10	0.31	SUC 12	13			
Micrococcus sp.		11.54	-.03		89			
Staphylococcus auricularis		7.31	0.09		1			
020210	**Low discrimination**							**020210**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Micrococcus bovicus		64.31	0.35	BGA 1	99		
Staphylococcus vitulinus		24.07	0.21		99		
Micrococcus sp.		3.75	-.03		1		
020211	**Acceptable identification**						**020211**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		93.49	0.29	BGA 1	1	1	
Aerococcus viridans		3.63	0.20	NIT 89	1	95	
Kocuria kristinae		2.88	-.04		99	93	
020212	**Good identification**						**020212**
Aerococcus viridans		100.0	0.32	ESL 90			
020213	**Good identification**						**020213**
Aerococcus viridans		100.0	0.40	ESL 90			
020220	**Low discrimination**						**020220**
					OXI		
Kocuria varians		71.91	0.33	SUC 12	13		
Kocuria rosea		19.22	0.12		35		
Micrococcus sp.		8.87	-.03		89		
020223	**Good identification**						**020223**
Aerococcus viridans		100.0	0.30	ESL 90			
020230	**Good identification**						**020230**
					NOV		
Staphylococcus vitulinus		98.63	0.29	BGA 1	99		
Kocuria rosea		1.37	-.08	NIT 91	1		
020232	**Good identification**						**020232**
Aerococcus viridans		100.0	0.32	ESL 90			
020233	**Good identification**						**020233**
Aerococcus viridans		100.0	0.40	ESL 90			
020240	**Low discrimination**						**020240**
					COA	OXI	
Staphylococcus auricularis		38.80	0.36	BGA 14	1	1	
Micrococcus sp.		21.73	0.15	NIT 90	1	89	
Staphylococcus aureus subsp. anaerobius		19.66	0.16	TRE 92	99	1	
020241	**Good identification**						**020241**
					OXI		
Kocuria kristinae		96.59	0.44	VPT 90	99		
Staphylococcus auricularis		2.05	0.16	TRE 95	1		
Aerococcus viridans		0.75	0.23		1		
020242	**Low discrimination**						**020242**
					PYR		
Aerococcus viridans		41.90	0.35	ESL 90	95		
Staphylococcus auricularis		29.56	0.16	TRE 95	40		
Kocuria varians		28.55	0.17		10		
020243	**Low discrimination**						**020243**
					OXI		
Kocuria kristinae		78.10	0.32	VPT 90	99		
Aerococcus viridans		21.91	0.42	TRE 95	1		
020250	**Acceptable identification**						**020250**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Micrococcus bovicus		94.70	0.43	BGA 1	99	99		
Micrococcus equipersicus		3.65	0.15		99	99		
Aerococcus viridans		1.65	0.25		1	1		
020252	**Low discrimination**							**020252**
					OXI	NOV	PYR	
Aerococcus viridans		78.22	0.45	ESL 90	1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		16.75	0.03	TRE 95	1	99	7	
Micrococcus bovicus		5.03	0.02		99	99	1	
020253	**Good identification**							**020253**
					OXI			
Aerococcus viridans		95.28	0.52	ESL 90	1			
Kocuria kristinae		4.72	0.03	TRE 95	99			
020262	**Low discrimination**							**020262**
					PYR			
Aerococcus viridans		80.68	0.35	ESL 90	95			
Kocuria varians		19.32	0.08	TRE 95	10			
020263	**Good identification**							**020263**
Aerococcus viridans		100.0	0.42	ESL 90				
020271	**Good identification**							**020271**
Aerococcus viridans		100.0	0.33	ESL 90				
020272	**Low discrimination**							**020272**
					NOV	YEP		
Aerococcus viridans		88.65	0.45	ESL 90	1	1		
Staphylococcus arlettae		11.35	0.15	TRE 95	99	99		
020273	**Acceptable identification**							**020273**
					NOV	YEP		
Aerococcus viridans		93.22	0.52	ESL 90	1	1		
Staphylococcus arlettae		6.78	0.18	TRE 95	99	99		
020300	**Low discrimination**							**020300**
					OXI			
Kocuria varians		83.97	0.46	SUC 12	13			
Dermacoccus nishinomiyaensis		10.49	0.20		93			
Micrococcus sp.		2.43	-.03		89			
020302	**Acceptable identification**							**020302**
					PYR			
Kocuria varians		94.91	0.35	SUC 12	10			
Aerococcus viridans		5.10	0.23		95			
020303	**Low discrimination**							**020303**
					OXI	PYR		
Aerococcus viridans		45.00	0.30	ESL 90	1	95		
Kocuria varians		31.24	0.13	TRE 95	13	10		
Kocuria kristinae		23.77	0.02		99	93		
020310	**Low discrimination**							**020310**
					NOV	OXI		
Micrococcus bovicus		87.94	0.28	BGA 1	99	99		
Micrococcus carouselicus		4.95	-.02		99	99		
Kocuria varians		4.18	0.05		1	13		
020312	**Good identification**							**020312**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.32	ESL 90			
020313	**Acceptable identification**						**020313**
					COA		
Aerococcus viridans		92.13	0.40	ESL 90	1		
Staphylococcus intermedius		7.87	-.01	TRE 95	99		
020320	**Good identification**						**020320**
Kocuria varians		100.0	0.36	SUC 12			
020323	**Low discrimination**						**020323**
					PYR		
Aerococcus viridans		80.39	0.30	ESL 90	95		
Kocuria varians		19.61	0.03	TRE 95	10		
020332	**Good identification**						**020332**
Aerococcus viridans		100.0	0.32	ESL 90			
020333	**Good identification**						**020333**
Aerococcus viridans		100.0	0.40	ESL 90			
020340	**Low discrimination**						**020340**
Kocuria varians		83.71	0.32	SUC 12			
Staphylococcus auricularis		16.29	0.16				
020341	**Good identification**						**020341**
					OXI	PYR	
Kocuria kristinae		95.08	0.34	VPT 90	99	93	
Kocuria varians		2.70	0.10	TRE 95	13	10	
Aerococcus viridans		2.22	0.23		1	95	
020350	**Acceptable identification**						**020350**
					OXI	NOV	
Macrococcus bovicus		92.66	0.35	BGA 1	99	99	
Aerococcus viridans		3.78	0.25		1	1	
Macrococcus equipersicus		3.57	0.08		99	99	
020351	**Low discrimination**						**020351**
					OXI		
Aerococcus viridans		62.73	0.33	ESL 90	1		
Kocuria kristinae		37.27	0.06	TRE 95	99		
020352	**Good identification**						**020352**
Aerococcus viridans		100.0	0.45	ESL 90			
020353	**Good identification**						**020353**
					OXI	COA	
Aerococcus viridans		95.37	0.52	ESL 90	1	1	
Staphylococcus intermedius		3.06	0.03	TRE 95	1	99	
Kocuria kristinae		1.57	-.06		99	1	
020362	**Low discrimination**						**020362**
					PYR		
Aerococcus viridans		73.58	0.35	ESL 90	95		
Kocuria varians		26.43	0.12	TRE 95	10		
020363	**Good identification**						**020363**
Aerococcus viridans		100.0	0.42	ESL 90			
020371	**Good identification**						**020371**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Aerococcus viridans		100.0	0.33	ESL 90					
020372	**Low discrimination**								**020372**
					NOV	YEP			
Aerococcus viridans		57.96	0.45	ESL 90	1	1			
Staphylococcus arlettae		42.04	0.31	TRE 95	99	99			
020373	**Low discrimination**								**020373**
					NOV	YEP			
Aerococcus viridans		70.82	0.52	ESL 90	1	1			
Staphylococcus arlettae		29.18	0.34	TRE 95	99	99			
020400	**Low discrimination**								**020400**
					NOV				
Staphylococcus auricularis		50.42	0.51	BGA 14	1				
Staphylococcus cohnii subsp. cohnii		19.04	0.46	NIT 90	99				
Kocuria varians		9.04	0.37		1				
020401	**Low discrimination**								**020401**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		68.99	0.47	BGA 1	99	1			
Staphylococcus auricularis		18.73	0.31		1	40			
Aerococcus viridans		6.13	0.37		1	95			
020402	**Low discrimination**								**020402**
					PYR				
Aerococcus viridans		45.85	0.49	ESL 90	95				
Staphylococcus auricularis		36.36	0.31		40				
Kocuria varians		16.54	0.26		10				
020403	**Good identification**								**020403**
					OXI				
Aerococcus viridans		92.61	0.56	ESL 90	1				
Staphylococcus auricularis		3.50	0.12		1				
Kocuria kristinae		1.48	-.03		99				
020410	**Low discrimination**								**020410**
					OXI	YEP			
Macrococcus bovicus		66.39	0.51	BGA 1	99	99			
Staphylococcus cohnii subsp. cohnii		22.04	0.48		1	12			
Macrococcus carouselicus		4.81	0.24		99	67			
020411	**Low discrimination**								**020411**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		80.82	0.49	BGA 1	1	99	1	12	
Aerococcus viridans		16.91	0.47		1	1	95	1	
Macrococcus bovicus		2.27	0.10		99	99	1	99	
020412	**Good identification**								**020412**
					OXI	NOV	YEP		
Aerococcus viridans		93.33	0.59	ESL 90	1	1	1		
Macrococcus bovicus		3.25	0.10		99	99	99		
Staphylococcus kloosii		2.34	0.26		1	99	99		
020413	**Very good identification**								**020413**
					NOV	PYR			
Aerococcus viridans		99.47	0.66	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		0.53	0.08		99	1			
020420	**Low discrimination**								**020420**
					PYR				
Kocuria varians		68.11	0.28	TRE 7	10				

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Aerococcus viridans		16.84	0.29		95				
Staphylococcus auricularis		10.93	0.10		40				
020421	**Low discrimination**								**020421**
					NOV	PYR			
Aerococcus viridans		79.08	0.37	ESL 90	1	95			
Kocuria varians		11.92	0.06	LAC 90	1	10			
Staphylococcus cohnii subsp. cohnii		9.00	0.05		99	1			
020422	**Low discrimination**								**020422**
					PYR				
Aerococcus viridans		88.75	0.49	ESL 90	95				
Kocuria varians		11.25	0.16		10				
020423	**Very good identification**								**020423**
Aerococcus viridans		100.0	0.56	ESL 90					
020430	**Low discrimination**								**020430**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		70.56	0.39	ESL 90	1	1	95	1	
Macrococcus bovicus		22.10	0.10	LAC 90	99	99	1	99	
Staphylococcus cohnii subsp. cohnii		7.34	0.07		1	99	1	12	
020431	**Good identification**								**020431**
					NOV	PYR			
Aerococcus viridans		95.40	0.47	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		4.60	0.08	LAC 90	99	1			
020432	**Very good identification**								**020432**
Aerococcus viridans		100.0	0.59	ESL 90					
020433	**Very good identification**								**020433**
Aerococcus viridans		100.0	0.66	ESL 90					
020440	**Low discrimination**								**020440**
					NOV	PYR			
Staphylococcus auricularis		63.13	0.59	BGA 14	1	40			
Staphylococcus cohnii subsp. cohnii		32.36	0.56	NIT 90	99	1			
Aerococcus viridans		1.69	0.42		1	95			
020441	**Low discrimination**								**020441**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		68.48	0.57	BGA 1	1	99			
Staphylococcus auricularis		13.70	0.39		1	1			
Kocuria kristinae		9.96	0.30		99	1			
020442	**Low discrimination**								**020442**
					NOV	PYR			
Aerococcus viridans		65.89	0.61	ESL 90	1	95			
Staphylococcus auricularis		30.48	0.39		1	40			
Staphylococcus cohnii subsp. cohnii		1.42	0.15		99	1			
020443	**Good identification**								**020443**
					OXI				
Aerococcus viridans		93.63	0.69	ESL 90	1				
Kocuria kristinae		3.37	0.17		99				
Staphylococcus auricularis		2.06	0.19		1				
020450	**Low discrimination**								**020450**
					OXI	YEP			
Macrococcus bovicus		60.90	0.59	BGA 1	99	99			

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		27.43	0.58		1	12			
Macrococcus equipersicus		3.95	0.36		99	99			
020451	**Low discrimination**								**020451**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		77.53	0.59	BGA 1	1	99	1	12	
Aerococcus viridans		20.50	0.59		1	1	95	1	
Macrococcus bovicus		1.61	0.18		99	99	1	99	
020452	**Low discrimination**								**020452**
					NOV	YEP	PYR		
Aerococcus viridans		79.44	0.71	ESL 90	1	1	95		
Staphylococcus kloosii		9.46	0.53		99	99	52		
Staphylococcus saprophyticus subsp. saprophyticus		8.77	0.23		99	16	7		
020453	**Very good identification**								**020453**
					NOV	PYR			
Aerococcus viridans		99.32	0.79	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		0.42	0.18		99	1			
Staphylococcus cohnii subsp. urealyticum		0.26	0.14		99	53			
020460	**Low discrimination**								**020460**
					PYR				
Aerococcus viridans		55.73	0.42	ESL 90	95				
Staphylococcus auricularis		21.09	0.18	LAC 90	40				
Kocuria varians		12.37	0.14		10				
020461	**Acceptable identification**								**020461**
					NOV	PYR			
Aerococcus viridans		91.74	0.49	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		8.26	0.16	LAC 90	99	1			
020462	**Very good identification**								**020462**
					PYR				
Aerococcus viridans		99.31	0.61	ESL 90	95				
Kocuria varians		0.69	0.03		10				
020463	**Very good identification**								**020463**
Aerococcus viridans		100.0	0.69	ESL 90					
020470	**Low discrimination**								**020470**
					OXI	NOV	YEP		
Aerococcus viridans		66.12	0.51	ESL 90	1	1	1		
Staphylococcus kloosii		16.35	0.40	LAC 90	1	99	99		
Macrococcus bovicus		12.08	0.18		99	99	99		
020471	**Very good identification**								**020471**
					NOV	PYR			
Aerococcus viridans		96.32	0.59	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		3.68	0.18	LAC 90	99	1			
020472	**Very good identification**								**020472**
					NOV	YEP			
Aerococcus viridans		96.48	0.71	ESL 90	1	1			
Staphylococcus kloosii		2.87	0.40		99	99			
Staphylococcus arlettae		0.65	0.15		99	99			
020473	**Very good identification**								**020473**
					NOV	YEP			
Aerococcus viridans		99.62	0.79	ESL 90	1	1			
Staphylococcus arlettae		0.38	0.18		99	99			
020500	**Low discrimination**								**020500**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					NOV	OXI			
Kocuria varians		51.34	0.41	TRE 7	1	13			
Staphylococcus auricularis		21.22	0.31		1	1			
Macrococcus carouselicus		13.42	0.20		99	99			
020501	**Low discrimination**								**020501**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		36.52	0.27	BGA 1	99	1			
Aerococcus viridans		29.18	0.37	GAL 10	1	95			
Kocuria varians		18.78	0.19		1	10			
020502	**Low discrimination**								**020502**
					PYR				
Aerococcus viridans		61.38	0.49	ESL 90	95				
Kocuria varians		33.21	0.29		10				
Staphylococcus auricularis		5.41	0.12		40				
020503	**Very good identification**								**020503**
					PYR				
Aerococcus viridans		98.02	0.56	ESL 90	95				
Kocuria varians		1.98	0.07		10				
020510	**Low discrimination**								**020510**
					OXI	NOV	PYR	YEP	
Macrococcus bovicus		75.75	0.44	BGA 1	99	99	1	99	
Staphylococcus cohnii subsp. cohnii		6.52	0.28		1	99	1	12	
Aerococcus viridans		5.70	0.39		1	1	95	1	
020511	**Low discrimination**								**020511**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		62.96	0.47	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		33.42	0.29	LAC 90	1	99	1	12	
Macrococcus bovicus		3.62	0.03		99	99	1	99	
020512	**Good identification**								**020512**
					OXI	NOV	YEP		
Aerococcus viridans		94.02	0.59	ESL 90	1	1	1		
Staphylococcus kloosii		4.58	0.32		1	99	99		
Macrococcus bovicus		1.40	0.03		99	99	99		
020513	**Very good identification**								**020513**
Aerococcus viridans		100.0	0.66	ESL 90					
020520	**Low discrimination**								**020520**
					PYR				
Kocuria varians		85.85	0.31	TRE 7	10				
Aerococcus viridans		14.15	0.29		95				
020521	**Low discrimination**								**020521**
					PYR				
Aerococcus viridans		81.56	0.37	ESL 90	95				
Kocuria varians		18.44	0.09	LAC 90	10				
020522	**Low discrimination**								**020522**
					PYR				
Aerococcus viridans		84.02	0.49	ESL 90	95				
Kocuria varians		15.98	0.20		10				
020523	**Very good identification**								**020523**
Aerococcus viridans		100.0	0.56	ESL 90					
020530	**Low discrimination**								**020530**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI	NOV	YEP		
Aerococcus viridans		80.95	0.39	ESL 90	1	1	1		
Macrococcus bovicus		10.87	0.03	LAC 90	99	99	99		
Staphylococcus kloosii		8.18	0.19		1	99	99		
020531	**Good identification**								**020531**
Aerococcus viridans		100.0	0.47	ESL 90					
020532	**Very good identification**								**020532**
					NOV	YEP			
Aerococcus viridans		98.80	0.59	ESL 90	1	1			
Staphylococcus kloosii		1.20	0.20		99	99			
020533	**Very good identification**								**020533**
Aerococcus viridans		100.0	0.66	ESL 90					
020540	**Low discrimination**								**020540**
					NOV	PYR			
Staphylococcus auricularis		48.31	0.39	BGA 14	1	40			
Staphylococcus cohnii subsp. cohnii		24.76	0.36	NIT 90	99	1			
Aerococcus viridans		11.61	0.42	GAL 10	1	95			
020541	**Low discrimination**								**020541**
					OXI	NOV	PYR		
Aerococcus viridans		37.65	0.49	ESL 90	1	1	95		
Staphylococcus cohnii subsp. cohnii		37.30	0.37	LAC 90	1	99	1		
Kocuria kristinae		16.27	0.20		99	1	93		
020542	**Good identification**								**020542**
					PYR				
Aerococcus viridans		91.90	0.61	ESL 90	95				
Staphylococcus auricularis		4.72	0.19		40				
Kocuria varians		2.73	0.16		10				
020543	**Very good identification**								**020543**
					OXI				
Aerococcus viridans		98.81	0.69	ESL 90	1				
Kocuria kristinae		1.19	0.07		99				
020550	**Low discrimination**								**020550**
					OXI	NOV	YEP		
Macrococcus bovicus		63.97	0.51	BGA 1	99	99	99		
Staphylococcus kloosii		15.85	0.58		1	99	99		
Aerococcus viridans		8.25	0.51		1	1	1		
020551	**Low discrimination**								**020551**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		68.39	0.59	ESL 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		28.75	0.39	LAC 90	1	99	1	12	
Macrococcus bovicus		2.30	0.10		99	99	1	99	
020552	**Low discrimination**								**020552**
					NOV	YEP	PYR		
Aerococcus viridans		80.04	0.71	ESL 90	1	1	95		
Staphylococcus kloosii		18.50	0.59		99	99	52		
Staphylococcus saprophyticus subsp. saprophyticus		0.77	0.02		99	16	7		
020553	**Very good identification**								**020553**
					NOV	YEP			
Aerococcus viridans		99.79	0.79	ESL 90	1	1			
Staphylococcus cohnii subsp. urealyticum		0.11	0.07		99	64			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		0.10	0.18		99	99		
020560	**Low discrimination**							**020560**
					PYR			
Aerococcus viridans		75.02	0.42	ESL 90	95			
Kocuria varians		24.98	0.18	LAC 90	10			
020561	**Good identification**							**020561**
Aerococcus viridans		100.0	0.49	ESL 90				
020562	**Very good identification**							**020562**
					PYR			
Aerococcus viridans		98.97	0.61	ESL 90	95			
Kocuria varians		1.03	0.06		10			
020563	**Very good identification**							**020563**
Aerococcus viridans		100.0	0.69	ESL 90				
020570	**Low discrimination**							**020570**
					OXI	NOV	YEP	
Aerococcus viridans		64.17	0.51	ESL 90	1	1	1	
Staphylococcus kloosii		30.81	0.46	LAC 90	1	99	99	
Macrococcus bovicus		5.02	0.10		99	99	99	
020571	**Very good identification**							**020571**
Aerococcus viridans		100.0	0.59	ESL 90				
020572	**Good identification**							**020572**
					NOV	YEP		
Aerococcus viridans		91.24	0.71	ESL 90	1	1		
Staphylococcus kloosii		5.27	0.46		99	99		
Staphylococcus arlettae		3.48	0.31		99	99		
020573	**Very good identification**							**020573**
					NOV	YEP		
Aerococcus viridans		97.88	0.79	ESL 90	1	1		
Staphylococcus arlettae		2.12	0.34		99	99		
020600	**Low discrimination**							**020600**
					NOV	OXI		
Staphylococcus auricularis		68.52	0.50	BGA 14	1	1		
Macrococcus carouselicus		24.71	0.34	NIT 90	99	99		
Kocuria varians		1.81	0.19		1	13		
020601	**Acceptable identification**							**020601**
					OXI			
Kocuria kristinae		93.54	0.51	VPT 90	99			
Staphylococcus auricularis		4.63	0.31	MLT 90	1			
Aerococcus viridans		1.64	0.37		1			
020602	**Low discrimination**							**020602**
					PYR			
Aerococcus viridans		54.99	0.49	ESL 90	95			
Staphylococcus auricularis		40.25	0.31		40			
Kocuria varians		2.71	0.08		10			
020603	**Low discrimination**							**020603**
					OXI			
Kocuria kristinae		60.25	0.39	VPT 90	99			
Aerococcus viridans		38.02	0.56	MLT 90	1			
Staphylococcus auricularis		1.33	0.11		1			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
020610	**Low discrimination**							**020610**
Macrococcus bovicus		52.12	0.39	BGA 1				
Macrococcus carouselicus		30.64	0.30					
Staphylococcus vitulinus		7.32	0.15					
020611	**Low discrimination**							**020611**
					OXI	NOV	PYR	
Aerococcus viridans		53.36	0.47	ESL 90	1	1	95	
Kocuria kristinae		42.28	0.23	LAC 90	99	1	93	
Staphylococcus cohnii subsp. cohnii		2.58	0.08		1	99	1	
020612	**Good identification**							**020612**
					NOV	PYR		
Aerococcus viridans		94.02	0.59	ESL 90	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		4.24	0.03		99	7		
Staphylococcus simulans		0.92	0.01		1	70		
020613	**Very good identification**							**020613**
					OXI			
Aerococcus viridans		96.52	0.66	ESL 90	1			
Kocuria kristinae		2.12	0.10		99			
Staphylococcus simulans		1.36	0.11		1			
020620	**Low discrimination**							**020620**
					PYR			
Aerococcus viridans		46.50	0.29	ESL 90	95			
Staphylococcus auricularis		27.85	0.09	LAC 90	40			
Kocuria varians		25.65	0.10		10			
020621	**Low discrimination**							**020621**
					OXI			
Aerococcus viridans		63.44	0.37	ESL 90	1			
Kocuria kristinae		36.56	0.10	LAC 90	99			
020622	**Good identification**							**020622**
Aerococcus viridans		100.0	0.49	ESL 90				
020623	**Very good identification**							**020623**
					OXI			
Aerococcus viridans		98.43	0.56	ESL 90	1			
Kocuria kristinae		1.58	-.03		99			
020631	**Good identification**							**020631**
Aerococcus viridans		100.0	0.47	ESL 90				
020632	**Very good identification**							**020632**
					NOV	YEP		
Aerococcus viridans		97.38	0.59	ESL 90	1	1		
Staphylococcus arlettae		2.62	0.15		99	99		
020633	**Very good identification**							**020633**
					NOV	YEP		
Aerococcus viridans		98.49	0.66	ESL 90	1	1		
Staphylococcus arlettae		1.51	0.18		99	99		
020640	**Low discrimination**							**020640**
					OXI			
Staphylococcus auricularis		63.78	0.58	BGA 14	1			
Staphylococcus petrasii subsp. pragensis		25.54	0.33	NIT 90	1			
Kocuria kristinae		5.58	0.30		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
020641	**Very good identification**							**020641**
					OXI			
Kocuria kristinae		97.87	0.71	VPT 90	99			
Staphylococcus auricularis		1.26	0.38		1			
Aerococcus viridans		0.76	0.49		1			
020642	**Low discrimination**							**020642**
					NOV	PYR		
Aerococcus viridans		58.46	0.61	ESL 90	1	95		
Staphylococcus auricularis		24.96	0.38		1	40		
Staphylococcus saprophyticus subsp. saprophyticus		8.15	0.16		99	7		
020643	**Low discrimination**							**020643**
					OXI			
Kocuria kristinae		77.75	0.58	VPT 90	99			
Aerococcus viridans		21.81	0.69		1			
Staphylococcus auricularis		0.44	0.19		1			
020650	**Low discrimination**							**020650**
					OXI	NOV	PYR	
Macrococcus bovicus		55.82	0.47	BGA 1	99	99	1	
Staphylococcus saprophyticus subsp. saprophyticus		21.88	0.29		1	99	7	
Aerococcus viridans		12.35	0.51		1	1	95	
020651	**Low discrimination**							**020651**
					OXI	NOV	PYR	
Kocuria kristinae		62.76	0.42	VPT 90	99	1	93	
Aerococcus viridans		35.21	0.59	MAN 4	1	1	95	
Staphylococcus cohnii subsp. cohnii		1.35	0.18		1	99	1	
020652	**Low discrimination**							**020652**
					NOV	YEP	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		51.93	0.44	URE 99	99	16	7	
Aerococcus viridans		46.55	0.71		1	1	95	
Staphylococcus kloosii		0.55	0.32		99	99	52	
020653	**Good identification**							**020653**
					OXI	NOV	PYR	
Aerococcus viridans		94.36	0.79	ESL 90	1	1	95	
Kocuria kristinae		4.67	0.30		99	1	93	
Staphylococcus saprophyticus subsp. saprophyticus		0.46	0.03		1	99	7	
020660	**Low discrimination**							**020660**
Aerococcus viridans		67.15	0.42	ESL 90				
Staphylococcus auricularis		23.46	0.17	LAC 90				
Staphylococcus petrasii subsp. pragensis		9.40	-.09					
020661	**Low discrimination**							**020661**
					OXI			
Kocuria kristinae		56.45	0.30	VPT 90	99			
Aerococcus viridans		43.55	0.49	XYL 1	1			
020662	**Very good identification**							**020662**
					NOV	YEP		
Aerococcus viridans		98.02	0.61	ESL 90	1	1		
Staphylococcus arlettae		1.98	0.15		99	99		
020663	**Very good identification**							**020663**
					OXI	NOV	YEP	
Aerococcus viridans		95.47	0.69	ESL 90	1	1	1	
Kocuria kristinae		3.44	0.17		99	1	15	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		1.10	0.18		1	99	99	
020670	**Low discrimination**							**020670**
					OXI	NOV	YEP	
Aerococcus viridans		88.44	0.51	ESL 90	1	1	1	
Staphylococcus arlettae		5.36	0.15	LAC 90	1	99	99	
Macrococcus bovicus		4.04	0.05		99	99	99	
020671	**Very good identification**							**020671**
					OXI	NOV	YEP	
Aerococcus viridans		95.02	0.59	ESL 90	1	1	1	
Staphylococcus arlettae		3.27	0.18	LAC 90	1	99	99	
Kocuria kristinae		1.71	0.01		99	1	15	
020672	**Low discrimination**							**020672**
					NOV	PYR		
Aerococcus viridans		59.49	0.71	ESL 90	1	95		
Staphylococcus arlettae		39.67	0.57		99	1		
Staphylococcus saprophyticus subsp. saprophyticus		0.67	0.03		99	7		
020673	**Low discrimination**							**020673**
					NOV	YEP		
Aerococcus viridans		72.52	0.79	ESL 90	1	1		
Staphylococcus arlettae		27.48	0.59		99	99		
020700	**Low discrimination**							**020700**
					NOV	OXI		
Macrococcus carouselicus		46.87	0.27	BGA 1	99	99		
Staphylococcus auricularis		33.69	0.31	ESL 94	1	1		
Kocuria varians		12.04	0.23		1	13		
020701	**Acceptable identification**							**020701**
					OXI			
Kocuria kristinae		92.48	0.41	VPT 90	99			
Aerococcus viridans		4.86	0.37	MLT 90	1			
Staphylococcus auricularis		1.53	0.11		1			
020702	**Low discrimination**							**020702**
					PYR			
Aerococcus viridans		86.57	0.49	ESL 90	95			
Staphylococcus auricularis		7.04	0.11		40			
Kocuria varians		6.39	0.11		10			
020703	**Low discrimination**							**020703**
					OXI	COA		
Aerococcus viridans		62.72	0.56	ESL 90	1	1		
Kocuria kristinae		33.13	0.29		99	1		
Staphylococcus intermedius		4.16	0.14		1	99		
020710	**Low discrimination**							**020710**
					OXI	NOV		
Macrococcus bovicus		52.33	0.31	BGA 1	99	99		
Macrococcus carouselicus		30.76	0.23		99	99		
Aerococcus viridans		15.75	0.39		1	1		
020711	**Low discrimination**							**020711**
					OXI	COA		
Aerococcus viridans		73.99	0.47	ESL 90	1	1		
Kocuria kristinae		19.54	0.13	LAC 90	99	1		
Staphylococcus intermedius		6.48	0.06		1	99		
020712	**Very good identification**							**020712**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.59	ESL 90				
020713	**Good identification**							**020713**
					OXI	COA	NOV	
Aerococcus viridans		92.77	0.66	ESL 90	1	1	1	
Staphylococcus intermedius		5.54	0.23		1	99	1	
Staphylococcus lentus		1.00	0.12		99	1	99	
020720	**Low discrimination**							**020720**
					PYR			
Aerococcus viridans		54.73	0.29	ESL 90	95			
Kocuria varians		45.27	0.13	LAC 90	10			
020721	**Low discrimination**							**020721**
					OXI			
Aerococcus viridans		83.89	0.37	ESL 90	1			
Kocuria kristinae		16.11	0.00	LAC 90	99			
020722	**Good identification**							**020722**
					PYR			
Aerococcus viridans		97.47	0.49	ESL 90	95			
Kocuria varians		2.53	0.02		10			
020723	**Very good identification**							**020723**
Aerococcus viridans		100.0	0.56	ESL 90				
020730	**Good identification**							**020730**
Aerococcus viridans		100.0	0.39	ESL 90				
020731	**Good identification**							**020731**
Aerococcus viridans		100.0	0.47	ESL 90				
020732	**Low discrimination**							**020732**
					NOV	YEP		
Aerococcus viridans		86.75	0.59	ESL 90	1	1		
Staphylococcus arlettae		13.25	0.31		99	99		
020733	**Good identification**							**020733**
					OXI	NOV	YEP	
Aerococcus viridans		91.11	0.66	ESL 90	1	1	1	
Staphylococcus arlettae		7.90	0.34		1	99	99	
Staphylococcus lentus		0.99	0.12		99	99	1	
020740	**Low discrimination**							**020740**
					OXI			
Staphylococcus auricularis		57.49	0.38	BGA 14	1			
Kocuria kristinae		15.09	0.20	NIT 90	99			
Aerococcus viridans		14.96	0.42	GAL 10	1			
020741	**Very good identification**							**020741**
					OXI			
Kocuria kristinae		97.06	0.61	VPT 90	99			
Aerococcus viridans		2.27	0.49		1			
Staphylococcus auricularis		0.42	0.19		1			
020742	**Good identification**							**020742**
					OXI			
Aerococcus viridans		90.32	0.61	ESL 90	1			
Staphylococcus auricularis		4.29	0.19		1			
Kocuria kristinae		2.53	0.07		99			
020743	**Low discrimination**							**020743**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	COA		
Kocuria kristinae		53.70	0.48	VPT 90	99	1		
Aerococcus viridans		45.18	0.69		1	1		
Staphylococcus intermedius		1.12	0.17		1	99		
020750	**Low discrimination**							**020750**
					OXI	NOV	YEP	
Macrococcus bovicus		54.05	0.39	BGA 1	99	99	99	
Aerococcus viridans		27.89	0.51		1	1	1	
Staphylococcus kloosii		5.30	0.37		1	99	99	
020751	**Low discrimination**							**020751**
					OXI	COA		
Aerococcus viridans		60.04	0.59	ESL 90	1	1		
Kocuria kristinae		35.67	0.32	LAC 90	99	1		
Staphylococcus intermedius		1.97	0.10		1	99		
020752	**Low discrimination**							**020752**
					NOV	PYR		
Aerococcus viridans		85.67	0.71	ESL 90	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		8.31	0.22		99	7		
Staphylococcus arlettae		3.27	0.31		99	1		
020753	**Good identification**							**020753**
					COA	NOV	YEP	
Aerococcus viridans		93.80	0.79	ESL 90	1	1	1	
Staphylococcus intermedius		2.10	0.26		99	1	1	
Staphylococcus arlettae		2.03	0.34		1	99	99	
020760	**Good identification**							**020760**
Aerococcus viridans		100.0	0.42	ESL 90				
020761	**Low discrimination**							**020761**
					OXI			
Aerococcus viridans		69.83	0.49	ESL 90	1			
Kocuria kristinae		30.18	0.20	LAC 90	99			
020762	**Low discrimination**							**020762**
					NOV	YEP		
Aerococcus viridans		89.73	0.61	ESL 90	1	1		
Staphylococcus arlettae		10.28	0.31		99	99		
020763	**Good identification**							**020763**
					OXI	NOV	YEP	
Aerococcus viridans		92.85	0.69	ESL 90	1	1	1	
Staphylococcus arlettae		6.04	0.34		1	99	99	
Kocuria kristinae		1.12	0.07		99	1	15	
020770	**Low discrimination**							**020770**
					NOV	YEP		
Aerococcus viridans		70.89	0.51	ESL 90	1	1		
Staphylococcus arlettae		24.35	0.31	LAC 90	99	99		
Staphylococcus kloosii		3.37	0.25		99	99		
020771	**Low discrimination**							**020771**
					NOV	YEP		
Aerococcus viridans		81.22	0.59	ESL 90	1	1		
Staphylococcus arlettae		15.85	0.34	LAC 90	99	99		
Staphylococcus xylosus		1.80	0.05		99	40		
020772	**Low discrimination**							**020772**
					NOV	YEP		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		78.98	0.72	GLR 92	99	99	
Aerococcus viridans		20.90	0.71		1	1	
Staphylococcus kloosii		0.12	0.26		99	99	
020773	**Low discrimination**						**020773**
					NOV	YEP	
Staphylococcus arlettae		67.95	0.75	GLR 92	99	99	
Aerococcus viridans		31.65	0.79		1	1	
Staphylococcus xylosus		0.22	0.14		99	40	
021000	**Low discrimination**						**021000**
					NOV		
Macrococcus carouselicus		40.60	0.28	BGA 1	99		
Micrococcus sp.		35.25	0.18	TRE 94	1		
Dermacoccus nishinomiyaensis		12.52	0.18		80		
021001	**Good identification**						**021001**
Aerococcus viridans		100.0	0.30	TRE 95			
021002	**Acceptable identification**						**021002**
					PYR		
Aerococcus viridans		90.73	0.42	TRE 95	95		
Kocuria varians		9.27	0.08		10		
021003	**Good identification**						**021003**
Aerococcus viridans		100.0	0.50	TRE 95			
021010	**Low discrimination**						**021010**
Macrococcus equipersicus		42.40	0.34	URE 86			
Macrococcus bovicus		26.38	0.28	BGA 1			
Macrococcus carouselicus		25.78	0.24				
021011	**Good identification**						**021011**
Aerococcus viridans		100.0	0.40	TRE 95			
021012	**Good identification**						**021012**
Aerococcus viridans		100.0	0.52	TRE 95			
021013	**Very good identification**						**021013**
Aerococcus viridans		100.0	0.60	TRE 95			
021021	**Good identification**						**021021**
Aerococcus viridans		100.0	0.30	TRE 95			
021022	**Good identification**						**021022**
Aerococcus viridans		100.0	0.42	TRE 95			
021023	**Good identification**						**021023**
Aerococcus viridans		100.0	0.50	TRE 95			
021030	**Low discrimination**						**021030**
					OXI	NOV	
Aerococcus viridans		76.84	0.32	TRE 95	1	1	
Staphylococcus vitulinus		23.17	-.03	LAC 90	99	99	
021031	**Good identification**						**021031**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.40	TRE 95			
021032	**Good identification**						**021032**
Aerococcus viridans		100.0	0.52	TRE 95			
021033	**Very good identification**						**021033**
Aerococcus viridans		100.0	0.60	TRE 95			
021040	**Low discrimination**						**021040**
					NOV	OXI	
Aerococcus viridans		42.12	0.35	TRE 95	1	1	
Macrococcus equipersicus		35.04	0.16	LAC 90	99	99	
Micrococcus sp.		15.30	-.05		1	89	
021041	**Good identification**						**021041**
Aerococcus viridans		100.0	0.42	TRE 95			
021042	**Very good identification**						**021042**
Aerococcus viridans		100.0	0.55	TRE 95			
021043	**Very good identification**						**021043**
Aerococcus viridans		100.0	0.62	TRE 95			
021050	**Low discrimination**						**021050**
					OXI	NOV	
Macrococcus equipersicus		57.91	0.43	URE 86	99	99	
Macrococcus bovicus		31.09	0.36	BGA 1	99	99	
Aerococcus viridans		10.99	0.45		1	1	
021051	**Good identification**						**021051**
					OXI	NOV	
Aerococcus viridans		97.77	0.52	TRE 95	1	1	
Macrococcus equipersicus		2.23	0.02	LAC 90	99	99	
021052	**Very good identification**						**021052**
					OXI	NOV	
Aerococcus viridans		99.41	0.64	TRE 95	1	1	
Macrococcus equipersicus		0.59	0.02		99	99	
021053	**Very good identification**						**021053**
Aerococcus viridans		100.0	0.72	TRE 95			
021060	**Good identification**						**021060**
Aerococcus viridans		100.0	0.35	TRE 95			
021061	**Good identification**						**021061**
Aerococcus viridans		100.0	0.42	TRE 95			
021062	**Very good identification**						**021062**
Aerococcus viridans		100.0	0.55	TRE 95			
021063	**Very good identification**						**021063**
Aerococcus viridans		100.0	0.62	TRE 95			
021070	**Acceptable identification**						**021070**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		94.95	0.45	TRE 95	1	1	
Macrococcus equipersicus		5.05	0.02	LAC 90	99	99	
021071	**Good identification**						**021071**
Aerococcus viridans		100.0	0.52	TRE 95			
021072	**Very good identification**						**021072**
Aerococcus viridans		100.0	0.64	TRE 95			
021073	**Very good identification**						**021073**
Aerococcus viridans		100.0	0.72	TRE 95			
021101	**Good identification**						**021101**
Aerococcus viridans		100.0	0.30	TRE 95			
021102	**Low discrimination**						**021102**
					PYR		
Aerococcus viridans		86.71	0.42	TRE 95	95		
Kocuria varians		13.29	0.11		10		
021103	**Good identification**						**021103**
Aerococcus viridans		100.0	0.50	TRE 95			
021111	**Good identification**						**021111**
Aerococcus viridans		100.0	0.40	TRE 95			
021112	**Good identification**						**021112**
Aerococcus viridans		100.0	0.52	TRE 95			
021113	**Very good identification**						**021113**
Aerococcus viridans		100.0	0.60	TRE 95			
021121	**Good identification**						**021121**
Aerococcus viridans		100.0	0.30	TRE 95			
021122	**Acceptable identification**						**021122**
					PYR		
Aerococcus viridans		94.89	0.42	TRE 95	95		
Kocuria varians		5.11	0.02		10		
021123	**Good identification**						**021123**
Aerococcus viridans		100.0	0.50	TRE 95			
021130	**Good identification**						**021130**
Aerococcus viridans		100.0	0.32	TRE 95			
021131	**Good identification**						**021131**
Aerococcus viridans		100.0	0.40	TRE 95			
021132	**Good identification**						**021132**
Aerococcus viridans		100.0	0.52	TRE 95			
021133	**Very good identification**						**021133**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.60	TRE 95				
021140	**Low discrimination**							**021140**
					NOV	PYR	OXI	
Aerococcus viridans		61.54	0.35	TRE 95	1	95	1	
Macrococcus equipersicus		21.94	0.09	LAC 90	99	1	99	
Kocuria varians		16.52	0.09		1	10	13	
021141	**Good identification**							**021141**
Aerococcus viridans		100.0	0.42	TRE 95				
021142	**Very good identification**							**021142**
Aerococcus viridans		100.0	0.55	TRE 95				
021143	**Very good identification**							**021143**
Aerococcus viridans		100.0	0.62	TRE 95				
021150	**Low discrimination**							**021150**
					OXI	NOV		
Macrococcus equipersicus		50.05	0.35	URE 86	99	99		
Macrococcus bovicus		26.87	0.28	BGA 1	99	99		
Aerococcus viridans		22.17	0.45		1	1		
021151	**Good identification**							**021151**
Aerococcus viridans		100.0	0.52	TRE 95				
021152	**Very good identification**							**021152**
					NOV	YEP		
Aerococcus viridans		99.51	0.64	TRE 95	1	1		
Staphylococcus kloosii		0.49	0.18		99	99		
021153	**Very good identification**							**021153**
Aerococcus viridans		100.0	0.72	TRE 95				
021160	**Good identification**							**021160**
Aerococcus viridans		100.0	0.35	TRE 95				
021161	**Good identification**							**021161**
Aerococcus viridans		100.0	0.42	TRE 95				
021162	**Very good identification**							**021162**
Aerococcus viridans		100.0	0.55	TRE 95				
021163	**Very good identification**							**021163**
Aerococcus viridans		100.0	0.62	TRE 95				
021170	**Good identification**							**021170**
Aerococcus viridans		100.0	0.45	TRE 95				
021171	**Good identification**							**021171**
Aerococcus viridans		100.0	0.52	TRE 95				
021172	**Very good identification**							**021172**
Aerococcus viridans		100.0	0.64	TRE 95				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
021173	**Very good identification**						**021173**
Aerococcus viridans		100.0	0.72	TRE 95			
021200	**Acceptable identification**						**021200**
					NOV	OXI	
Macrococcus carouselicus		93.95	0.34	BGA 1	99	99	
Micrococcus sp.		3.97	-.03	TRE 94	1	89	
Aerococcus viridans		2.08	0.23		1	1	
021202	**Good identification**						**021202**
Aerococcus viridans		100.0	0.42	TRE 95			
021203	**Good identification**						**021203**
					OXI		
Aerococcus viridans		95.78	0.50	TRE 95	1		
Kocuria kristinae		4.22	0.00		99		
021210	**Low discrimination**						**021210**
Staphylococcus vitulinus		46.69	0.29	BGA 1			
Macrococcus carouselicus		40.69	0.30	NIT 91			
Macrococcus bovicus		5.13	0.16				
021211	**Good identification**						**021211**
Aerococcus viridans		100.0	0.40	TRE 95			
021212	**Good identification**						**021212**
Aerococcus viridans		100.0	0.52	TRE 95			
021213	**Very good identification**						**021213**
Aerococcus viridans		100.0	0.60	TRE 95			
021221	**Good identification**						**021221**
Aerococcus viridans		100.0	0.30	TRE 95			
021222	**Good identification**						**021222**
Aerococcus viridans		100.0	0.42	TRE 95			
021223	**Good identification**						**021223**
Aerococcus viridans		100.0	0.50	TRE 95			
021230	**Good identification**						**021230**
					OXI	NOV	
Staphylococcus vitulinus		96.76	0.38	BGA 1	99	99	
Aerococcus viridans		3.24	0.32	NIT 91	1	1	
021231	**Low discrimination**						**021231**
					OXI	NOV	
Aerococcus viridans		88.56	0.40	TRE 95	1	1	
Staphylococcus vitulinus		11.44	-.03	LAC 90	99	99	
021232	**Good identification**						**021232**
					OXI	NOV	
Aerococcus viridans		96.76	0.52	TRE 95	1	1	
Staphylococcus vitulinus		3.24	-.03		99	99	
021233	**Very good identification**						**021233**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.60	TRE 95			
021240	**Good identification**						**021240**
Aerococcus viridans		100.0	0.35	TRE 95			
021241	**Low discrimination**						**021241**
					OXI		
Kocuria kristinae		78.10	0.32	VPT 90	99		
Aerococcus viridans		21.91	0.42	TRE 95	1		
021242	**Very good identification**						**021242**
Aerococcus viridans		100.0	0.55	TRE 95			
021243	**Good identification**						**021243**
					OXI		
Aerococcus viridans		90.99	0.62	TRE 95	1		
Kocuria kristinae		9.01	0.19		99		
021250	**Low discrimination**						**021250**
					OXI	NOV	
Aerococcus viridans		44.88	0.45	TRE 95	1	1	
Macrococcus bovicus		31.74	0.23	LAC 90	99	99	
Macrococcus equipersicus		23.38	0.22		99	99	
021251	**Good identification**						**021251**
					OXI		
Aerococcus viridans		95.28	0.52	TRE 95	1		
Kocuria kristinae		4.72	0.03	LAC 90	99		
021252	**Very good identification**						**021252**
Aerococcus viridans		100.0	0.64	TRE 95			
021253	**Very good identification**						**021253**
Aerococcus viridans		100.0	0.72	TRE 95			
021260	**Good identification**						**021260**
Aerococcus viridans		100.0	0.35	TRE 95			
021261	**Good identification**						**021261**
Aerococcus viridans		100.0	0.42	TRE 95			
021262	**Very good identification**						**021262**
Aerococcus viridans		100.0	0.55	TRE 95			
021263	**Very good identification**						**021263**
Aerococcus viridans		100.0	0.62	TRE 95			
021270	**Acceptable identification**						**021270**
					OXI	NOV	
Aerococcus viridans		92.99	0.45	TRE 95	1	1	
Staphylococcus vitulinus		7.01	-.03	LAC 90	99	99	
021271	**Good identification**						**021271**
Aerococcus viridans		100.0	0.52	TRE 95			
021272	**Very good identification**						**021272**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		98.60	0.64	TRE 95	1	1		
Staphylococcus arlettae		1.40	0.15		99	99		
021273	**Very good identification**							**021273**
					NOV	YEP		
Aerococcus viridans		99.20	0.72	TRE 95	1	1		
Staphylococcus arlettae		0.80	0.18		99	99		
021300	**Acceptable identification**							**021300**
					NOV	PYR	OXI	
Macrococcus carouselicus		92.07	0.27	BGA 1	99	1	99	
Aerococcus viridans		4.76	0.23	TRE 94	1	95	1	
Kocuria varians		3.17	0.05		1	10	13	
021301	**Low discrimination**							**021301**
					OXI			
Aerococcus viridans		65.44	0.30	TRE 95	1			
Kocuria kristinae		34.56	0.02	LAC 90	99			
021302	**Good identification**							**021302**
Aerococcus viridans		100.0	0.42	TRE 95				
021303	**Good identification**							**021303**
Aerococcus viridans		100.0	0.50	TRE 95				
021311	**Good identification**							**021311**
Aerococcus viridans		100.0	0.40	TRE 95				
021312	**Good identification**							**021312**
Aerococcus viridans		100.0	0.52	TRE 95				
021313	**Very good identification**							**021313**
					OXI	NOV		
Aerococcus viridans		97.77	0.60	TRE 95	1	1		
Staphylococcus lentus		2.23	0.12		99	99		
021321	**Good identification**							**021321**
Aerococcus viridans		100.0	0.30	TRE 95				
021322	**Good identification**							**021322**
Aerococcus viridans		100.0	0.42	TRE 95				
021323	**Good identification**							**021323**
Aerococcus viridans		100.0	0.50	TRE 95				
021330	**Low discrimination**							**021330**
					OXI	NOV		
Aerococcus viridans		76.84	0.32	TRE 95	1	1		
Staphylococcus vitulinus		23.17	-.03	LAC 90	99	99		
021331	**Good identification**							**021331**
Aerococcus viridans		100.0	0.40	TRE 95				
021332	**Good identification**							**021332**
Aerococcus viridans		100.0	0.52	TRE 95				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
021333	**Very good identification**							**021333**
					OXI	NOV		
Aerococcus viridans		97.77	0.60	TRE 95	1	1		
Staphylococcus lentus		2.23	0.12		99	99		
021340	**Good identification**							**021340**
Aerococcus viridans		100.0	0.35	TRE 95				
021342	**Very good identification**							**021342**
Aerococcus viridans		100.0	0.55	TRE 95				
021343	**Very good identification**							**021343**
					OXI			
Aerococcus viridans		96.80	0.62	TRE 95	1			
Kocuria kristinae		3.20	0.10		99			
021350	**Low discrimination**							**021350**
					OXI	NOV		
Aerococcus viridans		65.52	0.45	TRE 95	1	1		
Macrococcus bovicus		19.86	0.16	LAC 90	99	99		
Macrococcus equipersicus		14.63	0.14		99	99		
021351	**Good identification**							**021351**
					OXI	NOV	PYR	
Aerococcus viridans		95.61	0.52	TRE 95	1	1	95	
Staphylococcus lentus		2.81	0.07	LAC 90	99	99	6	
Kocuria kristinae		1.58	-.06		99	1	93	
021352	**Very good identification**							**021352**
Aerococcus viridans		100.0	0.64	TRE 95				
021353	**Very good identification**							**021353**
					OXI	NOV		
Aerococcus viridans		98.85	0.72	TRE 95	1	1		
Staphylococcus lentus		1.15	0.18		99	99		
021360	**Good identification**							**021360**
Aerococcus viridans		100.0	0.35	TRE 95				
021361	**Good identification**							**021361**
Aerococcus viridans		100.0	0.42	TRE 95				
021362	**Very good identification**							**021362**
Aerococcus viridans		100.0	0.55	TRE 95				
021363	**Very good identification**							**021363**
Aerococcus viridans		100.0	0.62	TRE 95				
021370	**Good identification**							**021370**
Aerococcus viridans		100.0	0.45	TRE 95				
021371	**Good identification**							**021371**
					OXI	NOV		
Aerococcus viridans		97.14	0.52	TRE 95	1	1		
Staphylococcus lentus		2.86	0.07	LAC 90	99	99		
021372	**Good identification**							**021372**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		92.54	0.64	TRE 95	1	1		
Staphylococcus arlettae		7.46	0.31		99	99		
021373	**Good identification**							**021373**
					OXI	NOV	YEP	
Aerococcus viridans		94.57	0.72	TRE 95	1	1	1	
Staphylococcus arlettae		4.33	0.34		1	99	99	
Staphylococcus lentus		1.10	0.18		99	99	1	
021400	**Acceptable identification**							**021400**
					OXI	NOV		
Macrococcus carouselicus		94.03	0.53	BGA 1	99	99		
Aerococcus viridans		5.13	0.49		1	1		
Staphylococcus auricularis		0.37	0.10		1	1		
021401	**Good identification**							**021401**
					OXI	NOV	PYR	
Aerococcus viridans		90.25	0.56	LAC 90	1	1	95	
Macrococcus carouselicus		7.16	0.12		99	99	1	
Kocuria kristinae		1.44	-.03		99	1	93	
021402	**Very good identification**							**021402**
					OXI	NOV		
Aerococcus viridans		97.98	0.68		1	1		
Macrococcus carouselicus		2.02	0.12		99	99		
021403	**Very good identification**							**021403**
Aerococcus viridans		100.0	0.76					
021410	**Low discrimination**							**021410**
					OXI	NOV		
Macrococcus carouselicus		68.13	0.49	BGA 1	99	99		
Aerococcus viridans		17.44	0.59		1	1		
Macrococcus equipersicus		7.15	0.34		99	99		
021411	**Very good identification**							**021411**
					OXI	NOV	PYR	
Aerococcus viridans		97.82	0.66	LAC 90	1	1	95	
Macrococcus carouselicus		1.66	0.08		99	99	1	
Staphylococcus cohnii subsp. cohnii		0.53	0.08		1	99	1	
021412	**Very good identification**							**021412**
					OXI	NOV	YEP	
Aerococcus viridans		99.29	0.78		1	1	1	
Macrococcus carouselicus		0.44	0.08		99	99	67	
Staphylococcus kloosii		0.28	0.26		1	99	99	
021413	**Very good identification**							**021413**
Aerococcus viridans		100.0	0.86					
021420	**Low discrimination**							**021420**
					OXI	NOV		
Aerococcus viridans		84.37	0.49	LAC 90	1	1		
Macrococcus carouselicus		15.63	0.12		99	99		
021421	**Very good identification**							**021421**
Aerococcus viridans		100.0	0.56	LAC 90				
021422	**Very good identification**							**021422**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Aerococcus viridans		100.0	0.68						
021423	**Very good identification**								**021423**
Aerococcus viridans		100.0	0.76						
021430	**Very good identification**								**021430**
					OXI	NOV			
Aerococcus viridans		95.38	0.59	LAC 90	1	1			
Macrococcus carouselicus		3.77	0.08		99	99			
Staphylococcus vitulinus		0.85	-.08		99	99			
021431	**Very good identification**								**021431**
Aerococcus viridans		100.0	0.66	LAC 90					
021432	**Very good identification**								**021432**
Aerococcus viridans		100.0	0.78						
021433	**Very good identification**								**021433**
Aerococcus viridans		100.0	0.86						
021440	**Low discrimination**								**021440**
					OXI	NOV			
Aerococcus viridans		86.17	0.61	LAC 90	1	1			
Macrococcus carouselicus		3.99	0.12		99	99			
Macrococcus equipersicus		3.77	0.16		99	99			
021441	**Very good identification**								**021441**
					OXI	NOV	PYR		
Aerococcus viridans		95.60	0.69	LAC 90	1	1	95		
Kocuria kristinae		3.44	0.17		99	1	93		
Staphylococcus cohnii subsp. cohnii		0.96	0.16		1	99	1		
021442	**Very good identification**								**021442**
Aerococcus viridans		100.0	0.81						
021443	**Very good identification**								**021443**
					OXI				
Aerococcus viridans		99.90	0.88		1				
Kocuria kristinae		0.10	0.05		99				
021450	**Low discrimination**								**021450**
					OXI	NOV			
Aerococcus viridans		61.37	0.71	LAC 90	1	1			
Macrococcus equipersicus		17.02	0.43		99	99			
Macrococcus bovicus		13.70	0.39		99	99			
021451	**Very good identification**								**021451**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		99.36	0.79	LAC 90	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		0.42	0.18		1	99	1	12	
Macrococcus equipersicus		0.12	0.02		99	99	1	99	
021452	**Very good identification**								**021452**
					OXI	NOV	YEP		
Aerococcus viridans		98.64	0.91		1	1	1		
Staphylococcus kloosii		1.31	0.53		1	99	99		
Macrococcus equipersicus		0.03	0.02		99	99	99		
021453	**Excellent identification**								**021453**
					NOV				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		99.99	0.98		1			
Staphylococcus cohnii subsp. urealyticum		0.01	0.06		99			
021460	**Very good identification**							**021460**
Aerococcus viridans		100.0	0.61	LAC 90				
021461	**Very good identification**							**021461**
Aerococcus viridans		100.0	0.69	LAC 90				
021462	**Very good identification**							**021462**
Aerococcus viridans		100.0	0.81					
021463	**Very good identification**							**021463**
Aerococcus viridans		100.0	0.88					
021470	**Very good identification**							**021470**
					OXI	NOV	YEP	
Aerococcus viridans		96.85	0.71	LAC 90	1	1	1	
Staphylococcus kloosii		2.66	0.40		1	99	99	
Macrococcus equipersicus		0.27	0.02		99	99	99	
021471	**Very good identification**							**021471**
Aerococcus viridans		100.0	0.79	LAC 90				
021472	**Very good identification**							**021472**
					NOV	YEP		
Aerococcus viridans		99.60	0.91		1	1		
Staphylococcus kloosii		0.33	0.40		99	99		
Staphylococcus arlettae		0.08	0.15		99	99		
021473	**Excellent identification**							**021473**
					NOV	YEP		
Aerococcus viridans		99.96	0.98		1	1		
Staphylococcus arlettae		0.04	0.18		99	99		
021500	**Low discrimination**							**021500**
					OXI	NOV		
Macrococcus carouselicus		88.43	0.45	BGA 1	99	99		
Aerococcus viridans		11.25	0.49		1	1		
Macrococcus equipersicus		0.31	0.00		99	99		
021501	**Very good identification**							**021501**
					OXI	NOV		
Aerococcus viridans		96.71	0.56	LAC 90	1	1		
Macrococcus carouselicus		3.29	0.04		99	99		
021502	**Very good identification**							**021502**
					OXI	NOV		
Aerococcus viridans		99.13	0.68		1	1		
Macrococcus carouselicus		0.87	0.04		99	99		
021503	**Very good identification**							**021503**
Aerococcus viridans		100.0	0.76					
021510	**Low discrimination**							**021510**
					OXI	NOV		
Macrococcus carouselicus		54.74	0.41	BGA 1	99	99		
Aerococcus viridans		32.69	0.59		1	1		
Macrococcus equipersicus		5.75	0.26		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
021511	**Very good identification**							**021511**
					OXI	NOV		
Aerococcus viridans		99.28	0.66	LAC 90	1	1		
Macrococcus carouselicus		0.72	0.00		99	99		
021512	**Very good identification**							**021512**
					OXI	NOV	YEP	
Aerococcus viridans		99.28	0.78		1	1	1	
Staphylococcus kloosii		0.54	0.32		1	99	99	
Macrococcus carouselicus		0.19	0.00		99	99	67	
021513	**Very good identification**							**021513**
					OXI	NOV		
Aerococcus viridans		99.88	0.86		1	1		
Staphylococcus lentus		0.12	0.12		99	99		
021520	**Acceptable identification**							**021520**
					OXI	NOV		
Aerococcus viridans		92.65	0.49	LAC 90	1	1		
Macrococcus carouselicus		7.35	0.04		99	99		
021521	**Very good identification**							**021521**
Aerococcus viridans		100.0	0.56	LAC 90				
021522	**Very good identification**							**021522**
Aerococcus viridans		100.0	0.68					
021523	**Very good identification**							**021523**
Aerococcus viridans		100.0	0.76					
021530	**Very good identification**							**021530**
					OXI	NOV	YEP	
Aerococcus viridans		97.26	0.59	LAC 90	1	1	1	
Macrococcus carouselicus		1.65	0.00		99	99	67	
Staphylococcus kloosii		1.09	0.19		1	99	99	
021531	**Very good identification**							**021531**
Aerococcus viridans		100.0	0.66	LAC 90				
021532	**Very good identification**							**021532**
					NOV	YEP		
Aerococcus viridans		99.87	0.78		1	1		
Staphylococcus kloosii		0.14	0.20		99	99		
021533	**Very good identification**							**021533**
					OXI	NOV		
Aerococcus viridans		99.88	0.86		1	1		
Staphylococcus lentus		0.12	0.12		99	99		
021540	**Very good identification**							**021540**
					OXI	NOV		
Aerococcus viridans		95.69	0.61	LAC 90	1	1		
Macrococcus carouselicus		1.90	0.04		99	99		
Macrococcus equipersicus		1.80	0.09		99	99		
021541	**Very good identification**							**021541**
					OXI			
Aerococcus viridans		98.81	0.69	LAC 90	1			
Kocuria kristinae		1.19	0.07		99			
021542	**Very good identification**							**021542**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		99.92	0.81		1	1		
Staphylococcus kloosii		0.08	0.18		99	99		
021543	**Very good identification**							**021543**
					OXI			
Aerococcus viridans		99.97	0.88		1			
Kocuria kristinae		0.03	-.05		99			
021550	**Low discrimination**							**021550**
					OXI	NOV	YEP	
Aerococcus viridans		69.83	0.71	LAC 90	1	1	1	
Staphylococcus kloosii		14.90	0.58		1	99	99	
Macrococcus equipersicus		8.30	0.35		99	99	99	
021551	**Very good identification**							**021551**
					OXI	NOV	YEP	
Aerococcus viridans		99.75	0.79	LAC 90	1	1	1	
Staphylococcus lentus		0.16	0.07		99	99	1	
Staphylococcus kloosii		0.09	0.17		1	99	99	
021552	**Very good identification**							**021552**
					NOV	YEP		
Aerococcus viridans		97.50	0.91		1	1		
Staphylococcus kloosii		2.50	0.59		99	99		
021553	**Excellent identification**							**021553**
					OXI	NOV	YEP	
Aerococcus viridans		99.93	0.98		1	1	1	
Staphylococcus lentus		0.06	0.18		99	99	1	
Staphylococcus kloosii		0.01	0.18		1	99	99	
021560	**Very good identification**							**021560**
Aerococcus viridans		100.0	0.61	LAC 90				
021561	**Very good identification**							**021561**
Aerococcus viridans		100.0	0.69	LAC 90				
021562	**Very good identification**							**021562**
Aerococcus viridans		100.0	0.81					
021563	**Very good identification**							**021563**
Aerococcus viridans		100.0	0.88					
021570	**Good identification**							**021570**
					NOV	YEP		
Aerococcus viridans		94.94	0.71	LAC 90	1	1		
Staphylococcus kloosii		5.06	0.46		99	99		
021571	**Very good identification**							**021571**
					OXI	NOV		
Aerococcus viridans		99.85	0.79	LAC 90	1	1		
Staphylococcus lentus		0.16	0.07		99	99		
021572	**Very good identification**							**021572**
					NOV	YEP		
Aerococcus viridans		98.95	0.91		1	1		
Staphylococcus kloosii		0.64	0.46		99	99		
Staphylococcus arlettae		0.42	0.31		99	99		
021573	**Excellent identification**							**021573**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	YEP	
Aerococcus viridans		99.70	0.98		1	1	1	
Staphylococcus arlettae		0.24	0.34		1	99	99	
Staphylococcus lentus		0.06	0.18		99	99	1	
021600	**Very good identification**							**021600**
					OXI	NOV		
Macrococcus carouselicus		97.12	0.59	BGA 1	99	99		
Aerococcus viridans		2.61	0.49		1	1		
Staphylococcus auricularis		0.17	0.09		1	1		
021601	**Low discrimination**							**021601**
					OXI	NOV	PYR	
Kocuria kristinae		57.71	0.39	VPT 90	99	1	93	
Aerococcus viridans		36.42	0.56	MLT 90	1	1	95	
Macrococcus carouselicus		5.87	0.18		99	99	1	
021602	**Very good identification**							**021602**
					OXI	NOV		
Aerococcus viridans		95.99	0.68		1	1		
Macrococcus carouselicus		4.01	0.18		99	99		
021603	**Very good identification**							**021603**
					OXI			
Aerococcus viridans		95.78	0.76		1			
Kocuria kristinae		4.22	0.26		99			
021610	**Low discrimination**							**021610**
					OXI	NOV		
Macrococcus carouselicus		84.43	0.55	BGA 1	99	99		
Aerococcus viridans		10.64	0.59		1	1		
Staphylococcus vitulinus		3.48	0.24		99	99		
021611	**Good identification**							**021611**
					OXI	NOV	PYR	
Aerococcus viridans		94.28	0.66	LAC 90	1	1	95	
Macrococcus carouselicus		3.24	0.14		99	99	1	
Kocuria kristinae		2.08	0.10		99	1	93	
021612	**Very good identification**							**021612**
					OXI	NOV		
Aerococcus viridans		99.12	0.78		1	1		
Macrococcus carouselicus		0.88	0.14		99	99		
021613	**Very good identification**							**021613**
					OXI	NOV	PYR	
Aerococcus viridans		99.82	0.86		1	1	95	
Staphylococcus lentus		0.12	0.12		99	99	6	
Kocuria kristinae		0.06	-.02		99	1	93	
021620	**Low discrimination**							**021620**
					OXI	NOV		
Aerococcus viridans		71.29	0.49	LAC 90	1	1		
Macrococcus carouselicus		26.80	0.18		99	99		
Staphylococcus vitulinus		1.91	-.08		99	99		
021621	**Very good identification**							**021621**
					OXI			
Aerococcus viridans		98.43	0.56	LAC 90	1			
Kocuria kristinae		1.58	-.03		99			
021622	**Very good identification**							**021622**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.68					
021623	**Very good identification**							**021623**
Aerococcus viridans		100.0	0.76					
021630	**Low discrimination**							**021630**
					OXI	NOV		
Aerococcus viridans		50.92	0.59	LAC 90	1	1		
Staphylococcus vitulinus		45.00	0.33		99	99		
Macrococcus carouselicus		4.08	0.14		99	99		
021631	**Very good identification**							**021631**
					OXI	NOV		
Aerococcus viridans		99.62	0.66	LAC 90	1	1		
Staphylococcus vitulinus		0.38	-.08		99	99		
021632	**Very good identification**							**021632**
					OXI	NOV	YEP	
Aerococcus viridans		99.60	0.78		1	1	1	
Staphylococcus arlettae		0.30	0.15		1	99	99	
Staphylococcus vitulinus		0.10	-.08		99	99	75	
021633	**Very good identification**							**021633**
					OXI	NOV	YEP	
Aerococcus viridans		99.71	0.86		1	1	1	
Staphylococcus arlettae		0.17	0.18		1	99	99	
Staphylococcus lentus		0.12	0.12		99	99	1	
021640	**Low discrimination**							**021640**
					OXI	NOV	PYR	
Aerococcus viridans		81.14	0.61	LAC 90	1	1	95	
Macrococcus carouselicus		7.63	0.18		99	99	1	
Kocuria kristinae		6.82	0.17		99	1	93	
021641	**Low discrimination**							**021641**
					OXI			
Kocuria kristinae		78.10	0.58	VPT 90	99			
Aerococcus viridans		21.91	0.69		1			
021642	**Very good identification**							**021642**
					OXI			
Aerococcus viridans		99.77	0.81		1			
Kocuria kristinae		0.23	0.05		99			
021643	**Good identification**							**021643**
					OXI			
Aerococcus viridans		90.99	0.88		1			
Kocuria kristinae		9.01	0.46		99			
021650	**Low discrimination**							**021650**
					OXI	NOV		
Aerococcus viridans		89.75	0.71	LAC 90	1	1		
Macrococcus bovicus		5.01	0.27		99	99		
Macrococcus equipersicus		2.46	0.22		99	99		
021651	**Good identification**							**021651**
					OXI	NOV	PYR	
Aerococcus viridans		94.77	0.79	LAC 90	1	1	95	
Kocuria kristinae		4.69	0.30		99	1	93	
Staphylococcus sciuri		0.39	0.13		99	99	1	
021652	**Very good identification**							**021652**
					NOV	YEP	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		99.67	0.91		1	1	95	
Staphylococcus kloosii		0.13	0.32		99	99	52	
Staphylococcus saprophyticus subsp. saprophyticus		0.13	0.03		99	16	7	
021653	**Excellent identification**							**021653**
					OXI	NOV	PYR	
Aerococcus viridans		99.76	0.98		1	1	95	
Kocuria kristinae		0.14	0.17		99	1	93	
Staphylococcus lentus		0.06	0.18		99	99	6	
021660	**Very good identification**							**021660**
Aerococcus viridans		100.0	0.61	LAC 90				
021661	**Very good identification**							**021661**
					OXI			
Aerococcus viridans		96.52	0.69	LAC 90	1			
Kocuria kristinae		3.48	0.17		99			
021662	**Very good identification**							**021662**
					NOV	YEP		
Aerococcus viridans		99.78	0.81		1	1		
Staphylococcus arlettae		0.22	0.15		99	99		
021663	**Very good identification**							**021663**
					OXI	NOV	YEP	
Aerococcus viridans		99.77	0.88		1	1	1	
Staphylococcus arlettae		0.13	0.18		1	99	99	
Kocuria kristinae		0.10	0.05		99	1	15	
021670	**Very good identification**							**021670**
					NOV	YEP		
Aerococcus viridans		98.85	0.71	LAC 90	1	1		
Staphylococcus arlettae		0.67	0.15		99	99		
Staphylococcus kloosii		0.27	0.19		99	99		
021671	**Very good identification**							**021671**
					OXI	NOV	YEP	
Aerococcus viridans		99.16	0.79	LAC 90	1	1	1	
Staphylococcus arlettae		0.38	0.18		1	99	99	
Staphylococcus sciuri		0.31	0.11		99	99	57	
021672	**Good identification**							**021672**
					NOV	YEP		
Aerococcus viridans		93.07	0.91		1	1		
Staphylococcus arlettae		6.90	0.57		99	99		
Staphylococcus kloosii		0.03	0.20		99	99		
021673	**Very good identification**							**021673**
					OXI	NOV	YEP	
Aerococcus viridans		95.90	0.98		1	1	1	
Staphylococcus arlettae		4.04	0.59		1	99	99	
Staphylococcus lentus		0.06	0.18		99	99	1	
021700	**Acceptable identification**							**021700**
					OXI	NOV		
Macrococcus carouselicus		93.52	0.51	BGA 1	99	99		
Aerococcus viridans		5.86	0.49		1	1		
Rothia mucilaginosa		0.61	0.03		1	1		
021701	**Low discrimination**							**021701**
					OXI			
Aerococcus viridans		57.74	0.56	LAC 90	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		30.50	0.29		99			
Rothia mucilaginosa		7.77	0.13		1			
021702	**Very good identification**							**021702**
					OXI	NOV		
Aerococcus viridans		98.24	0.68		1	1		
Macrococcus carouselicus		1.76	0.10		99	99		
021703	**Very good identification**							**021703**
					OXI	NOV	PYR	
Aerococcus viridans		98.21	0.76		1	1	95	
Kocuria kristinae		1.44	0.16		99	1	93	
Staphylococcus lentus		0.35	0.12		99	99	6	
021710	**Low discrimination**							**021710**
					OXI	NOV		
Macrococcus carouselicus		76.26	0.47	BGA 1	99	99		
Aerococcus viridans		22.43	0.59		1	1		
Macrococcus bovicus		0.92	0.12		99	99		
021711	**Low discrimination**							**021711**
					OXI	NOV		
Aerococcus viridans		75.04	0.66	LAC 90	1	1		
Staphylococcus lentus		22.68	0.41		99	99		
Macrococcus carouselicus		1.10	0.06		99	99		
021712	**Very good identification**							**021712**
					OXI	NOV		
Aerococcus viridans		99.34	0.78		1	1		
Macrococcus carouselicus		0.38	0.06		99	99		
Staphylococcus lentus		0.28	0.12		99	99		
021713	**Low discrimination**							**021713**
					OXI	NOV		
Aerococcus viridans		89.36	0.86		1	1		
Staphylococcus lentus		10.64	0.53		99	99		
021720	**Low discrimination**							**021720**
					OXI	NOV		
Aerococcus viridans		86.12	0.49	LAC 90	1	1		
Macrococcus carouselicus		13.88	0.10		99	99		
021721	**Very good identification**							**021721**
Aerococcus viridans		100.0	0.56	LAC 90				
021722	**Very good identification**							**021722**
Aerococcus viridans		100.0	0.68					
021723	**Very good identification**							**021723**
					OXI	NOV		
Aerococcus viridans		99.64	0.76		1	1		
Staphylococcus lentus		0.36	0.12		99	99		
021730	**Very good identification**							**021730**
					OXI	NOV		
Aerococcus viridans		95.85	0.59	LAC 90	1	1		
Macrococcus carouselicus		3.29	0.06		99	99		
Staphylococcus vitulinus		0.86	-.08		99	99		
021731	**Low discrimination**							**021731**
					OXI	NOV		
Aerococcus viridans		76.42	0.66	LAC 90	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		23.09	0.41		99	99		
Staphylococcus sciuri		0.49	0.05		99	99		
021732	**Very good identification**							**021732**
					OXI	NOV	YEP	
Aerococcus viridans		98.06	0.78		1	1	1	
Staphylococcus arlettae		1.66	0.31		1	99	99	
Staphylococcus lentus		0.28	0.12		99	99	1	
021733	**Low discrimination**							**021733**
					OXI	NOV	YEP	
Aerococcus viridans		88.60	0.86		1	1	1	
Staphylococcus lentus		10.55	0.53		99	99	1	
Staphylococcus arlettae		0.85	0.34		1	99	99	
021740	**Low discrimination**							**021740**
					OXI	NOV		
Aerococcus viridans		88.55	0.61	LAC 90	1	1		
Rothia mucilaginosa		5.41	0.10		1	1		
Macrococcus carouselicus		3.57	0.10		99	99		
021741	**Low discrimination**							**021741**
					OXI			
Kocuria kristinae		52.32	0.48	VPT 90	99			
Aerococcus viridans		44.02	0.69		1			
Rothia mucilaginosa		3.46	0.20		1			
021742	**Very good identification**							**021742**
					OXI			
Aerococcus viridans		99.92	0.81		1			
Kocuria kristinae		0.08	-.05		99			
021743	**Very good identification**							**021743**
					OXI	NOV	PYR	
Aerococcus viridans		96.63	0.88		1	1	95	
Kocuria kristinae		3.19	0.36		99	1	93	
Staphylococcus lentus		0.18	0.18		99	99	6	
021750	**Good identification**							**021750**
					OXI	NOV	YEP	
Aerococcus viridans		93.35	0.71	LAC 90	1	1	1	
Macrococcus bovicus		2.23	0.19		99	99	99	
Staphylococcus kloosii		1.97	0.37		1	99	99	
021751	**Low discrimination**							**021751**
					OXI	NOV	PYR	
Aerococcus viridans		84.90	0.79	LAC 90	1	1	95	
Staphylococcus lentus		13.02	0.48		99	99	6	
Kocuria kristinae		1.40	0.20		99	1	93	
021752	**Very good identification**							**021752**
					NOV	YEP		
Aerococcus viridans		99.19	0.91		1	1		
Staphylococcus arlettae		0.42	0.31		99	99		
Staphylococcus kloosii		0.25	0.38		99	99		
021753	**Good identification**							**021753**
					OXI	NOV	YEP	
Aerococcus viridans		94.04	0.98		1	1	1	
Staphylococcus lentus		5.68	0.59		99	99	1	
Staphylococcus arlettae		0.23	0.34		1	99	99	
021760	**Very good identification**							**021760**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.61	LAC 90				
021761	**Very good identification**							**021761**
					OXI	NOV	PYR	
Aerococcus viridans		98.36	0.69	LAC 90	1	1	95	
Kocuria kristinae		1.18	0.07		99	1	93	
Staphylococcus lentus		0.46	0.07		99	99	6	
021762	**Very good identification**							**021762**
					NOV	YEP		
Aerococcus viridans		98.74	0.81		1	1		
Staphylococcus arlettae		1.26	0.31		99	99		
021763	**Very good identification**							**021763**
					OXI	NOV	YEP	
Aerococcus viridans		99.07	0.88		1	1	1	
Staphylococcus arlettae		0.72	0.34		1	99	99	
Staphylococcus lentus		0.18	0.18		99	99	1	
021770	**Very good identification**							**021770**
					NOV	YEP		
Aerococcus viridans		95.51	0.71	LAC 90	1	1		
Staphylococcus arlettae		3.65	0.31		99	99		
Staphylococcus kloosii		0.50	0.25		99	99		
021771	**Low discrimination**							**021771**
					OXI	NOV	YEP	
Aerococcus viridans		84.54	0.79	LAC 90	1	1	1	
Staphylococcus lentus		12.97	0.48		99	99	1	
Staphylococcus arlettae		1.83	0.34		1	99	99	
021772	**Low discrimination**							**021772**
					OXI	NOV	YEP	
Aerococcus viridans		70.33	0.91		1	1	1	
Staphylococcus arlettae		29.53	0.72		1	99	99	
Staphylococcus lentus		0.10	0.18		99	99	1	
021773	**Low discrimination**							**021773**
					OXI	NOV	YEP	
Aerococcus viridans		76.93	0.98		1	1	1	
Staphylococcus arlettae		18.35	0.75		1	99	99	
Staphylococcus lentus		4.65	0.59		99	99	1	
022000	**Low discrimination**							**022000**
					OXI			
Kocuria varians		53.44	0.74		13			
Kocuria rosea		26.48	0.59		35			
Dermacoccus nishinomiyaensis		12.94	0.54		93			
022001	**Low discrimination**							**022001**
Kocuria varians		67.71	0.52	MNS 8				
Kocuria rosea		20.31	0.33					
Staphylococcus auricularis		5.82	0.29					
022002	**Good identification**							**022002**
Kocuria varians		94.77	0.63					
Staphylococcus auricularis		2.51	0.29					
Kocuria rosea		1.68	0.18					
022003	**Acceptable identification**							**022003**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		94.64	0.41	MNS 8				
Staphylococcus auricularis		3.21	0.10					
Staphylococcus carnosus subsp. carnosus		2.16	0.02					
022010	**Low discrimination**							**022010**
					NOV	YEP	OXI	
Kocuria rosea		69.25	0.39	BGA 1	1	1	35	
Kocuria varians		12.70	0.33	MAN 10	1	80	13	
Macrococcus bovicus		11.61	0.28		99	99	99	
022011	**Acceptable identification**							**022011**
Staphylococcus capitis subsp. capitis		92.73	0.42	BGA 1				
Kocuria rosea		4.86	0.13					
Kocuria varians		1.47	0.11					
022020	**Low discrimination**							**022020**
					OXI			
Kocuria varians		61.91	0.65		13			
Kocuria rosea		37.43	0.51		35			
Dermacoccus nishinomiyaensis		0.43	0.13		93			
022021	**Low discrimination**							**022021**
Kocuria varians		73.21	0.43	MNS 8				
Kocuria rosea		26.79	0.25					
022022	**Very good identification**							**022022**
Kocuria varians		97.88	0.54					
Kocuria rosea		2.12	0.10					
022023	**Good identification**							**022023**
Kocuria varians		100.0	0.32	MNS 8				
022030	**Low discrimination**							**022030**
					NOV	OXI		
Kocuria rosea		80.56	0.32	BGA 1	1	35		
Kocuria varians		12.12	0.24	MAN 10	1	13		
Staphylococcus vitulinus		7.33	0.09		99	99		
022040	**Low discrimination**							**022040**
Kocuria varians		56.45	0.61					
Staphylococcus auricularis		40.40	0.56					
Kocuria rosea		1.29	0.18					
022041	**Low discrimination**							**022041**
Kocuria varians		52.24	0.39	MNS 8				
Staphylococcus auricularis		47.76	0.37					
022042	**Low discrimination**							**022042**
Kocuria varians		78.01	0.49					
Staphylococcus auricularis		21.99	0.37					
022043	**Low discrimination**							**022043**
Kocuria varians		73.52	0.28	MNS 8				
Staphylococcus auricularis		26.48	0.17					

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
022050	**Low discrimination**							**022050**
					NOV	OXI		
Macrococcus bovicus		83.16	0.36	BGA 1	99	99		
Kocuria varians		8.56	0.20	NIT 10	1	13		
Staphylococcus auricularis		6.13	0.15		1	1		
022060	**Good identification**							**022060**
Kocuria varians		95.38	0.51					
Kocuria rosea		2.65	0.10					
Staphylococcus auricularis		1.96	0.15					
022061	**Good identification**							**022061**
Kocuria varians		100.0	0.29	MNS 8				
022062	**Good identification**							**022062**
Kocuria varians		100.0	0.40					
022100	**Very good identification**							**022100**
					OXI			
Kocuria varians		96.20	0.78		13			
Dermacoccus nishinomiyaensis		2.96	0.39		93			
Staphylococcus auricularis		0.48	0.29		1			
022101	**Very good identification**							**022101**
					OXI			
Kocuria varians		99.02	0.56	MNS 8	13			
Staphylococcus auricularis		0.63	0.10		1			
Dermacoccus nishinomiyaensis		0.35	-.02		93			
022102	**Very good identification**							**022102**
					OXI			
Kocuria varians		99.69	0.67		13			
Staphylococcus auricularis		0.20	0.10		1			
Dermacoccus nishinomiyaensis		0.11	-.02		93			
022103	**Good identification**							**022103**
Kocuria varians		100.0	0.45	MNS 8				
022110	**Low discrimination**							**022110**
					NOV	YEP	OXI	
Kocuria varians		75.27	0.37	MAN 1	1	80	13	
Macrococcus bovicus		19.65	0.21		99	99	99	
Kocuria rosea		2.76	-.02		1	1	35	
022120	**Very good identification**							**022120**
					OXI			
Kocuria varians		99.51	0.68		13			
Kocuria rosea		0.41	0.10		35			
Dermacoccus nishinomiyaensis		0.09	-.02		93			
022121	**Good identification**							**022121**
Kocuria varians		100.0	0.47	MNS 8				
022122	**Very good identification**							**022122**
Kocuria varians		100.0	0.57					
022123	**Good identification**							**022123**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.35	MNS 8				
022130	**Good identification**							**022130**
Kocuria varians		100.0	0.27	MAN 1				
022140	**Good identification**							**022140**
					OXI			
Kocuria varians		94.84	0.64		13			
Staphylococcus auricularis		5.03	0.37		1			
Dermacoccus nishinomiyaensis		0.13	-.02		93			
022141	**Acceptable identification**							**022141**
Kocuria varians		93.66	0.42	MNS 8				
Staphylococcus auricularis		6.34	0.17					
022142	**Very good identification**							**022142**
Kocuria varians		97.96	0.53					
Staphylococcus auricularis		2.05	0.17					
022143	**Good identification**							**022143**
Kocuria varians		100.0	0.31	MNS 8				
022150	**Low discrimination**							**022150**
					NOV	PYR	OXI	
Macrococcus bovicus		70.09	0.28	BGA 1	99	1	99	
Kocuria varians		25.26	0.23	NIT 10	1	10	13	
Staphylococcus saprophyticus subsp. bovis		4.65	0.05		99	99	1	
022160	**Very good identification**							**022160**
Kocuria varians		100.0	0.55					
022161	**Good identification**							**022161**
Kocuria varians		100.0	0.33	MNS 8				
022162	**Good identification**							**022162**
Kocuria varians		100.0	0.44					
022200	**Low discrimination**							**022200**
Kocuria varians		49.38	0.56	SUC 12				
Staphylococcus auricularis		22.50	0.48					
Kocuria rosea		19.94	0.39					
022201	**Low discrimination**							**022201**
Kocuria varians		39.39	0.35	SUC 12				
Staphylococcus capitis subsp. capitis		25.70	0.24	MNS 8				
Staphylococcus auricularis		22.94	0.29					
022202	**Low discrimination**							**022202**
Kocuria varians		83.75	0.45	SUC 12				
Staphylococcus auricularis		15.04	0.29					
Kocuria rosea		1.21	-.02					
022210	**Low discrimination**							**022210**
					NOV	YEP		
Staphylococcus vitulinus		87.26	0.41	BGA 1	99	75		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		6.80	0.20		1	1	
Macrococcus bovicus		2.56	0.16		99	99	
022211	**Good identification**						**022211**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		98.93	0.47	BGA 1	1	1	
Staphylococcus vitulinus		0.76	0.00		99	99	
Kocuria rosea		0.31	-.07		1	35	
022220	**Low discrimination**						**022220**
					NOV	OXI	
Kocuria varians		64.50	0.47	SUC 12	1	13	
Kocuria rosea		31.77	0.32		1	35	
Staphylococcus vitulinus		2.89	0.09		99	99	
022222	**Good identification**						**022222**
Kocuria varians		100.0	0.36	SUC 12			
022230	**Good identification**						**022230**
					NOV	OXI	
Staphylococcus vitulinus		98.56	0.50	BGA 1	99	99	
Kocuria rosea		1.22	0.12		1	35	
Kocuria varians		0.23	0.06		1	13	
022240	**Low discrimination**						**022240**
					PYR		
Staphylococcus auricularis		80.26	0.56	BGA 14	40		
Kocuria varians		16.57	0.43	TRE 92	10		
Staphylococcus haemolyticus		2.40	0.25		98		
022241	**Low discrimination**						**022241**
					OXI	PYR	
Staphylococcus auricularis		64.27	0.36	BGA 14	1	40	
Kocuria kristinae		25.35	0.21	TRE 92	99	93	
Kocuria varians		10.38	0.21	MNS 10	13	10	
022242	**Low discrimination**						**022242**
					PYR		
Staphylococcus auricularis		58.15	0.36	BGA 14	40		
Kocuria varians		30.47	0.32	TRE 92	10		
Staphylococcus haemolyticus		8.79	0.20	LAC 10	98		
022260	**Low discrimination**						**022260**
Kocuria varians		87.78	0.33	SUC 12			
Staphylococcus auricularis		12.22	0.15				
022300	**Very good identification**						**022300**
					OXI		
Kocuria varians		95.16	0.60	SUC 12	13		
Staphylococcus auricularis		3.21	0.29		1		
Dermacoccus nishinomiyaensis		1.37	0.14		93		
022301	**Acceptable identification**						**022301**
Kocuria varians		92.26	0.38	SUC 12			
Staphylococcus auricularis		3.98	0.09	MNS 8			
Rothia mucilaginosa		2.18	-.07				
022302	**Good identification**						**022302**
Kocuria varians		98.69	0.49	SUC 12			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		1.31	0.09				
022303	**Low discrimination**						**022303**
					PYR	COA	
Kocuria varians		69.52	0.27	SUC 12	10	1	
Staphylococcus intermedius		25.84	0.14	MNS 8	90	99	
Staphylococcus hyicus		4.64	-.07		1	24	
022320	**Good identification**						**022320**
Kocuria varians		100.0	0.51	SUC 12			
022321	**Good identification**						**022321**
Kocuria varians		100.0	0.29	SUC 12			
022322	**Good identification**						**022322**
Kocuria varians		100.0	0.39	SUC 12			
022340	**Low discrimination**						**022340**
					PYR		
Kocuria varians		66.27	0.46	SUC 12	10		
Staphylococcus auricularis		23.78	0.36		40		
Staphylococcus haemolyticus		7.21	0.26		98		
022342	**Low discrimination**						**022342**
					PYR		
Kocuria varians		64.45	0.35	SUC 12	10		
Staphylococcus haemolyticus		13.98	0.21		98		
Staphylococcus auricularis		9.11	0.16		40		
022360	**Good identification**						**022360**
Kocuria varians		100.0	0.37	SUC 12			
022400	**Low discrimination**						**022400**
Staphylococcus auricularis		88.60	0.71	BGA 14			
Kocuria varians		8.61	0.51				
Staphylococcus carnosus subsp. utilis		1.48	0.18				
022401	**Low discrimination**						**022401**
					NOV		
Staphylococcus auricularis		89.84	0.51	BGA 14	1		
Kocuria varians		6.84	0.29	MNS 10	1		
Staphylococcus cohnii subsp. cohnii		2.77	0.23		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
022402	**Low discrimination**							**022402**
Staphylococcus auricularis		80.21	0.51	BGA 14				
Kocuria varians		19.80	0.40	LAC 10				
022403	**Low discrimination**							**022403**
Staphylococcus auricularis		81.69	0.31	BGA 14				
Kocuria varians		15.78	0.18	MNS 10				
Staphylococcus simulans		2.54	-.01	LAC 10				
022410	**Low discrimination**							**022410**
					OXI	NOV	YEP	
Macrococcus bovicus		51.05	0.32	BGA 1	99	99	99	
Staphylococcus auricularis		28.84	0.30	NIT 10	1	1	1	
Staphylococcus cohnii subsp. cohnii		11.48	0.25		1	99	12	
022420	**Low discrimination**							**022420**
Kocuria varians		72.63	0.42	TRE 7				
Staphylococcus auricularis		21.48	0.30					
Kocuria rosea		5.89	0.10					
022422	**Low discrimination**							**022422**
Kocuria varians		89.57	0.30	TRE 7				
Staphylococcus auricularis		10.43	0.10					
022440	**Very good identification**							**022440**
					NOV			
Staphylococcus auricularis		98.63	0.78	BGA 14	1			
Kocuria varians		0.90	0.38		1			
Staphylococcus cohnii subsp. cohnii		0.42	0.33		99			
022441	**Good identification**							**022441**
					NOV			
Staphylococcus auricularis		94.78	0.59	BGA 14	1			
Staphylococcus cohnii subsp. cohnii		3.96	0.34	MNS 10	99			
Kocuria varians		0.68	0.16		1			
022442	**Very good identification**							**022442**
					PYR			
Staphylococcus auricularis		97.35	0.59	BGA 14	40			
Kocuria varians		2.26	0.26	LAC 10	10			
Aerococcus viridans		0.24	0.20		95			
022443	**Acceptable identification**							**022443**
					PYR			
Staphylococcus auricularis		92.35	0.39	BGA 14	40			
Aerococcus viridans		4.71	0.28	MNS 10	95			
Kocuria varians		1.68	0.04	LAC 10	10			
022450	**Low discrimination**							**022450**
					OXI	NOV	PYR	
Macrococcus bovicus		42.56	0.39	BGA 1	99	99	1	
Staphylococcus auricularis		24.04	0.37	NIT 10	1	1	40	
Staphylococcus saprophyticus subsp. bovis		19.96	0.34		1	99	99	
022451	**Low discrimination**							**022451**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		62.09	0.36	BGA 1	99	1		
Staphylococcus saprophyticus subsp. bovis		22.03	0.22	NIT 7	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		11.79	0.18		1	40	
022453	**Low discrimination**						**022453**
					NOV		
Aerococcus viridans		69.73	0.38	ESL 90	1		
Staphylococcus saprophyticus subsp. bovis		19.37	0.09	NIT 1	99		
Staphylococcus simulans		10.90	0.04		1		
022460	**Low discrimination**						**022460**
Staphylococcus auricularis		75.86	0.37	BGA 14			
Kocuria varians		24.14	0.28	XYL 1			
022463	**Good identification**						**022463**
Aerococcus viridans		100.0	0.28	ESL 90			
022472	**Good identification**						**022472**
Aerococcus viridans		100.0	0.30	ESL 90			
022473	**Good identification**						**022473**
Aerococcus viridans		100.0	0.38	ESL 90			
022500	**Low discrimination**						**022500**
					OXI		
Kocuria varians		56.52	0.55	TRE 7	13		
Staphylococcus auricularis		43.06	0.51		1		
Dermacoccus nishinomiyaensis		0.23	-.02		93		
022501	**Low discrimination**						**022501**
Kocuria varians		47.60	0.33	TRE 7			
Staphylococcus auricularis		46.34	0.31	MNS 8			
Staphylococcus schleiferi subsp. schleiferi		2.60	0.02				
022502	**Low discrimination**						**022502**
Kocuria varians		76.92	0.43	TRE 7			
Staphylococcus auricularis		23.09	0.31				
022520	**Good identification**						**022520**
Kocuria varians		97.86	0.45	TRE 7			
Staphylococcus auricularis		2.14	0.10				
022522	**Good identification**						**022522**
Kocuria varians		100.0	0.34	TRE 7			
022540	**Low discrimination**						**022540**
					NOV	OXI	
Staphylococcus auricularis		86.38	0.59	BGA 14	1	1	
Kocuria varians		10.67	0.41	GAL 10	1	13	
Macrococcus caseolyticus		1.89	0.14		99	99	
022541	**Low discrimination**						**022541**
Staphylococcus auricularis		83.29	0.39	BGA 14			
Kocuria varians		8.05	0.19	GAL 10			
Rothia mucilaginosa		3.67	0.01	MNS 10			
022542	**Low discrimination**						**022542**
					NOV	OXI	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		60.70	0.39	BGA 14	1	1		
Kocuria varians		19.03	0.30	GAL 10	1	13		
Macrococcus caseolyticus		17.95	0.18	LAC 10	99	99		
022550	**Low discrimination**							**022550**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		77.43	0.47	URE 99	1	99	99	
Macrococcus bovicus		17.69	0.32		99	99	1	
Staphylococcus auricularis		2.59	0.18		1	1	40	
022551	**Low discrimination**							**022551**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		87.89	0.34	URE 99	1	99		
Staphylococcus cohnii subsp. cohnii		6.88	0.16		1	1		
Staphylococcus lentus		3.03	0.07		99	6		
022552	**Low discrimination**							**022552**
					NOV	YEP		
Staphylococcus saprophyticus subsp. bovis		89.39	0.34	URE 99	99	80		
Aerococcus viridans		8.62	0.30		1	1		
Staphylococcus kloosii		1.99	0.18		99	99		
022560	**Low discrimination**							**022560**
Kocuria varians		81.12	0.32	TRE 7				
Staphylococcus auricularis		18.89	0.18					
022572	**Good identification**							**022572**
Aerococcus viridans		100.0	0.30	ESL 90				
022600	**Very good identification**							**022600**
					NOV	OXI		
Staphylococcus auricularis		98.12	0.70	BGA 14	1	1		
Kocuria varians		1.41	0.33		1	13		
Macrococcus carouselicus		0.25	0.10		99	99		
022601	**Low discrimination**							**022601**
					OXI	PYR		
Staphylococcus auricularis		84.35	0.50	BGA 14	1	40		
Kocuria kristinae		14.25	0.28	MNS 10	99	93		
Kocuria varians		0.95	0.11		13	10		
022602	**Acceptable identification**							**022602**
Staphylococcus auricularis		93.69	0.50	BGA 14				
Kocuria varians		3.42	0.22	LAC 10				
Staphylococcus simulans		2.51	0.17					
022603	**Low discrimination**							**022603**
					OXI			
Staphylococcus auricularis		44.94	0.31	BGA 14	1			
Staphylococcus simulans		36.27	0.27	MNS 10	1			
Kocuria kristinae		17.09	0.15	LAC 10	99			
022610	**Low discrimination**							**022610**
					OXI	NOV		
Staphylococcus vitulinus		77.54	0.36	BGA 1	99	99		
Staphylococcus auricularis		12.66	0.29		1	1		
Macrococcus bovicus		6.07	0.19		99	99		
022612	**Low discrimination**							**022612**
					OXI	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		84.05	0.29	URE 98	1	1	
Staphylococcus auricularis		7.93	0.09	ARG 92	1	1	
Staphylococcus vitulinus		4.41	-.05		99	99	
022613	**Acceptable identification**						**022613**
					OXI	NOV	
Staphylococcus simulans		92.10	0.40	URE 98	1	1	
Staphylococcus lentus		2.95	0.12	ARG 92	99	99	
Aerococcus viridans		2.76	0.25		1	1	
022630	**Good identification**						**022630**
Staphylococcus vitulinus		100.0	0.45	BGA 1			
022640	**Very good identification**						**022640**
Staphylococcus auricularis		95.77	0.78	BGA 14			
Staphylococcus haemolyticus		3.31	0.49				
Staphylococcus petrasii subsp. pragensis		0.47	0.13				
022641	**Low discrimination**						**022641**
					OXI		
Staphylococcus auricularis		60.00	0.58	BGA 14	1		
Kocuria kristinae		39.11	0.47	MNS 10	99		
Staphylococcus haemolyticus		0.58	0.17		1		
022642	**Low discrimination**						**022642**
					PYR		
Staphylococcus auricularis		83.40	0.58	BGA 14	40		
Staphylococcus haemolyticus		14.57	0.43	LAC 10	98		
Staphylococcus hominis subsp. hominis		0.60	0.16		1		
022643	**Low discrimination**						**022643**
					OXI		
Kocuria kristinae		52.85	0.35	VPT 90	99		
Staphylococcus auricularis		36.04	0.38	NIT 7	1		
Staphylococcus simulans		5.60	0.20		1		
022650	**Low discrimination**						**022650**
					NOV		
Staphylococcus saprophyticus subsp. bovis		49.90	0.47	URE 99	99		
Staphylococcus haemolyticus		27.83	0.44		2		
Staphylococcus auricularis		13.87	0.36		1		
022651	**Low discrimination**						**022651**
					OXI	NOV	
Staphylococcus saprophyticus subsp. bovis		61.38	0.34	URE 99	1	99	
Kocuria kristinae		20.38	0.19		99	1	
Staphylococcus auricularis		7.58	0.17		1	1	
022652	**Low discrimination**						**022652**
					NOV		
Staphylococcus haemolyticus		43.69	0.39	VPT 86	2		
Staphylococcus saprophyticus subsp. bovis		34.82	0.34	ARG 92	99		
Staphylococcus simulans		8.78	0.22		1		
022653	**Low discrimination**						**022653**
					NOV		
Staphylococcus simulans		51.79	0.33	URE 98	1		
Staphylococcus saprophyticus subsp. bovis		15.33	0.22	ARG 92	99		
Aerococcus viridans		13.80	0.38		1		
022660	**Acceptable identification**						**022660**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					PYR				
Staphylococcus auricularis		92.46	0.36	BGA 14	40				
Kocuria varians		4.35	0.10	XYL 1	10				
Staphylococcus haemolyticus		3.19	0.07		98				
022663	**Low discrimination**								**022663**
					OXI				
Aerococcus viridans		78.84	0.28	ESL 90	1				
Kocuria kristinae		21.16	-.06	NIT 1	99				
022672	**Low discrimination**								**022672**
					NOV	YEP			
Aerococcus viridans		59.99	0.30	ESL 90	1	1			
Staphylococcus arlettae		40.01	0.15	NIT 1	99	99			
022673	**Low discrimination**								**022673**
					OXI	NOV			
Aerococcus viridans		40.37	0.38	ESL 90	1	1			
Staphylococcus xylosus		22.39	0.12	NIT 1	1	99			
Staphylococcus lentus		21.95	0.18		99	99			
022700	**Low discrimination**								**022700**
Staphylococcus auricularis		72.11	0.50	BGA 14					
Kocuria varians		13.98	0.37	GAL 10					
Rothia mucilaginosa		12.60	0.24						
022701	**Low discrimination**								**022701**
					OXI				
Rothia mucilaginosa		68.67	0.34	VPT 95	1				
Staphylococcus auricularis		14.55	0.31	ESL 90	1				
Kocuria kristinae		7.38	0.18		99				
022702	**Low discrimination**								**022702**
					PYR				
Staphylococcus auricularis		65.19	0.31	BGA 14	40				
Kocuria varians		32.09	0.26	GAL 10	10				
Staphylococcus haemolyticus		2.72	0.03	LAC 10	98				
022703	**Low discrimination**								**022703**
					OXI	PYR	COA		
Staphylococcus intermedius		74.95	0.37	URE 90	1	90	99		
Staphylococcus hyicus		11.65	0.15	PHS 97	1	1	24		
Kocuria kristinae		3.38	0.06		99	93	1		
022711	**Low discrimination**								**022711**
					OXI	COA	NOV	PYR	
Staphylococcus lentus		67.48	0.41	ESL 99	99	1	99	6	
Staphylococcus intermedius		28.85	0.29		1	99	1	90	
Staphylococcus saprophyticus subsp. bovis		1.62	0.05		1	1	99	99	
022713	**Low discrimination**								**022713**
					OXI	COA	NOV		
Staphylococcus lentus		56.05	0.53	ESL 99	99	1	99		
Staphylococcus intermedius		41.53	0.46		1	99	1		
Staphylococcus simulans		1.54	0.18		1	1	1		
022720	**Low discrimination**								**022720**
Kocuria varians		87.09	0.28	SUC 12					
Staphylococcus auricularis		12.91	0.09	TRE 7					
022731	**Good identification**								**022731**

Code	Taxon	ID score	T index	tests against	Differentiation tests				Code
					OXI				
Staphylococcus lentus		98.06	0.41	ESL 99	99				
Staphylococcus xylosus		1.94	0.00		1				
022733	**Good identification**								**022733**
					OXI	NOV			
Staphylococcus lentus		96.82	0.53	ESL 99	99	99			
Staphylococcus xylosus		1.54	0.10		1	99			
Aerococcus viridans		0.91	0.25		1	1			
022740	**Low discrimination**								**022740**
Staphylococcus auricularis		63.09	0.58	BGA 14					
Staphylococcus haemolyticus		22.10	0.50	GAL 10					
Rothia mucilaginosa		11.03	0.32						
022741	**Low discrimination**								**022741**
					OXI				
Rothia mucilaginosa		58.01	0.42	VPT 95	1				
Kocuria kristinae		24.03	0.38	ESL 90	99				
Staphylococcus auricularis		12.29	0.38		1				
022742	**Low discrimination**								**022742**
					OXI	NOV			
Staphylococcus haemolyticus		55.57	0.44	VPT 86	1	2			
Staphylococcus auricularis		31.34	0.38	ARG 92	1	1			
Macrococcus caseolyticus		6.69	0.14		99	99			
022743	**Low discrimination**								**022743**
					OXI	COA			
Staphylococcus intermedius		67.26	0.40	URE 90	1	99			
Kocuria kristinae		18.22	0.25	PHS 97	99	1			
Staphylococcus auricularis		4.14	0.19		1	1			
022750	**Low discrimination**								**022750**
					OXI	NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		84.39	0.59	URE 99	1	99	99		
Staphylococcus haemolyticus		13.27	0.45		1	2	98		
Macrococcus bovicus		1.21	0.19		99	99	1		
022751	**Low discrimination**								**022751**
					OXI	COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		44.94	0.47	URE 99	1	1	99	99	
Staphylococcus lentus		38.39	0.48		99	1	99	6	
Staphylococcus intermedius		12.13	0.33		1	99	1	90	
022752	**Low discrimination**								**022752**
					OXI	NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		69.90	0.47	URE 99	1	99	99		
Staphylococcus haemolyticus		24.72	0.40		1	2	98		
Staphylococcus lentus		2.14	0.18		99	99	6		
022753	**Low discrimination**								**022753**
					OXI	COA	NOV	PYR	
Staphylococcus lentus		59.90	0.59	ESL 99	99	1	99	6	
Staphylococcus intermedius		32.78	0.49		1	99	1	90	
Staphylococcus saprophyticus subsp. bovis		4.95	0.34		1	1	99	99	
022771	**Low discrimination**								**022771**
					OXI				
Staphylococcus lentus		85.60	0.48	ESL 99	99				
Staphylococcus xylosus		11.11	0.23		1				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		1.65	0.11		99			
022772	**Low discrimination**							**022772**
					OXI	NOV	YEP	
Staphylococcus arlettae		58.44	0.31	GLR 92	1	99	99	
Staphylococcus lentus		19.62	0.18	NIT 1	99	99	1	
Aerococcus viridans		15.46	0.30		1	1	1	
022773	**Low discrimination**							**022773**
					OXI	YEP		
Staphylococcus lentus		83.26	0.59	ESL 99	99	1		
Staphylococcus xylosus		8.67	0.33		1	40		
Staphylococcus arlettae		3.32	0.34		1	99		
023000	**Low discrimination**							**023000**
					OXI			
Kocuria varians		51.07	0.33	ESL 1	13			
Kocuria rosea		25.31	0.18		35			
Dermacoccus nishinomiyaensis		12.37	0.13		93			
023053	**Good identification**							**023053**
Aerococcus viridans		100.0	0.31	NIT 1				
023073	**Good identification**							**023073**
Aerococcus viridans		100.0	0.31	NIT 1				
023100	**Acceptable identification**							**023100**
					OXI	NOV		
Kocuria varians		94.68	0.37	ESL 1	13	1		
Dermacoccus nishinomiyaensis		2.91	-.02		93	80		
Macrococcus carouselicus		2.41	-.04		99	99		
023120	**Good identification**							**023120**
Kocuria varians		100.0	0.27	ESL 1				
023210	**Good identification**							**023210**
Staphylococcus vitulinus		99.45	0.50	BGA 1				
Macrococcus carouselicus		0.55	0.06					
023230	**Very good identification**							**023230**
Staphylococcus vitulinus		100.0	0.59	BGA 1				
023311	**Good identification**							**023311**
Staphylococcus lentus		95.94	0.41	TRE 99				
Staphylococcus sciuri		4.06	0.11					
023313	**Good identification**							**023313**
					OXI	NOV		
Staphylococcus lentus		99.56	0.53	TRE 99	99	99		
Aerococcus viridans		0.45	0.18		1	1		
023331	**Good identification**							**023331**
Staphylococcus lentus		96.91	0.41	TRE 99				
Staphylococcus sciuri		3.09	0.09					
023333	**Good identification**							**023333**
					OXI	NOV		
Staphylococcus lentus		99.56	0.53	TRE 99	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		0.45	0.18		1	1	
023351	**Acceptable identification**						**023351**
Staphylococcus lentus		92.73	0.48	TRE 99			
Staphylococcus sciuri		7.27	0.23				
023353	**Very good identification**						**023353**
					OXI	NOV	
Staphylococcus lentus		98.78	0.59	TRE 99	99	99	
Aerococcus viridans		0.87	0.31		1	1	
Staphylococcus sciuri		0.36	0.07		99	99	
023371	**Acceptable identification**						**023371**
Staphylococcus lentus		94.42	0.48	TRE 99			
Staphylococcus sciuri		5.58	0.21				
023373	**Very good identification**						**023373**
					OXI	NOV	
Staphylococcus lentus		98.86	0.59	TRE 99	99	99	
Aerococcus viridans		0.87	0.31		1	1	
Staphylococcus sciuri		0.27	0.04		99	99	
023400	**Low discrimination**						**023400**
					NOV	OXI	
Macrococcus carouselicus		62.18	0.28	BGA 1	99	99	
Staphylococcus auricularis		34.47	0.30	NIT 6	1	1	
Kocuria varians		3.35	0.10		1	13	
023402	**Low discrimination**						**023402**
Aerococcus viridans		55.77	0.27	NIT 1			
Staphylococcus auricularis		44.23	0.10				
023403	**Good identification**						**023403**
Aerococcus viridans		100.0	0.35	NIT 1			
023412	**Good identification**						**023412**
Aerococcus viridans		100.0	0.37	NIT 1			
023413	**Low discrimination**						**023413**
					OXI	NOV	
Aerococcus viridans		89.36	0.45	NIT 1	1	1	
Staphylococcus lentus		10.64	0.12		99	99	
023422	**Good identification**						**023422**
Aerococcus viridans		100.0	0.27	NIT 1			
023423	**Good identification**						**023423**
Aerococcus viridans		100.0	0.35	NIT 1			
023432	**Good identification**						**023432**
Aerococcus viridans		100.0	0.37	NIT 1			
023433	**Low discrimination**						**023433**
					OXI	NOV	
Aerococcus viridans		89.36	0.45	NIT 1	1	1	
Staphylococcus lentus		10.64	0.12		99	99	
023440	**Good identification**						**023440**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		97.40	0.37	BGA 14			
Aerococcus viridans		2.60	0.20	ESL 1			
023442	**Low discrimination**						**023442**
Aerococcus viridans		68.37	0.40	NIT 1			
Staphylococcus auricularis		31.63	0.18				
023443	**Good identification**						**023443**
Aerococcus viridans		100.0	0.47	NIT 1			
023451	**Low discrimination**						**023451**
					OXI	NOV	
Aerococcus viridans		73.88	0.38	NIT 1	1	1	
Staphylococcus sciuri		14.79	0.07	LAC 90	99	99	
Staphylococcus lentus		11.33	0.07		99	99	
023452	**Good identification**						**023452**
Aerococcus viridans		100.0	0.50	NIT 1			
023453	**Good identification**						**023453**
					OXI	NOV	
Aerococcus viridans		94.30	0.57	NIT 1	1	1	
Staphylococcus lentus		5.70	0.18		99	99	
023461	**Good identification**						**023461**
Aerococcus viridans		100.0	0.28	NIT 1			
023462	**Good identification**						**023462**
Aerococcus viridans		100.0	0.40	NIT 1			
023463	**Good identification**						**023463**
Aerococcus viridans		100.0	0.47	NIT 1			
023470	**Good identification**						**023470**
Aerococcus viridans		100.0	0.30	NIT 1			
023471	**Low discrimination**						**023471**
					OXI	NOV	
Aerococcus viridans		76.66	0.38	NIT 1	1	1	
Staphylococcus lentus		11.76	0.07	LAC 90	99	99	
Staphylococcus sciuri		11.58	0.05		99	99	
023472	**Good identification**						**023472**
Aerococcus viridans		100.0	0.50	NIT 1			
023473	**Good identification**						**023473**
					OXI	NOV	
Aerococcus viridans		94.30	0.57	NIT 1	1	1	
Staphylococcus lentus		5.70	0.18		99	99	
023502	**Low discrimination**						**023502**
					PYR		
Aerococcus viridans		77.31	0.27	NIT 1	95		
Kocuria varians		22.69	0.02		10		
023503	**Low discrimination**						**023503**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		73.68	0.35	NIT 1	1	1		
Staphylococcus lentus		26.32	0.12		99	99		
023511	**Good identification**							**023511**
					OXI	NOV		
Staphylococcus lentus		95.49	0.41	SUC 99	99	99		
Aerococcus viridans		3.19	0.25		1	1		
Staphylococcus sciuri		1.32	0.01		99	99		
023512	**Low discrimination**							**023512**
					OXI	NOV		
Aerococcus viridans		78.26	0.37	NIT 1	1	1		
Staphylococcus lentus		21.74	0.12		99	99		
023513	**Acceptable identification**							**023513**
					OXI	NOV		
Staphylococcus lentus		92.18	0.53	SUC 99	99	99		
Aerococcus viridans		7.82	0.45		1	1		
023522	**Good identification**							**023522**
Aerococcus viridans		100.0	0.27	NIT 1				
023523	**Low discrimination**							**023523**
					OXI	NOV		
Aerococcus viridans		73.68	0.35	NIT 1	1	1		
Staphylococcus lentus		26.32	0.12		99	99		
023531	**Good identification**							**023531**
					OXI	NOV		
Staphylococcus lentus		96.77	0.41	SUC 99	99	99		
Aerococcus viridans		3.23	0.25		1	1		
023532	**Low discrimination**							**023532**
					OXI	NOV		
Aerococcus viridans		78.26	0.37	NIT 1	1	1		
Staphylococcus lentus		21.74	0.12		99	99		
023533	**Acceptable identification**							**023533**
					OXI	NOV		
Staphylococcus lentus		92.18	0.53	SUC 99	99	99		
Aerococcus viridans		7.82	0.45		1	1		
023542	**Good identification**							**023542**
Aerococcus viridans		100.0	0.40	NIT 1				
023543	**Low discrimination**							**023543**
					OXI	NOV		
Aerococcus viridans		84.65	0.47	NIT 1	1	1		
Staphylococcus lentus		15.35	0.18		99	99		
023551	**Acceptable identification**							**023551**
					OXI	NOV		
Staphylococcus lentus		91.62	0.48	SUC 99	99	99		
Aerococcus viridans		6.03	0.38		1	1		
Staphylococcus sciuri		2.35	0.13		99	99		
023552	**Low discrimination**							**023552**
					OXI	NOV	YEP	
Aerococcus viridans		85.71	0.50	NIT 1	1	1	1	
Staphylococcus lentus		12.09	0.18		99	99	1	
Staphylococcus kloosii		2.20	0.18		1	99	99	
023553	**Low discrimination**							**023553**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Staphylococcus lentus		85.68	0.59	SUC 99	99	99		
Aerococcus viridans		14.32	0.57		1	1		
023561	**Low discrimination**							**023561**
					OXI	NOV		
Aerococcus viridans		68.49	0.28	NIT 1	1	1		
Staphylococcus lentus		31.51	0.07	LAC 90	99	99		
023562	**Good identification**							**023562**
Aerococcus viridans		100.0	0.40	NIT 1				
023563	**Low discrimination**							**023563**
					OXI	NOV		
Aerococcus viridans		84.65	0.47	NIT 1	1	1		
Staphylococcus lentus		15.35	0.18		99	99		
023570	**Low discrimination**							**023570**
					OXI	NOV		
Aerococcus viridans		73.64	0.30	NIT 1	1	1		
Staphylococcus lentus		26.36	0.07	LAC 90	99	99		
023571	**Acceptable identification**							**023571**
					OXI	NOV		
Staphylococcus lentus		92.15	0.48	SUC 99	99	99		
Aerococcus viridans		6.07	0.38		1	1		
Staphylococcus sciuri		1.78	0.11		99	99		
023572	**Low discrimination**							**023572**
					OXI	NOV		
Aerococcus viridans		87.64	0.50	NIT 1	1	1		
Staphylococcus lentus		12.36	0.18		99	99		
023573	**Low discrimination**							**023573**
					OXI	NOV		
Staphylococcus lentus		85.68	0.59	SUC 99	99	99		
Aerococcus viridans		14.32	0.57		1	1		
023600	**Low discrimination**							**023600**
					OXI	NOV		
Macrococcus carouselicus		74.72	0.34	BGA 1	99	99		
Staphylococcus auricularis		18.83	0.29	NIT 6	1	1		
Rothia mucilaginosa		3.29	0.03		1	1		
023602	**Low discrimination**							**023602**
Aerococcus viridans		57.74	0.27	NIT 1				
Staphylococcus auricularis		42.26	0.09					
023603	**Low discrimination**							**023603**
					OXI	NOV	PYR	
Aerococcus viridans		59.34	0.35	NIT 1	1	1	95	
Staphylococcus lentus		21.19	0.12		99	99	6	
Kocuria kristinae		19.46	0.03		99	1	93	
023610	**Low discrimination**							**023610**
Staphylococcus vitulinus		84.88	0.45	BGA 1				
Macrococcus carouselicus		13.01	0.30					
Staphylococcus sciuri		1.58	0.18					
023611	**Low discrimination**							**023611**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		56.01	0.41	GAL 99				
Staphylococcus sciuri		39.47	0.36					
Staphylococcus vitulinus		2.65	0.04					
023612	**Low discrimination**							**023612**
					OXI	NOV		
Aerococcus viridans		56.98	0.37	NIT 1	1	1		
Staphylococcus vitulinus		20.92	0.04		99	99		
Staphylococcus lentus		15.83	0.12		99	99		
023613	**Low discrimination**							**023613**
					OXI	NOV		
Staphylococcus lentus		89.51	0.53	GAL 99	99	99		
Aerococcus viridans		7.60	0.45		1	1		
Staphylococcus sciuri		2.90	0.20		99	99		
023622	**Good identification**							**023622**
Aerococcus viridans		100.0	0.27	NIT 1				
023623	**Low discrimination**							**023623**
					OXI	NOV		
Aerococcus viridans		73.68	0.35	NIT 1	1	1		
Staphylococcus lentus		26.32	0.12		99	99		
023630	**Very good identification**							**023630**
					OXI	NOV		
Staphylococcus vitulinus		99.37	0.54	BGA 1	99	99		
Staphylococcus sciuri		0.52	0.15		99	99		
Aerococcus viridans		0.11	0.18		1	1		
023631	**Low discrimination**							**023631**
Staphylococcus lentus		59.07	0.41	GAL 99				
Staphylococcus sciuri		31.40	0.34					
Staphylococcus vitulinus		7.56	0.13					
023632	**Low discrimination**							**023632**
					OXI	NOV		
Aerococcus viridans		44.04	0.37	NIT 1	1	1		
Staphylococcus vitulinus		43.72	0.13		99	99		
Staphylococcus lentus		12.23	0.12		99	99		
023633	**Acceptable identification**							**023633**
					OXI	NOV		
Staphylococcus lentus		90.15	0.53	GAL 99	99	99		
Aerococcus viridans		7.65	0.45		1	1		
Staphylococcus sciuri		2.20	0.18		99	99		
023640	**Low discrimination**							**023640**
Staphylococcus auricularis		79.60	0.36	BGA 14				
Rothia mucilaginosa		13.91	0.10	ESL 1				
Staphylococcus haemolyticus		2.75	0.07					
023641	**Low discrimination**							**023641**
					OXI	NOV	PYR	
Kocuria kristinae		67.62	0.35	VPT 90	99	1	93	
Rothia mucilaginosa		19.79	0.20	NIT 7	1	1	75	
Staphylococcus sciuri		4.68	0.17		99	99	1	
023642	**Low discrimination**							**023642**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		66.60	0.40	NIT 1				
Staphylococcus auricularis		28.44	0.17					
Staphylococcus haemolyticus		4.97	0.02					
023643	**Low discrimination**							**023643**
					OXI	NOV	PYR	
Aerococcus viridans		51.22	0.47	NIT 1	1	1	95	
Kocuria kristinae		37.80	0.23		99	1	93	
Staphylococcus lentus		9.28	0.18		99	99	6	
023650	**Low discrimination**							**023650**
					OXI	NOV		
Staphylococcus sciuri		62.36	0.30	BGA 1	99	99		
Aerococcus viridans		10.91	0.30	MNS 89	1	1		
Staphylococcus vitulinus		9.01	0.04		99	99		
023651	**Low discrimination**							**023651**
					OXI	NOV		
Staphylococcus sciuri		54.50	0.48	BGA 1	99	99		
Staphylococcus lentus		41.74	0.48		99	99		
Aerococcus viridans		2.75	0.38		1	1		
023652	**Low discrimination**							**023652**
					OXI	NOV		
Aerococcus viridans		80.36	0.50	NIT 1	1	1		
Staphylococcus lentus		11.33	0.18		99	99		
Staphylococcus sciuri		8.31	0.13		99	99		
023653	**Low discrimination**							**023653**
					OXI	NOV		
Staphylococcus lentus		81.38	0.59	GAL 99	99	99		
Aerococcus viridans		13.60	0.57		1	1		
Staphylococcus sciuri		4.88	0.32		99	99		
023662	**Good identification**							**023662**
Aerococcus viridans		100.0	0.40	NIT 1				
023663	**Low discrimination**							**023663**
					OXI	NOV		
Aerococcus viridans		84.65	0.47	NIT 1	1	1		
Staphylococcus lentus		15.35	0.18		99	99		
023670	**Low discrimination**							**023670**
					OXI	NOV		
Staphylococcus sciuri		54.56	0.27	BGA 1	99	99		
Staphylococcus vitulinus		28.26	0.13	MNS 89	99	99		
Aerococcus viridans		12.65	0.30		1	1		
023671	**Low discrimination**							**023671**
					OXI	NOV		
Staphylococcus lentus		48.43	0.48	GAL 99	99	99		
Staphylococcus sciuri		47.70	0.46		99	99		
Aerococcus viridans		3.19	0.38		1	1		
023672	**Low discrimination**							**023672**
					OXI	NOV		
Aerococcus viridans		77.33	0.50	NIT 1	1	1		
Staphylococcus lentus		10.90	0.18		99	99		
Staphylococcus sciuri		6.03	0.11		99	99		
023673	**Low discrimination**							**023673**
					OXI	NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		81.16	0.59	GAL 99	99	99		
Aerococcus viridans		13.57	0.57		1	1		
Staphylococcus sciuri		3.67	0.30		99	99		
023700	**Acceptable identification**							**023700**
					OXI	NOV		
Rothia mucilaginosa		90.52	0.44	VPT 95	1	1		
Macrococcus carouselicus		8.90	0.27		99	99		
Staphylococcus auricularis		0.58	0.09		1	1		
023701	**Good identification**							**023701**
					OXI	NOV	PYR	
Rothia mucilaginosa		93.12	0.54	VPT 95	1	1	75	
Staphylococcus lentus		6.34	0.41		99	99	6	
Kocuria kristinae		0.28	0.06		99	1	93	
023703	**Acceptable identification**							**023703**
					OXI	NOV		
Staphylococcus lentus		93.44	0.53	MAN 99	99	99		
Rothia mucilaginosa		3.91	0.13		1	1		
Aerococcus viridans		2.64	0.35		1	1		
023710	**Low discrimination**							**023710**
Staphylococcus lentus		62.97	0.41	MNS 99				
Macrococcus carouselicus		19.38	0.23					
Staphylococcus sciuri		10.65	0.24					
023711	**Very good identification**							**023711**
					OXI	NOV		
Staphylococcus lentus		98.46	0.82		99	99		
Staphylococcus sciuri		1.36	0.42		99	99		
Rothia mucilaginosa		0.15	0.13		1	1		
023712	**Good identification**							**023712**
					OXI	NOV		
Staphylococcus lentus		95.77	0.53	MNS 99	99	99		
Aerococcus viridans		3.48	0.37		1	1		
Staphylococcus sciuri		0.74	0.07		99	99		
023713	**Excellent identification**							**023713**
					OXI	NOV		
Staphylococcus lentus		99.84	0.94		99	99		
Aerococcus viridans		0.09	0.45		1	1		
Staphylococcus sciuri		0.06	0.26		99	99		
023721	**Low discrimination**							**023721**
					OXI	NOV		
Staphylococcus lentus		84.72	0.41	MAN 99	99	99		
Rothia mucilaginosa		12.58	0.13		1	1		
Staphylococcus sciuri		2.70	0.09		99	99		
023722	**Low discrimination**							**023722**
					OXI	NOV		
Aerococcus viridans		54.55	0.27	NIT 1	1	1		
Staphylococcus lentus		45.46	0.12		99	99		
023723	**Good identification**							**023723**
					OXI	NOV		
Staphylococcus lentus		97.25	0.53	MAN 99	99	99		
Aerococcus viridans		2.75	0.35		1	1		
023730	**Low discrimination**							**023730**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		78.75	0.41	MNS 99				
Staphylococcus vitulinus		10.08	0.13					
Staphylococcus sciuri		10.04	0.21					
023731	**Very good identification**							**023731**
					OXI	NOV		
Staphylococcus lentus		98.91	0.82		99	99		
Staphylococcus sciuri		1.03	0.40		99	99		
Aerococcus viridans		0.03	0.25		1	1		
023732	**Good identification**							**023732**
					OXI	NOV		
Staphylococcus lentus		95.95	0.53	MNS 99	99	99		
Aerococcus viridans		3.49	0.37		1	1		
Staphylococcus sciuri		0.56	0.05		99	99		
023733	**Excellent identification**							**023733**
					OXI	NOV		
Staphylococcus lentus		99.84	0.94		99	99		
Aerococcus viridans		0.09	0.45		1	1		
Staphylococcus sciuri		0.05	0.24		99	99		
023740	**Good identification**							**023740**
Rothia mucilaginosa		98.63	0.52	VPT 95				
Staphylococcus auricularis		0.63	0.17					
Staphylococcus haemolyticus		0.22	0.09					
023741	**Good identification**							**023741**
					OXI	NOV	PYR	
Rothia mucilaginosa		92.88	0.61	VPT 95	1	1	75	
Staphylococcus lentus		5.50	0.48		99	99	6	
Kocuria kristinae		1.07	0.25		99	1	93	
023742	**Low discrimination**							**023742**
					OXI	NOV		
Aerococcus viridans		45.90	0.40	NIT 1	1	1		
Rothia mucilaginosa		30.83	0.10		1	1		
Staphylococcus lentus		19.41	0.18		99	99		
023743	**Low discrimination**							**023743**
					OXI	NOV		
Staphylococcus lentus		89.03	0.59	MAN 99	99	99		
Aerococcus viridans		4.96	0.47		1	1		
Rothia mucilaginosa		4.28	0.20		1	1		
023750	**Low discrimination**							**023750**
					OXI	NOV		
Staphylococcus lentus		68.47	0.48	MNS 99	99	99		
Staphylococcus sciuri		21.45	0.36		99	99		
Rothia mucilaginosa		3.89	0.10		1	1		
023751	**Very good identification**							**023751**
					OXI	NOV		
Staphylococcus lentus		97.26	0.89		99	99		
Staphylococcus sciuri		2.49	0.54		99	99		
Rothia mucilaginosa		0.17	0.20		1	1		
023752	**Good identification**							**023752**
					OXI	NOV		
Staphylococcus lentus		91.80	0.59	MNS 99	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		6.58	0.50		1	1		
Staphylococcus sciuri		1.32	0.19		99	99		
023753	**Excellent identification**							**023753**
					OXI	NOV		
Staphylococcus lentus		99.71	1.00		99	99		
Aerococcus viridans		0.17	0.57		1	1		
Staphylococcus sciuri		0.12	0.38		99	99		
023761	**Low discrimination**							**023761**
					OXI	NOV		
Staphylococcus lentus		79.21	0.48	MAN 99	99	99		
Rothia mucilaginosa		13.52	0.20		1	1		
Staphylococcus sciuri		4.68	0.21		99	99		
023762	**Low discrimination**							**023762**
					OXI	NOV		
Aerococcus viridans		70.28	0.40	NIT 1	1	1		
Staphylococcus lentus		29.73	0.18		99	99		
023763	**Good identification**							**023763**
					OXI	NOV	PYR	
Staphylococcus lentus		93.68	0.59	MAN 99	99	99	6	
Aerococcus viridans		5.22	0.47		1	1	95	
Staphylococcus gallinarum		0.69	0.08		1	99	12	
023770	**Low discrimination**							**023770**
					OXI	NOV		
Staphylococcus lentus		79.08	0.48	MNS 99	99	99		
Staphylococcus sciuri		18.69	0.33		99	99		
Aerococcus viridans		2.23	0.30		1	1		
023771	**Very good identification**							**023771**
					OXI			
Staphylococcus lentus		97.77	0.89		99			
Staphylococcus sciuri		1.89	0.52		99			
Staphylococcus equorum		0.14	0.25		1			
023772	**Low discrimination**							**023772**
					OXI	NOV	YEP	
Staphylococcus lentus		89.88	0.59	MNS 99	99	99	1	
Aerococcus viridans		6.44	0.50		1	1	1	
Staphylococcus arlettae		2.70	0.31		1	99	99	
023773	**Excellent identification**							**023773**
					OXI	NOV	PYR	
Staphylococcus lentus		99.50	1.00		99	99	6	
Aerococcus viridans		0.17	0.57		1	1	95	
Staphylococcus equorum		0.10	0.34		1	99	5	
024000	**Low discrimination**							**024000**
					OXI			
Dermacoccus nishinomiyaensis		60.38	0.33	BGA 1	93			
Micrococcus sp.		32.62	0.18	PHS 5	89			
Kocuria varians		5.51	0.19		13			
024050	**Low discrimination**							**024050**
					OXI			
Staphylococcus kloosii		89.04	0.52	TRE 99	1			
Macrococcus bovicus		10.96	0.14		99			
024052	**Good identification**							**024052**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.53	TRE 99		
024070	**Good identification**					**024070**
Staphylococcus kloosii		100.0	0.40	TRE 99		
024072	**Good identification**					**024072**
Staphylococcus kloosii		100.0	0.40	TRE 99		
024110	**Low discrimination**					**024110**
					OXI	
Staphylococcus kloosii		81.88	0.32	TRE 99	1	
Macrococcus bovicus		18.12	-.01	MLT 95	99	
024112	**Good identification**					**024112**
Staphylococcus kloosii		100.0	0.32	TRE 99		
024150	**Very good identification**					**024150**
					OXI	
Staphylococcus kloosii		97.35	0.58	TRE 99	1	
Macrococcus bovicus		2.65	0.07		99	
024152	**Very good identification**					**024152**
Staphylococcus kloosii		100.0	0.59	TRE 99		
024170	**Good identification**					**024170**
Staphylococcus kloosii		100.0	0.46	TRE 99		
024172	**Good identification**					**024172**
Staphylococcus kloosii		100.0	0.46	TRE 99		
024240	**Very good identification**					**024240**
Staphylococcus aureus subsp. anaerobius		100.0	0.57	BGA 1		
024252	**Good identification**					**024252**
Staphylococcus kloosii		100.0	0.32	SUC 9		
024313	**Good identification**					**024313**
Staphylococcus intermedius		100.0	0.31	URE 90		
024350	**Good identification**					**024350**
Staphylococcus kloosii		100.0	0.37	SUC 9		
024352	**Good identification**					**024352**
Staphylococcus kloosii		100.0	0.38	SUC 9		
024353	**Good identification**					**024353**
Staphylococcus intermedius		100.0	0.34	URE 90		
024372	**Acceptable identification**					**024372**
Staphylococcus arlettae		90.92	0.35	GLR 92		
Staphylococcus kloosii		9.08	0.26	TRE 99		
024373	**Good identification**					**024373**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		100.0	0.37	GLR 92				
024410	**Good identification**							**024410**
					OXI	YEP		
Staphylococcus kloosii		93.94	0.67	MLT 95	1	99		
Staphylococcus cohnii subsp. cohnii		4.64	0.27		1	12		
Macrococcus bovicus		1.43	0.10		99	99		
024411	**Low discrimination**							**024411**
					YEP			
Staphylococcus cohnii subsp. cohnii		84.11	0.28	BGA 1	12			
Staphylococcus kloosii		15.89	0.26	PHS 9	99			
024412	**Very good identification**							**024412**
					NOV	YEP		
Staphylococcus kloosii		99.60	0.67	MLT 95	99	99		
Aerococcus viridans		0.41	0.18		1	1		
024430	**Very good identification**							**024430**
Staphylococcus kloosii		100.0	0.54	MLT 95				
024432	**Very good identification**							**024432**
					NOV	YEP		
Staphylococcus kloosii		98.40	0.55	MLT 95	99	99		
Aerococcus viridans		1.60	0.18		1	1		
024440	**Low discrimination**							**024440**
					NOV	YEP		
Staphylococcus kloosii		56.58	0.52	MAN 99	99	99		
Staphylococcus cohnii subsp. cohnii		36.21	0.35		99	12		
Staphylococcus auricularis		7.22	0.18		1	1		
024441	**Good identification**							**024441**
Staphylococcus cohnii subsp. cohnii		100.0	0.36	BGA 1				
024442	**Good identification**							**024442**
					NOV	YEP		
Staphylococcus kloosii		97.25	0.53	MAN 99	99	99		
Aerococcus viridans		2.75	0.20		1	1		
024443	**Good identification**							**024443**
Aerococcus viridans		100.0	0.28	ESL 90				
024450	**Very good identification**							**024450**
					OXI	YEP		
Staphylococcus kloosii		99.00	0.93		1	99		
Staphylococcus cohnii subsp. cohnii		0.82	0.38		1	12		
Macrococcus bovicus		0.19	0.18		99	99		
024451	**Low discrimination**							**024451**
					NOV	YEP	PYR	
Staphylococcus kloosii		52.46	0.52	MNS 1	99	99	52	
Staphylococcus cohnii subsp. cohnii		46.29	0.38		99	12	1	
Aerococcus viridans		1.25	0.18		1	1	95	
024452	**Excellent identification**							**024452**
					NOV	YEP		
Staphylococcus kloosii		99.91	0.94		99	99		
Aerococcus viridans		0.09	0.30		1	1		
024453	**Low discrimination**							**024453**
					NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		78.47	0.53	MNS 1	99	99	
Aerococcus viridans		15.53	0.38		1	1	
Staphylococcus cohnii subsp. urealyticum		6.00	0.18		99	64	
024460	**Good identification**						**024460**
Staphylococcus kloosii		100.0	0.40	MAN 99			
024462	**Low discrimination**						**024462**
					NOV	YEP	
Staphylococcus kloosii		89.84	0.40	MAN 99	99	99	
Aerococcus viridans		10.16	0.20		1	1	
024463	**Good identification**						**024463**
Aerococcus viridans		100.0	0.28	ESL 90			
024470	**Very good identification**						**024470**
Staphylococcus kloosii		100.0	0.81				
024471	**Acceptable identification**						**024471**
					NOV	YEP	
Staphylococcus kloosii		91.30	0.40	MNS 1	99	99	
Aerococcus viridans		8.70	0.18		1	1	
024472	**Very good identification**						**024472**
					NOV	YEP	
Staphylococcus kloosii		99.32	0.81		99	99	
Aerococcus viridans		0.34	0.30		1	1	
Staphylococcus arlettae		0.34	0.19		99	99	
024473	**Low discrimination**						**024473**
					NOV	YEP	
Staphylococcus kloosii		44.61	0.40	MNS 1	99	99	
Aerococcus viridans		35.32	0.38		1	1	
Staphylococcus arlettae		20.07	0.22		99	99	
024500	**Low discrimination**						**024500**
					YEP		
Staphylococcus kloosii		81.98	0.32	MAN 99	99		
Staphylococcus cohnii subsp. cohnii		18.02	0.05	MLT 95	12		
024502	**Good identification**						**024502**
Staphylococcus kloosii		100.0	0.32	MAN 99			
024510	**Very good identification**						**024510**
					OXI	YEP	
Staphylococcus kloosii		99.39	0.73	MLT 95	1	99	
Macrococcus bovicus		0.33	0.03		99	99	
Staphylococcus cohnii subsp. cohnii		0.28	0.08		1	12	
024511	**Low discrimination**						**024511**
					YEP		
Staphylococcus kloosii		76.74	0.32	MLT 95	99		
Staphylococcus cohnii subsp. cohnii		23.26	0.08	MNS 1	12		
024512	**Very good identification**						**024512**
					NOV	YEP	
Staphylococcus kloosii		99.79	0.73	MLT 95	99	99	
Aerococcus viridans		0.21	0.18		1	1	
024513	**Low discrimination**						**024513**
					COA	NOV	YEP

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		41.33	0.32	MLT 95	1	99	99	
Staphylococcus intermedius		38.66	0.13	MNS 1	99	1	1	
Aerococcus viridans		20.01	0.25		1	1	1	
024530	**Very good identification**							**024530**
Staphylococcus kloosii		100.0	0.60	MLT 95				
024532	**Very good identification**							**024532**
					NOV	YEP		
Staphylococcus kloosii		99.17	0.61	MLT 95	99	99		
Aerococcus viridans		0.83	0.18		1	1		
024540	**Very good identification**							**024540**
					YEP			
Staphylococcus kloosii		96.47	0.58	MAN 99	99			
Staphylococcus cohnii subsp. cohnii		3.53	0.16		12			
024542	**Very good identification**							**024542**
					NOV	YEP		
Staphylococcus kloosii		98.56	0.59	MAN 99	99	99		
Aerococcus viridans		1.44	0.20		1	1		
024543	**Low discrimination**							**024543**
					COA	NOV	YEP	
Aerococcus viridans		47.60	0.28	ESL 90	1	1	1	
Staphylococcus intermedius		38.26	0.07	PHS 1	99	1	1	
Staphylococcus kloosii		14.15	0.18		1	99	99	
024550	**Excellent identification**							**024550**
					OXI	YEP		
Staphylococcus kloosii		99.91	0.99		1	99		
Staphylococcus cohnii subsp. cohnii		0.05	0.18		1	12		
Macrococcus bovicus		0.04	0.10		99	99		
024551	**Good identification**							**024551**
					COA	YEP	NOV	
Staphylococcus kloosii		92.95	0.58	MNS 1	1	99	99	
Staphylococcus cohnii subsp. cohnii		4.70	0.19		1	12	99	
Staphylococcus intermedius		1.21	0.00		99	1	1	
024552	**Excellent identification**							**024552**
					NOV	YEP		
Staphylococcus kloosii		99.96	1.00		99	99		
Aerococcus viridans		0.04	0.30		1	1		
024553	**Low discrimination**							**024553**
					COA	NOV	YEP	
Staphylococcus kloosii		83.84	0.59	MNS 1	1	99	99	
Aerococcus viridans		8.55	0.38		1	1	1	
Staphylococcus intermedius		6.19	0.16		99	1	1	
024560	**Good identification**							**024560**
Staphylococcus kloosii		100.0	0.46	MAN 99				
024562	**Acceptable identification**							**024562**
					NOV	YEP		
Staphylococcus kloosii		94.49	0.46	MAN 99	99	99		
Aerococcus viridans		5.51	0.20		1	1		
024570	**Very good identification**							**024570**
Staphylococcus kloosii		100.0	0.87					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
024571	**Good identification**							**024571**
					NOV	YEP		
Staphylococcus kloosii		95.32	0.46	MNS 1	99	99		
Aerococcus viridans		4.68	0.18		1	1		
024572	**Very good identification**							**024572**
					NOV	YEP		
Staphylococcus kloosii		98.84	0.87		99	99		
Staphylococcus arlettae		0.99	0.35		99	99		
Aerococcus viridans		0.17	0.30		1	1		
024573	**Low discrimination**							**024573**
					COA	NOV		
Staphylococcus arlettae		38.13	0.37	GLR 92	1	99		
Staphylococcus kloosii		29.03	0.46	SUC 99	1	99		
Staphylococcus lutrae		21.00	0.18		99	1		
024610	**Good identification**							**024610**
					OXI			
Staphylococcus kloosii		96.30	0.46	SUC 9	1			
Macrococcus bovicus		3.70	-.02	MLT 95	99			
024612	**Good identification**							**024612**
					NOV	YEP		
Staphylococcus kloosii		96.05	0.47	SUC 9	99	99		
Aerococcus viridans		3.95	0.18	MLT 95	1	1		
024630	**Good identification**							**024630**
Staphylococcus kloosii		100.0	0.34	SUC 9				
024632	**Low discrimination**							**024632**
					NOV	YEP		
Staphylococcus kloosii		54.85	0.34	SUC 9	99	99		
Staphylococcus arlettae		36.12	0.19	MLT 95	99	99		
Aerococcus viridans		9.03	0.18		1	1		
024641	**Good identification**							**024641**
Kocuria kristinae		100.0	0.30	VPT 90				
024642	**Low discrimination**							**024642**
					NOV	YEP		
Staphylococcus kloosii		77.77	0.32	SUC 9	99	99		
Aerococcus viridans		22.23	0.20	MAN 99	1	1		
024650	**Very good identification**							**024650**
					OXI			
Staphylococcus kloosii		99.53	0.72	SUC 9	1			
Macrococcus bovicus		0.47	0.05		99			
024651	**Low discrimination**							**024651**
					OXI	COA	NOV	
Staphylococcus kloosii		51.92	0.31	SUC 9	1	1	99	
Kocuria kristinae		22.30	0.01	MNS 1	99	1	1	
Staphylococcus intermedius		13.28	0.00		1	99	1	
024652	**Very good identification**							**024652**
					NOV	YEP	PYR	
Staphylococcus kloosii		97.37	0.73	SUC 9	99	99	52	
Staphylococcus saprophyticus subsp. saprophyticus		0.94	0.03		99	16	7	
Aerococcus viridans		0.84	0.30		1	1	95	
024653	**Low discrimination**							**024653**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
					COA	NOV	YEP	
Aerococcus viridans		35.81	0.38	ESL 90	1	1	1	
Staphylococcus intermedius		25.94	0.16	PHS 1	99	1	1	
Staphylococcus arlettae		20.35	0.22		1	99	99	
024670	**Very good identification**							**024670**
Staphylococcus kloosii		96.38	0.60	SUC 9				
Staphylococcus arlettae		3.62	0.19					
024672	**Low discrimination**							**024672**
					NOV	YEP		
Staphylococcus arlettae		76.84	0.60	GLR 92	99	99		
Staphylococcus kloosii		22.39	0.61		99	99		
Aerococcus viridans		0.78	0.30		1	1		
024673	**Very good identification**							**024673**
					NOV	YEP		
Staphylococcus arlettae		98.04	0.63	GLR 92	99	99		
Aerococcus viridans		1.74	0.38		1	1		
Staphylococcus kloosii		0.22	0.20		99	99		
024701	**Acceptable identification**							**024701**
					OXI	COA		
Staphylococcus intermedius		94.24	0.29	URE 90	1	99		
Kocuria kristinae		5.76	0.00	NIT 93	99	1		
024703	**Good identification**							**024703**
					PYR			
Staphylococcus intermedius		99.26	0.45	URE 90	90			
Staphylococcus hyicus		0.74	-.05	NIT 93	1			
024710	**Good identification**							**024710**
Staphylococcus kloosii		100.0	0.52	SUC 9				
024711	**Good identification**							**024711**
Staphylococcus intermedius		100.0	0.37	URE 90				
024712	**Low discrimination**							**024712**
					COA	NOV	YEP	
Staphylococcus kloosii		89.55	0.53	SUC 9	1	99	99	
Staphylococcus intermedius		8.56	0.13	MLT 95	99	1	1	
Aerococcus viridans		1.90	0.18		1	1	1	
024713	**Very good identification**							**024713**
					COA			
Staphylococcus intermedius		99.13	0.54	URE 90	99			
Aerococcus viridans		0.52	0.25	NIT 93	1			
Staphylococcus felis		0.35	-.04		1			
024730	**Good identification**							**024730**
Staphylococcus kloosii		100.0	0.40	SUC 9				
024732	**Low discrimination**							**024732**
					NOV	YEP		
Staphylococcus arlettae		63.93	0.35	GLR 92	99	99		
Staphylococcus kloosii		33.25	0.40	MLT 99	99	99		
Aerococcus viridans		2.82	0.18		1	1		
024733	**Low discrimination**							**024733**
					COA	NOV	YEP	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		81.46	0.37	GLR 92	1	99	99	
Staphylococcus intermedius		12.22	0.13	MLT 99	99	1	1	
Aerococcus viridans		6.32	0.25		1	1	1	
024740	**Good identification**							**024740**
Staphylococcus kloosii		100.0	0.37	SUC 9				
024741	**Low discrimination**							**024741**
					OXI	COA		
Staphylococcus intermedius		73.16	0.32	URE 90	1	99		
Kocuria kristinae		26.85	0.20	NIT 93	99	1		
024742	**Low discrimination**							**024742**
					COA	NOV	YEP	
Staphylococcus kloosii		70.25	0.38	SUC 9	1	99	99	
Staphylococcus intermedius		19.40	0.07	MAN 99	99	1	1	
Aerococcus viridans		10.35	0.20		1	1	1	
024743	**Good identification**							**024743**
					OXI	COA		
Staphylococcus intermedius		97.32	0.48	URE 90	1	99		
Kocuria kristinae		1.45	0.07	NIT 93	99	1		
Aerococcus viridans		1.22	0.28		1	1		
024750	**Very good identification**							**024750**
					COA	NOV		
Staphylococcus kloosii		99.56	0.78	SUC 9	1	99		
Staphylococcus saprophyticus subsp. bovis		0.20	0.05		1	99		
Staphylococcus intermedius		0.13	0.00		99	1		
024751	**Acceptable identification**							**024751**
					COA	NOV	YEP	
Staphylococcus intermedius		92.06	0.41	URE 90	99	1	1	
Staphylococcus kloosii		7.06	0.37	NIT 93	1	99	99	
Aerococcus viridans		0.88	0.18	LAC 86	1	1	1	
024752	**Very good identification**							**024752**
					COA	NOV		
Staphylococcus kloosii		96.40	0.79	SUC 9	1	99		
Staphylococcus arlettae		2.44	0.35		1	99		
Staphylococcus intermedius		0.73	0.16		99	1		
024753	**Good identification**							**024753**
					COA	NOV	YEP	
Staphylococcus intermedius		93.09	0.57	URE 90	99	1	1	
Staphylococcus arlettae		4.18	0.37	NIT 93	1	99	99	
Aerococcus viridans		1.30	0.38		1	1	1	
024762	**Low discrimination**							**024762**
					NOV	YEP		
Staphylococcus arlettae		86.31	0.35	GLR 92	99	99		
Staphylococcus kloosii		8.62	0.26	MAN 99	99	99		
Aerococcus viridans		5.08	0.20		1	1		
024763	**Low discrimination**							**024763**
					COA	NOV	PYR	
Staphylococcus arlettae		68.01	0.37	GLR 92	1	99	1	
Staphylococcus lutrae		19.30	0.12	MAN 99	99	1	10	
Aerococcus viridans		7.04	0.28		1	1	95	
024770	**Good identification**							**024770**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		90.12	0.66	SUC 9				
Staphylococcus arlettae		9.88	0.35					
024771	**Low discrimination**							**024771**
					COA	NOV		
Staphylococcus arlettae		87.67	0.37	GLR 92	1	99		
Staphylococcus kloosii		6.09	0.25	LAC 99	1	99		
Staphylococcus intermedius		3.21	0.00		99	1		
024772	**Good identification**							**024772**
					NOV	YEP		
Staphylococcus arlettae		90.78	0.76	GLR 92	99	99		
Staphylococcus kloosii		9.06	0.67		99	99		
Aerococcus viridans		0.16	0.30		1	1		
024773	**Very good identification**							**024773**
					COA	NOV	PYR	
Staphylococcus arlettae		98.74	0.78	GLR 92	1	99	1	
Staphylococcus lutrae		0.55	0.18		99	1	10	
Aerococcus viridans		0.31	0.38		1	1	95	
025050	**Good identification**							**025050**
					OXI			
Staphylococcus kloosii		97.52	0.52	TRE 99	1			
Macrococcus equipersicus		2.49	0.02		99			
025052	**Good identification**							**025052**
					NOV	YEP		
Staphylococcus kloosii		96.14	0.53	TRE 99	99	99		
Aerococcus viridans		3.86	0.23		1	1		
025053	**Good identification**							**025053**
Aerococcus viridans		100.0	0.31	PHS 1				
025070	**Good identification**							**025070**
Staphylococcus kloosii		100.0	0.40	TRE 99				
025072	**Low discrimination**							**025072**
					NOV	YEP		
Staphylococcus kloosii		86.15	0.40	TRE 99	99	99		
Aerococcus viridans		13.85	0.23		1	1		
025073	**Good identification**							**025073**
Aerococcus viridans		100.0	0.31	PHS 1				
025110	**Good identification**							**025110**
Staphylococcus kloosii		100.0	0.32	TRE 99				
025112	**Good identification**							**025112**
Staphylococcus kloosii		100.0	0.32	TRE 99				
025150	**Very good identification**							**025150**
Staphylococcus kloosii		100.0	0.58	TRE 99				
025152	**Very good identification**							**025152**
					NOV	YEP		
Staphylococcus kloosii		97.97	0.59	TRE 99	99	99		
Aerococcus viridans		2.03	0.23		1	1		
025153	**Low discrimination**							**025153**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		82.70	0.31	PHS 1	1	1		
Staphylococcus kloosii		17.30	0.18	TRE 95	99	99		
025170	**Good identification**							**025170**
Staphylococcus kloosii		100.0	0.46	TRE 99				
025172	**Acceptable identification**							**025172**
					NOV	YEP		
Staphylococcus kloosii		92.35	0.46	TRE 99	99	99		
Aerococcus viridans		7.65	0.23		1	1		
025173	**Good identification**							**025173**
Aerococcus viridans		100.0	0.31	PHS 1				
025250	**Good identification**							**025250**
Staphylococcus kloosii		100.0	0.31	SUC 9				
025252	**Low discrimination**							**025252**
					NOV	YEP		
Staphylococcus kloosii		71.11	0.32	SUC 9	99	99		
Aerococcus viridans		28.89	0.23	TRE 99	1	1		
025253	**Good identification**							**025253**
Aerococcus viridans		100.0	0.31	PHS 1				
025350	**Good identification**							**025350**
Staphylococcus kloosii		100.0	0.37	SUC 9				
025352	**Low discrimination**							**025352**
					NOV	YEP		
Staphylococcus kloosii		82.69	0.38	SUC 9	99	99		
Aerococcus viridans		17.31	0.23	TRE 99	1	1		
025353	**Low discrimination**							**025353**
					OXI	NOV		
Aerococcus viridans		77.70	0.31	PHS 1	1	1		
Staphylococcus lentus		22.30	0.05	TRE 95	99	99		
025372	**Low discrimination**							**025372**
					NOV	YEP		
Staphylococcus arlettae		84.50	0.35	GLR 92	99	99		
Staphylococcus kloosii		8.44	0.26	TRE 99	99	99		
Aerococcus viridans		7.06	0.23		1	1		
025373	**Low discrimination**							**025373**
					OXI	NOV	YEP	
Staphylococcus arlettae		84.08	0.37	GLR 92	1	99	99	
Aerococcus viridans		12.37	0.31	TRE 99	1	1	1	
Staphylococcus lentus		3.55	0.05		99	99	1	
025402	**Low discrimination**							**025402**
					NOV	YEP		
Aerococcus viridans		54.73	0.27	PHS 1	1	1		
Staphylococcus kloosii		45.28	0.26		99	99		
025403	**Good identification**							**025403**
Aerococcus viridans		100.0	0.35	PHS 1				
025410	**Very good identification**							**025410**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	YEP	
Staphylococcus kloosii		97.88	0.67	MLT 95	1	99	99	
Macrococcus carouselicus		1.69	0.08		99	99	67	
Aerococcus viridans		0.43	0.18		1	1	1	
025412	**Very good identification**							**025412**
					NOV	YEP		
Staphylococcus kloosii		96.47	0.67	MLT 95	99	99		
Aerococcus viridans		3.53	0.37		1	1		
025413	**Low discrimination**							**025413**
					NOV	YEP		
Aerococcus viridans		89.43	0.45	PHS 1	1	1		
Staphylococcus kloosii		10.57	0.26		99	99		
025422	**Good identification**							**025422**
Aerococcus viridans		100.0	0.27	PHS 1				
025423	**Good identification**							**025423**
Aerococcus viridans		100.0	0.35	PHS 1				
025430	**Very good identification**							**025430**
					NOV	YEP		
Staphylococcus kloosii		98.27	0.54	MLT 95	99	99		
Aerococcus viridans		1.73	0.18		1	1		
025432	**Low discrimination**							**025432**
					NOV	YEP		
Staphylococcus kloosii		87.22	0.55	MLT 95	99	99		
Aerococcus viridans		12.78	0.37		1	1		
025433	**Good identification**							**025433**
Aerococcus viridans		100.0	0.45	PHS 1				
025440	**Good identification**							**025440**
					NOV	YEP		
Staphylococcus kloosii		97.03	0.52	MAN 99	99	99		
Aerococcus viridans		2.97	0.20		1	1		
025441	**Good identification**							**025441**
Aerococcus viridans		100.0	0.28	PHS 1				
025442	**Low discrimination**							**025442**
					NOV	YEP		
Staphylococcus kloosii		79.72	0.53	MAN 99	99	99		
Aerococcus viridans		20.29	0.40		1	1		
025443	**Good identification**							**025443**
Aerococcus viridans		100.0	0.47	PHS 1				
025450	**Very good identification**							**025450**
					OXI	NOV	YEP	
Staphylococcus kloosii		99.86	0.93		1	99	99	
Aerococcus viridans		0.09	0.30		1	1	1	
Macrococcus equipersicus		0.03	0.02		99	99	99	
025451	**Low discrimination**							**025451**
					NOV	YEP		
Staphylococcus kloosii		82.34	0.52	MNS 1	99	99		
Aerococcus viridans		17.66	0.38		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
025452	**Excellent identification**						**025452**
					NOV	YEP	
Staphylococcus kloosii		99.24	0.94		99	99	
Aerococcus viridans		0.77	0.50		1	1	
025453	**Low discrimination**						**025453**
					NOV	YEP	
Aerococcus viridans		63.37	0.57	PHS 1	1	1	
Staphylococcus kloosii		35.57	0.53		99	99	
Staphylococcus cohnii subsp. urealyticum		1.06	0.09		99	64	
025460	**Low discrimination**						**025460**
					NOV	YEP	
Staphylococcus kloosii		89.09	0.40	MAN 99	99	99	
Aerococcus viridans		10.92	0.20		1	1	
025461	**Good identification**						**025461**
Aerococcus viridans		100.0	0.28	PHS 1			
025462	**Low discrimination**						**025462**
					NOV	YEP	
Aerococcus viridans		50.44	0.40	PHS 1	1	1	
Staphylococcus kloosii		49.56	0.40		99	99	
025463	**Good identification**						**025463**
Aerococcus viridans		100.0	0.47	PHS 1			
025470	**Very good identification**						**025470**
					NOV	YEP	
Staphylococcus kloosii		99.63	0.81		99	99	
Aerococcus viridans		0.37	0.30		1	1	
025471	**Low discrimination**						**025471**
					NOV	YEP	
Staphylococcus kloosii		53.83	0.40	MNS 1	99	99	
Aerococcus viridans		46.17	0.38		1	1	
025472	**Very good identification**						**025472**
					NOV	YEP	
Staphylococcus kloosii		96.69	0.81		99	99	
Aerococcus viridans		2.98	0.50		1	1	
Staphylococcus arlettae		0.33	0.19		99	99	
025473	**Low discrimination**						**025473**
					NOV	YEP	
Aerococcus viridans		83.09	0.57	PHS 1	1	1	
Staphylococcus kloosii		11.66	0.40		99	99	
Staphylococcus arlettae		5.25	0.22		99	99	
025500	**Low discrimination**						**025500**
					OXI		
Staphylococcus kloosii		62.94	0.32	MAN 99	1		
Macrococcus carouselicus		37.07	0.04	MLT 95	99		
025502	**Low discrimination**						**025502**
					NOV	YEP	
Staphylococcus kloosii		61.63	0.32	MAN 99	99	99	
Aerococcus viridans		38.37	0.27	MLT 95	1	1	
025503	**Good identification**						**025503**
Aerococcus viridans		100.0	0.35	PHS 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
025510	**Very good identification**							**025510**
					OXI	NOV	YEP	
Staphylococcus kloosii		99.40	0.73	MLT 95	1	99	99	
Macrococcus carouselicus		0.38	0.00		99	99	67	
Aerococcus viridans		0.23	0.18		1	1	1	
025511	**Low discrimination**							**025511**
					NOV	YEP		
Staphylococcus kloosii		65.59	0.32	MLT 95	99	99		
Aerococcus viridans		34.41	0.25	MNS 1	1	1		
025512	**Very good identification**							**025512**
					NOV	YEP		
Staphylococcus kloosii		98.15	0.73	MLT 95	99	99		
Aerococcus viridans		1.85	0.37		1	1		
025513	**Low discrimination**							**025513**
					NOV	YEP		
Aerococcus viridans		81.34	0.45	PHS 1	1	1		
Staphylococcus kloosii		18.66	0.32		99	99		
025522	**Low discrimination**							**025522**
					NOV	YEP		
Aerococcus viridans		71.35	0.27	PHS 1	1	1		
Staphylococcus kloosii		28.65	0.20		99	99		
025523	**Good identification**							**025523**
Aerococcus viridans		100.0	0.35	PHS 1				
025530	**Very good identification**							**025530**
					NOV	YEP		
Staphylococcus kloosii		99.10	0.60	MLT 95	99	99		
Aerococcus viridans		0.90	0.18		1	1		
025532	**Good identification**							**025532**
					NOV	YEP		
Staphylococcus kloosii		92.98	0.61	MLT 95	99	99		
Aerococcus viridans		7.02	0.37		1	1		
025533	**Acceptable identification**							**025533**
					NOV	YEP		
Aerococcus viridans		94.58	0.45	PHS 1	1	1		
Staphylococcus kloosii		5.42	0.20		99	99		
025540	**Very good identification**							**025540**
					NOV	YEP		
Staphylococcus kloosii		98.45	0.58	MAN 99	99	99		
Aerococcus viridans		1.55	0.20		1	1		
025541	**Low discrimination**							**025541**
					NOV	YEP		
Aerococcus viridans		78.47	0.28	PHS 1	1	1		
Staphylococcus kloosii		21.53	0.17	LAC 90	99	99		
025542	**Low discrimination**							**025542**
					NOV	YEP		
Staphylococcus kloosii		88.41	0.59	MAN 99	99	99		
Aerococcus viridans		11.59	0.40		1	1		
025543	**Good identification**							**025543**
					NOV	YEP		
Aerococcus viridans		96.80	0.47	PHS 1	1	1		
Staphylococcus kloosii		3.20	0.18		99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
025550	**Excellent identification**							**025550**
					NOV	YEP		
Staphylococcus kloosii		99.95	0.99		99	99		
Aerococcus viridans		0.05	0.30		1	1		
025551	**Good identification**							**025551**
					NOV	YEP		
Staphylococcus kloosii		90.05	0.58	MNS 1	99	99		
Aerococcus viridans		9.95	0.38		1	1		
025552	**Excellent identification**							**025552**
					NOV	YEP		
Staphylococcus kloosii		99.60	1.00		99	99		
Aerococcus viridans		0.40	0.50		1	1		
025553	**Low discrimination**							**025553**
					OXI	NOV	YEP	
Staphylococcus kloosii		51.77	0.59	MNS 1	1	99	99	
Aerococcus viridans		47.51	0.57		1	1	1	
Staphylococcus lentus		0.72	0.05		99	99	1	
025560	**Acceptable identification**							**025560**
					NOV	YEP		
Staphylococcus kloosii		94.06	0.46	MAN 99	99	99		
Aerococcus viridans		5.94	0.20		1	1		
025561	**Good identification**							**025561**
Aerococcus viridans		100.0	0.28	PHS 1				
025562	**Low discrimination**							**025562**
					NOV	YEP		
Staphylococcus kloosii		65.60	0.46	MAN 99	99	99		
Aerococcus viridans		34.40	0.40		1	1		
025563	**Good identification**							**025563**
Aerococcus viridans		100.0	0.47	PHS 1				
025570	**Very good identification**							**025570**
					NOV	YEP		
Staphylococcus kloosii		99.81	0.87		99	99		
Aerococcus viridans		0.19	0.30		1	1		
025571	**Low discrimination**							**025571**
					NOV	YEP		
Staphylococcus kloosii		69.36	0.46	MNS 1	99	99		
Aerococcus viridans		30.64	0.38		1	1		
025572	**Very good identification**							**025572**
					NOV	YEP		
Staphylococcus kloosii		97.48	0.87		99	99		
Aerococcus viridans		1.55	0.50		1	1		
Staphylococcus arlettae		0.98	0.35		99	99		
025573	**Low discrimination**							**025573**
					NOV	YEP		
Aerococcus viridans		60.78	0.57	PHS 1	1	1		
Staphylococcus arlettae		21.75	0.37		99	99		
Staphylococcus kloosii		16.56	0.46		99	99		
025602	**Good identification**							**025602**
Aerococcus viridans		100.0	0.27	PHS 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
025603	**Good identification**							**025603**
Aerococcus viridans		100.0	0.35	PHS 1				
025610	**Low discrimination**							**025610**
					OXI	NOV	YEP	
Staphylococcus kloosii		71.52	0.46	SUC 9	1	99	99	
Macrococcus carouselicus		25.29	0.14	MLT 95	99	99	67	
Aerococcus viridans		3.19	0.18		1	1	1	
025612	**Low discrimination**							**025612**
					NOV	YEP		
Staphylococcus kloosii		72.97	0.47	SUC 9	99	99		
Aerococcus viridans		27.03	0.37	MLT 95	1	1		
025613	**Good identification**							**025613**
Aerococcus viridans		100.0	0.45	PHS 1				
025622	**Good identification**							**025622**
Aerococcus viridans		100.0	0.27	PHS 1				
025623	**Good identification**							**025623**
Aerococcus viridans		100.0	0.35	PHS 1				
025630	**Low discrimination**							**025630**
					OXI	NOV	YEP	
Staphylococcus kloosii		74.86	0.34	SUC 9	1	99	99	
Aerococcus viridans		13.35	0.18	MLT 95	1	1	1	
Staphylococcus vitulinus		11.80	-.08		99	99	75	
025632	**Low discrimination**							**025632**
					NOV	YEP		
Aerococcus viridans		47.18	0.37	PHS 1	1	1		
Staphylococcus kloosii		31.85	0.34		99	99		
Staphylococcus arlettae		20.98	0.19		99	99		
025633	**Low discrimination**							**025633**
					NOV	YEP		
Aerococcus viridans		79.84	0.45	PHS 1	1	1		
Staphylococcus arlettae		20.16	0.22		99	99		
025640	**Low discrimination**							**025640**
					NOV	YEP		
Staphylococcus kloosii		76.35	0.31	SUC 9	99	99		
Aerococcus viridans		23.65	0.20	MAN 99	1	1		
025642	**Low discrimination**							**025642**
					NOV	YEP		
Aerococcus viridans		72.01	0.40	PHS 1	1	1		
Staphylococcus kloosii		27.99	0.32		99	99		
025643	**Acceptable identification**							**025643**
					OXI			
Aerococcus viridans		90.99	0.47	PHS 1	1			
Kocuria kristinae		9.01	0.05		99			
025650	**Very good identification**							**025650**
					OXI	NOV	YEP	
Staphylococcus kloosii		98.89	0.72	SUC 9	1	99	99	
Aerococcus viridans		0.93	0.30		1	1	1	
Staphylococcus sciuri		0.18	-.01		99	99	57	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
025651	**Low discrimination**							**025651**
					OXI	NOV	YEP	
Aerococcus viridans		46.51	0.38	PHS 1	1	1	1	
Staphylococcus sciuri		32.04	0.18	LAC 90	99	99	57	
Staphylococcus kloosii		21.45	0.31		1	99	99	
025652	**Good identification**							**025652**
					NOV	YEP		
Staphylococcus kloosii		92.03	0.73	SUC 9	99	99		
Aerococcus viridans		7.18	0.50		1	1		
Staphylococcus arlettae		0.80	0.19		99	99		
025653	**Low discrimination**							**025653**
					NOV	YEP		
Aerococcus viridans		87.24	0.57	PHS 1	1	1		
Staphylococcus arlettae		5.51	0.22		99	99		
Staphylococcus kloosii		4.84	0.32		99	99		
025661	**Good identification**							**025661**
Aerococcus viridans		100.0	0.28	PHS 1				
025662	**Low discrimination**							**025662**
					NOV	YEP		
Aerococcus viridans		69.90	0.40	PHS 1	1	1		
Staphylococcus arlettae		23.31	0.19		99	99		
Staphylococcus kloosii		6.79	0.20		99	99		
025663	**Low discrimination**							**025663**
					NOV	YEP		
Aerococcus viridans		84.07	0.47	PHS 1	1	1		
Staphylococcus arlettae		15.93	0.22		99	99		
025670	**Good identification**							**025670**
					NOV	YEP		
Staphylococcus kloosii		93.02	0.60	SUC 9	99	99		
Staphylococcus arlettae		3.49	0.19		99	99		
Aerococcus viridans		3.49	0.30		1	1		
025671	**Low discrimination**							**025671**
					OXI	NOV	YEP	
Aerococcus viridans		45.39	0.38	PHS 1	1	1	1	
Staphylococcus arlettae		25.79	0.22	LAC 90	1	99	99	
Staphylococcus sciuri		23.59	0.16		99	99	57	
025672	**Low discrimination**							**025672**
					NOV	YEP		
Staphylococcus arlettae		72.34	0.60	GLR 92	99	99		
Staphylococcus kloosii		21.08	0.61		99	99		
Aerococcus viridans		6.58	0.50		1	1		
025673	**Low discrimination**							**025673**
					OXI	NOV	YEP	
Staphylococcus arlettae		85.87	0.63	GLR 92	1	99	99	
Aerococcus viridans		13.74	0.57		1	1	1	
Staphylococcus lentus		0.21	0.05		99	99	1	
025702	**Good identification**							**025702**
Aerococcus viridans		100.0	0.27	PHS 1				
025703	**Low discrimination**							**025703**
					COA			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		80.77	0.35	PHS 1	1			
Staphylococcus intermedius		19.24	0.04		99			
025710	**Acceptable identification**							**025710**
					OXI	NOV	YEP	
Staphylococcus kloosii		90.82	0.52	SUC 9	1	99	99	
Macrococcus carouselicus		7.09	0.06	MLT 95	99	99	67	
Aerococcus viridans		2.09	0.18		1	1	1	
025711	**Low discrimination**							**025711**
					OXI	NOV		
Staphylococcus lentus		75.54	0.29	NIT 99	99	99		
Staphylococcus sciuri		14.36	0.12		99	99		
Aerococcus viridans		10.10	0.25		1	1		
025712	**Low discrimination**							**025712**
					NOV	YEP		
Staphylococcus kloosii		83.98	0.53	SUC 9	99	99		
Aerococcus viridans		16.02	0.37	MLT 95	1	1		
025713	**Low discrimination**							**025713**
					OXI	COA	NOV	
Staphylococcus lentus		70.81	0.40	NIT 99	99	1	99	
Aerococcus viridans		24.03	0.45		1	1	1	
Staphylococcus intermedius		5.16	0.13		1	99	1	
025722	**Good identification**							**025722**
Aerococcus viridans		100.0	0.27	PHS 1				
025723	**Good identification**							**025723**
Aerococcus viridans		100.0	0.35	PHS 1				
025730	**Acceptable identification**							**025730**
					NOV	YEP		
Staphylococcus kloosii		91.59	0.40	SUC 9	99	99		
Aerococcus viridans		8.41	0.18	MLT 95	1	1		
025731	**Low discrimination**							**025731**
					OXI	NOV		
Staphylococcus lentus		78.30	0.29	NIT 99	99	99		
Staphylococcus sciuri		11.23	0.10		99	99		
Aerococcus viridans		10.47	0.25		1	1		
025732	**Low discrimination**							**025732**
					NOV	YEP		
Staphylococcus arlettae		52.16	0.35	GLR 92	99	99		
Staphylococcus kloosii		27.13	0.40	MLT 99	99	99		
Aerococcus viridans		20.71	0.37		1	1		
025733	**Low discrimination**							**025733**
					OXI	NOV	YEP	
Staphylococcus lentus		54.79	0.40	NIT 99	99	99	1	
Staphylococcus arlettae		26.61	0.37		1	99	99	
Aerococcus viridans		18.60	0.45		1	1	1	
025740	**Low discrimination**							**025740**
					NOV	YEP		
Staphylococcus kloosii		86.24	0.37	SUC 9	99	99		
Aerococcus viridans		13.76	0.20	MAN 99	1	1		
025742	**Low discrimination**							**025742**
					NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		57.00	0.40	PHS 1	1	1		
Staphylococcus kloosii		43.00	0.38		99	99		
025743	**Low discrimination**							**025743**
					OXI	COA	NOV	
Aerococcus viridans		85.64	0.47	PHS 1	1	1	1	
Staphylococcus intermedius		7.65	0.07		1	99	1	
Staphylococcus lentus		3.88	0.05		99	1	99	
025750	**Very good identification**							**025750**
					OXI	NOV	YEP	
Staphylococcus kloosii		99.34	0.78	SUC 9	1	99	99	
Aerococcus viridans		0.48	0.30		1	1	1	
Staphylococcus sciuri		0.19	0.05		99	99	57	
025751	**Low discrimination**							**025751**
					OXI	NOV		
Staphylococcus lentus		53.12	0.35	NIT 99	99	99		
Staphylococcus sciuri		18.71	0.24		99	99		
Aerococcus viridans		13.99	0.38		1	1		
025752	**Good identification**							**025752**
					NOV	YEP		
Staphylococcus kloosii		93.73	0.79	SUC 9	99	99		
Aerococcus viridans		3.77	0.50		1	1		
Staphylococcus arlettae		2.37	0.35		99	99		
025753	**Low discrimination**							**025753**
					OXI	NOV	YEP	
Staphylococcus lentus		48.78	0.47	NIT 99	99	99	1	
Aerococcus viridans		32.62	0.57		1	1	1	
Staphylococcus arlettae		11.67	0.37		1	99	99	
025761	**Low discrimination**							**025761**
					NOV	PYR		
Aerococcus viridans		67.23	0.28	PHS 1	1	95		
Staphylococcus gallinarum		32.77	-.01	LAC 90	99	12		
025762	**Low discrimination**							**025762**
					NOV	YEP		
Staphylococcus arlettae		61.39	0.35	GLR 92	99	99		
Aerococcus viridans		32.49	0.40	MAN 99	1	1		
Staphylococcus kloosii		6.13	0.26		99	99		
025763	**Low discrimination**							**025763**
					NOV	PYR		
Staphylococcus arlettae		47.68	0.37	GLR 92	99	1		
Aerococcus viridans		44.42	0.47	MAN 99	1	95		
Staphylococcus gallinarum		5.89	0.08		99	12		
025770	**Low discrimination**							**025770**
					NOV	YEP		
Staphylococcus kloosii		88.14	0.66	SUC 9	99	99		
Staphylococcus arlettae		9.66	0.35		99	99		
Aerococcus viridans		1.71	0.30		1	1		
025771	**Low discrimination**							**025771**
					OXI	YEP		
Staphylococcus lentus		33.98	0.35	NIT 99	99	1		
Staphylococcus arlettae		28.82	0.37		1	99		
Staphylococcus gallinarum		14.70	0.20		1	78		
025772	**Low discrimination**							**025772**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus arlettae		89.57	0.76	GLR 92	99	99		
Staphylococcus kloosii		8.94	0.67		99	99		
Aerococcus viridans		1.44	0.50		1	1		
025773	**Good identification**							**025773**
					OXI	NOV	YEP	
Staphylococcus arlettae		92.03	0.78	GLR 92	1	99	99	
Staphylococcus lentus		3.89	0.47		99	99	1	
Aerococcus viridans		2.60	0.57		1	1	1	
026000	**Low discrimination**							**026000**
					OXI			
Dermacoccus nishinomiyaensis		43.69	0.28	BGA 1	93			
Kocuria varians		34.62	0.33	PHS 5	13			
Kocuria rosea		17.16	0.18		35			
026001	**Low discrimination**							**026001**
Staphylococcus schleiferi subsp. schleiferi		53.69	0.33	VPT 99				
Staphylococcus carnosus subsp. carnosus		42.07	0.31	ARG 86				
Kocuria varians		2.67	0.11					
026003	**Good identification**							**026003**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	ARG 99				
026011	**Low discrimination**							**026011**
Staphylococcus carnosus subsp. carnosus		89.01	0.43	ARG 99				
Staphylococcus capitis subsp. capitis		10.99	0.23					
026013	**Very good identification**							**026013**
Staphylococcus carnosus subsp. carnosus		100.0	0.55	ARG 99				
026100	**Low discrimination**							**026100**
					OXI			
Kocuria varians		86.19	0.37	PHS 1	13			
Dermacoccus nishinomiyaensis		13.81	0.13		93			
026101	**Low discrimination**							**026101**
Staphylococcus schleiferi subsp. schleiferi		78.71	0.33	VPT 99				
Staphylococcus carnosus subsp. carnosus		15.42	0.18	ARG 86				
Kocuria varians		5.87	0.15					
026103	**Acceptable identification**							**026103**
					COA	PYR		
Staphylococcus carnosus subsp. carnosus		93.39	0.31	ARG 99	1	72		
Staphylococcus intermedius		4.11	0.04		99	90		
Kocuria varians		2.51	0.04		1	10		
026111	**Good identification**							**026111**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	ARG 99				
026113	**Good identification**							**026113**
					COA			
Staphylococcus carnosus subsp. carnosus		97.11	0.43	ARG 99	1			
Staphylococcus intermedius		2.89	0.13		99			
026120	**Good identification**							**026120**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.27	PHS 1				
026211	**Good identification**							**026211**
Staphylococcus capitis subsp. capitis		95.78	0.29	BGA 1				
Staphylococcus carnosus subsp. carnosus		4.22	0.02	PHS 11				
026301	**Good identification**							**026301**
					PYR			
Staphylococcus intermedius		96.70	0.29	URE 90	90			
Staphylococcus hyicus		3.30	-.07	TRE 93	1			
026303	**Acceptable identification**							**026303**
					YEP	PYR	COA	
Staphylococcus intermedius		91.93	0.45	URE 90	1	90	99	
Staphylococcus hyicus		5.87	0.15	TRE 93	1	1	24	
Staphylococcus chromogenes		2.21	0.10		95	45	1	
026311	**Good identification**							**026311**
Staphylococcus intermedius		100.0	0.37	URE 90				
026313	**Very good identification**							**026313**
					COA			
Staphylococcus intermedius		99.06	0.54	URE 90	99			
Staphylococcus felis		0.35	-.04	TRE 93	1			
Staphylococcus carnosus subsp. carnosus		0.34	0.02		1			
026341	**Low discrimination**							**026341**
					COA	PYR		
Staphylococcus intermedius		72.76	0.32	URE 90	99	90		
Staphylococcus epidermidis		27.24	0.26	TRE 93	1	1		
026342	**Acceptable identification**							**026342**
					COA	PYR		
Staphylococcus epidermidis		91.14	0.30	VPT 96	1	1		
Staphylococcus intermedius		8.86	0.07	URE 92	99	90		
026343	**Low discrimination**							**026343**
					COA	YEP	PYR	
Staphylococcus intermedius		82.71	0.48	URE 90	99	1	90	
Staphylococcus epidermidis		15.97	0.36	TRE 93	1	1	1	
Staphylococcus chromogenes		1.32	0.10		1	95	45	
026351	**Good identification**							**026351**
Staphylococcus intermedius		100.0	0.41	URE 90				
026353	**Very good identification**							**026353**
					OXI	COA	YEP	
Staphylococcus intermedius		99.65	0.57	URE 90	1	99	1	
Staphylococcus lentus		0.19	0.05	TRE 93	99	1	1	
Staphylococcus chromogenes		0.17	-.01		1	1	95	
026400	**Low discrimination**							**026400**
Staphylococcus auricularis		85.79	0.30	BGA 14				
Kocuria varians		8.34	0.10	PHS 1				
Staphylococcus schleiferi subsp. schleiferi		5.87	0.02					
026401	**Low discrimination**							**026401**
Staphylococcus schleiferi subsp. schleiferi		88.63	0.43	VPT 99				
Staphylococcus carnosus subsp. carnosus		9.92	0.23	ARG 86				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		1.45	0.10				
026403	**Acceptable identification**						**026403**
Staphylococcus carnosus subsp. carnosus		92.99	0.35	ARG 99			
Staphylococcus felis		4.92	0.03				
Staphylococcus schleiferi subsp. schleiferi		2.10	0.02				
026411	**Good identification**						**026411**
					NOV		
Staphylococcus carnosus subsp. carnosus		96.47	0.35	ARG 99	1		
Staphylococcus schleiferi subsp. schleiferi		2.18	0.02		1		
Staphylococcus cohnii subsp. cohnii		1.36	0.05		99		
026413	**Acceptable identification**						**026413**
Staphylococcus carnosus subsp. carnosus		91.53	0.48	ARG 99			
Staphylococcus felis		8.09	0.20				
Staphylococcus simulans		0.38	0.02				
026440	**Good identification**						**026440**
					NOV		
Staphylococcus auricularis		95.97	0.37	BGA 14	1		
Staphylococcus cohnii subsp. cohnii		4.03	0.12	PHS 1	99		
026450	**Low discrimination**						**026450**
					PYR	YEP	
Staphylococcus kloosii		75.75	0.52	NIT 1	52	99	
Staphylococcus saprophyticus subsp. bovis		18.05	0.22		99	80	
Staphylococcus cohnii subsp. cohnii		4.64	0.14		1	12	
026452	**Acceptable identification**						**026452**
Staphylococcus kloosii		94.79	0.53	NIT 1			
Staphylococcus saprophyticus subsp. bovis		5.21	0.09				
026470	**Good identification**						**026470**
Staphylococcus kloosii		100.0	0.40	NIT 1			
026472	**Good identification**						**026472**
Staphylococcus kloosii		100.0	0.40	NIT 1			
026501	**Acceptable identification**						**026501**
					COA		
Staphylococcus schleiferi subsp. schleiferi		94.98	0.43	VPT 99	1		
Staphylococcus carnosus subsp. carnosus		2.66	0.11	ARG 86	1		
Staphylococcus intermedius		2.36	0.11		99		
026503	**Low discrimination**						**026503**
					COA		
Staphylococcus intermedius		42.13	0.27	URE 90	99		
Staphylococcus carnosus subsp. carnosus		30.89	0.23	SUC 99	1		
Staphylococcus felis		21.87	0.14		1		
026510	**Good identification**						**026510**
Staphylococcus kloosii		100.0	0.32	NIT 1			
026512	**Good identification**						**026512**
Staphylococcus kloosii		100.0	0.32	NIT 1			
026513	**Low discrimination**						**026513**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA			
Staphylococcus felis		34.71	0.31	URE 99	1			
Staphylococcus carnosus subsp. carnosus		29.31	0.35	ARG 92	1			
Staphylococcus intermedius		27.02	0.36		99			
026543	**Low discrimination**							**026543**
					PYR			
Staphylococcus intermedius		79.58	0.30	URE 90	90			
Staphylococcus lutrae		20.43	0.12	SUC 99	10			
026550	**Low discrimination**							**026550**
Staphylococcus kloosii		67.07	0.58	NIT 1				
Staphylococcus saprophyticus subsp. bovis		32.93	0.34					
026552	**Low discrimination**							**026552**
					COA	NOV		
Staphylococcus kloosii		89.03	0.59	NIT 1	1	99		
Staphylococcus saprophyticus subsp. bovis		10.09	0.22		1	99		
Staphylococcus intermedius		0.88	-.02		99	1		
026553	**Low discrimination**							**026553**
					COA	PYR	NOV	
Staphylococcus intermedius		80.80	0.39	URE 90	99	90	1	
Staphylococcus lutrae		14.89	0.18	SUC 99	99	10	1	
Staphylococcus saprophyticus subsp. bovis		2.33	0.09		1	99	99	
026563	**Very good identification**							**026563**
Staphylococcus lutrae		100.0	0.53	URE 99				
026570	**Good identification**							**026570**
Staphylococcus kloosii		100.0	0.46	NIT 1				
026572	**Low discrimination**							**026572**
					COA	NOV		
Staphylococcus kloosii		58.03	0.46	NIT 1	1	99		
Staphylococcus lutrae		41.97	0.18		99	1		
026573	**Very good identification**							**026573**
					OXI	PYR	COA	
Staphylococcus lutrae		99.87	0.59	URE 99	1	10	99	
Staphylococcus lentus		0.08	0.05		99	6	1	
Staphylococcus intermedius		0.06	-.02		1	90	99	
026603	**Low discrimination**							**026603**
					COA	PYR		
Staphylococcus intermedius		43.25	0.27	URE 90	99	90		
Staphylococcus hyicus		20.58	0.15	GAL 99	24	1		
Staphylococcus felis		13.61	0.09		1	20		
026613	**Low discrimination**							**026613**
					COA			
Staphylococcus intermedius		43.85	0.36	URE 90	99			
Staphylococcus felis		34.16	0.26	GAL 99	1			
Staphylococcus simulans		19.03	0.30		1			
026620	**Very good identification**							**026620**
Staphylococcus muscae		100.0	0.59	BGA 1				
026640	**Good identification**							**026640**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		96.66	0.36	BGA 14				
Staphylococcus haemolyticus		3.34	0.07	PHS 1				
026643	**Low discrimination**							**026643**
					COA	YEP		
Staphylococcus intermedius		79.04	0.30	URE 90	99	1		
Staphylococcus chromogenes		12.03	0.12	GAL 99	1	95		
Staphylococcus simulans		6.94	0.10		1	1		
026650	**Low discrimination**							**026650**
					NOV			
Staphylococcus saprophyticus subsp. bovis		88.79	0.34	URE 99	99			
Staphylococcus kloosii		9.22	0.31		99			
Staphylococcus haemolyticus		2.00	0.03		2			
026653	**Low discrimination**							**026653**
					COA	NOV		
Staphylococcus intermedius		84.12	0.39	URE 90	99	1		
Staphylococcus simulans		10.93	0.23	GAL 99	1	1		
Staphylococcus saprophyticus subsp. bovis		2.43	0.09		1	99		
026701	**Good identification**							**026701**
					YEP	PYR	COA	
Staphylococcus intermedius		95.15	0.52	URE 90	1	90	99	
Staphylococcus hyicus		2.81	0.15	LAC 86	1	1	24	
Staphylococcus chromogenes		1.35	0.12		95	45	1	
026702	**Acceptable identification**							**026702**
					PYR			
Staphylococcus intermedius		94.76	0.27	URE 90	90			
Staphylococcus hyicus		5.24	-.05	MNS 99	1			
026703	**Good identification**							**026703**
					YEP	PYR	COA	
Staphylococcus intermedius		92.02	0.68	URE 90	1	90	99	
Staphylococcus hyicus		5.09	0.37		1	1	24	
Staphylococcus chromogenes		1.91	0.32		95	45	1	
026711	**Very good identification**							**026711**
					OXI	COA		
Staphylococcus intermedius		97.76	0.61	URE 90	1	99		
Staphylococcus lentus		1.77	0.29	LAC 86	99	1		
Staphylococcus sciuri		0.17	0.06		99	1		
026712	**Good identification**							**026712**
					COA			
Staphylococcus intermedius		97.43	0.36	URE 90	99			
Staphylococcus felis		2.57	-.04	MNS 99	1			
026713	**Very good identification**							**026713**
					OXI	COA	NOV	
Staphylococcus intermedius		95.73	0.77	URE 90	1	99	1	
Staphylococcus felis		2.52	0.37		1	1	1	
Staphylococcus lentus		1.00	0.40		99	1	99	
026723	**Low discrimination**							**026723**
					PYR			
Staphylococcus intermedius		69.43	0.27	URE 90	90			
Staphylococcus lutrae		26.73	0.12	XYL 1	10			
Staphylococcus hyicus		3.84	-.05		1			
026731	**Low discrimination**							**026731**
					OXI	COA		

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus lentus		61.37	0.29	ESL 99	99	1			
Staphylococcus intermedius		34.28	0.19		1	99			
Staphylococcus sciuri		4.36	0.04		99	1			
026733	**Low discrimination**								**026733**
					OXI	PYR	COA		
Staphylococcus lentus		43.88	0.40	ESL 99	99	6	1		
Staphylococcus intermedius		42.47	0.36		1	90	99		
Staphylococcus lutrae		11.74	0.18		1	10	99		
026741	**Very good identification**								**026741**
					COA	YEP			
Staphylococcus intermedius		98.37	0.55	URE 90	99	1			
Staphylococcus chromogenes		0.93	0.12	LAC 86	1	95			
Rothia mucilaginosa		0.49	0.01		1	1			
026742	**Good identification**								**026742**
					COA				
Staphylococcus intermedius		97.05	0.30	URE 90	99				
Staphylococcus haemolyticus		2.95	0.03	MNS 99	1				
026743	**Very good identification**								**026743**
					COA	PYR	YEP		
Staphylococcus intermedius		98.31	0.72	URE 90	99	90	1		
Staphylococcus chromogenes		1.36	0.32		1	45	95		
Staphylococcus lutrae		0.26	0.12		99	10	1		
026750	**Low discrimination**								**026750**
					COA	NOV			
Staphylococcus saprophyticus subsp. bovis		87.35	0.47	URE 99	1	99			
Staphylococcus intermedius		7.70	0.23		99	1			
Staphylococcus kloosii		4.40	0.37		1	99			
026751	**Good identification**								**026751**
					OXI	COA	NOV	PYR	
Staphylococcus intermedius		94.48	0.64	URE 90	1	99	1	90	
Staphylococcus saprophyticus subsp. bovis		2.71	0.34	LAC 86	1	1	99	99	
Staphylococcus lentus		2.31	0.35		99	1	99	6	
026752	**Low discrimination**								**026752**
					COA	NOV			
Staphylococcus intermedius		63.42	0.39	URE 90	99	1			
Staphylococcus saprophyticus subsp. bovis		29.29	0.34	MNS 99	1	99			
Staphylococcus kloosii		6.39	0.38		1	99			
026753	**Very good identification**								**026753**
					OXI	PYR	COA		
Staphylococcus intermedius		98.12	0.80	URE 90	1	90	99		
Staphylococcus lentus		1.39	0.47		99	6	1		
Staphylococcus lutrae		0.18	0.18		1	10	99		
026763	**Very good identification**								**026763**
					COA	PYR	NOV		
Staphylococcus lutrae		95.57	0.53	URE 99	99	10	1		
Staphylococcus intermedius		3.76	0.30		99	90	1		
Staphylococcus gallinarum		0.42	0.08		1	12	99		
026771	**Low discrimination**								**026771**
					OXI	COA	NOV		
Staphylococcus lentus		38.60	0.35	ESL 99	99	1	99		
Staphylococcus lutrae		18.03	0.18		1	99	1		
Staphylococcus gallinarum		16.70	0.20		1	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
026772	**Low discrimination**							**026772**
					COA	NOV		
Staphylococcus arlettae		51.93	0.35	GLR 92	1	99		
Staphylococcus lutrae		37.91	0.18	NIT 1	99	1		
Staphylococcus kloosii		5.18	0.26		1	99		
026773	**Low discrimination**							**026773**
					OXI	PYR	COA	
Staphylococcus lutrae		84.62	0.59	URE 99	1	10	99	
Staphylococcus lentus		6.49	0.47		99	6	1	
Staphylococcus intermedius		4.64	0.39		1	90	99	
027013	**Good identification**							**027013**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ARG 99				
027311	**Low discrimination**							**027311**
Staphylococcus lentus		77.63	0.29	TRE 99				
Staphylococcus sciuri		22.37	0.16					
027313	**Acceptable identification**							**027313**
					OXI	COA		
Staphylococcus lentus		92.08	0.40	TRE 99	99	1		
Staphylococcus intermedius		6.71	0.13		1	99		
Staphylococcus sciuri		1.22	0.00		99	1		
027331	**Low discrimination**							**027331**
Staphylococcus lentus		82.15	0.29	TRE 99				
Staphylococcus sciuri		17.85	0.14					
027333	**Good identification**							**027333**
Staphylococcus lentus		100.0	0.40	TRE 99				
027351	**Low discrimination**							**027351**
					OXI	COA		
Staphylococcus lentus		63.90	0.35	TRE 99	99	1		
Staphylococcus sciuri		34.11	0.28		99	1		
Staphylococcus intermedius		1.99	0.00		1	99		
027353	**Acceptable identification**							**027353**
					OXI	COA		
Staphylococcus lentus		92.74	0.47	TRE 99	99	1		
Staphylococcus intermedius		4.99	0.16		1	99		
Staphylococcus sciuri		2.27	0.12		99	1		
027371	**Low discrimination**							**027371**
					OXI			
Staphylococcus lentus		54.48	0.35	TRE 99	99			
Staphylococcus gallinarum		23.58	0.20		1			
Staphylococcus sciuri		21.94	0.25		99			
027373	**Low discrimination**							**027373**
					OXI			
Staphylococcus lentus		75.91	0.47	TRE 99	99			
Staphylococcus gallinarum		22.68	0.28		1			
Staphylococcus sciuri		1.40	0.09		99			
027413	**Good identification**							**027413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ARG 99				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
027450	**Good identification**						**027450**
Staphylococcus kloosii		100.0	0.52	NIT 1			
027452	**Good identification**						**027452**
Staphylococcus kloosii		100.0	0.53	NIT 1			
027470	**Good identification**						**027470**
Staphylococcus kloosii		100.0	0.40	NIT 1			
027472	**Good identification**						**027472**
Staphylococcus kloosii		100.0	0.40	NIT 1			
027510	**Good identification**						**027510**
Staphylococcus kloosii		100.0	0.32	NIT 1			
027511	**Low discrimination**						**027511**
					OXI	NOV	
Staphylococcus lentus		83.51	0.29	SUC 99	99	99	
Staphylococcus carnosus subsp. carnosus		8.64	0.03		1	1	
Staphylococcus sciuri		7.86	0.06		99	99	
027512	**Good identification**						**027512**
Staphylococcus kloosii		100.0	0.32	NIT 1			
027513	**Low discrimination**						**027513**
					OXI	NOV	
Staphylococcus lentus		89.55	0.40	SUC 99	99	99	
Staphylococcus carnosus subsp. carnosus		10.45	0.16		1	1	
027531	**Acceptable identification**						**027531**
Staphylococcus lentus		93.37	0.29	SUC 99			
Staphylococcus sciuri		6.63	0.04				
027533	**Good identification**						**027533**
Staphylococcus lentus		100.0	0.40	SUC 99			
027550	**Very good identification**						**027550**
Staphylococcus kloosii		100.0	0.58	NIT 1			
027551	**Low discrimination**						**027551**
					OXI	YEP	
Staphylococcus lentus		83.44	0.35	SUC 99	99	1	
Staphylococcus sciuri		14.55	0.18		99	57	
Staphylococcus kloosii		2.01	0.17		1	99	
027552	**Very good identification**						**027552**
					OXI	YEP	
Staphylococcus kloosii		98.63	0.59	NIT 1	1	99	
Staphylococcus lentus		1.37	0.05		99	1	
027553	**Good identification**						**027553**
					OXI	YEP	
Staphylococcus lentus		97.15	0.47	SUC 99	99	1	
Staphylococcus sciuri		0.78	0.02		99	57	
Staphylococcus kloosii		0.72	0.18		1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
027570	**Good identification**							**027570**
Staphylococcus kloosii		100.0	0.46	NIT 1				
027571	**Low discrimination**							**027571**
					OXI			
Staphylococcus lentus		63.93	0.35	SUC 99	99			
Staphylococcus gallinarum		27.66	0.20		1			
Staphylococcus sciuri		8.41	0.15		99			
027572	**Acceptable identification**							**027572**
					OXI	YEP		
Staphylococcus kloosii		94.75	0.46	NIT 1	1	99		
Staphylococcus lentus		5.25	0.05		99	1		
027573	**Low discrimination**							**027573**
					OXI	COA	NOV	
Staphylococcus lentus		69.57	0.47	SUC 99	99	1	99	
Staphylococcus gallinarum		20.79	0.28		1	1	99	
Staphylococcus lutrae		9.17	0.18		1	99	1	
027611	**Low discrimination**							**027611**
Staphylococcus sciuri		82.76	0.41	BGA 1				
Staphylococcus lentus		17.25	0.29					
027613	**Low discrimination**							**027613**
Staphylococcus lentus		81.94	0.40	GAL 99				
Staphylococcus sciuri		18.06	0.25					
027631	**Low discrimination**							**027631**
Staphylococcus sciuri		78.36	0.39	BGA 1				
Staphylococcus lentus		21.64	0.29					
027633	**Low discrimination**							**027633**
Staphylococcus lentus		85.75	0.40	GAL 99				
Staphylococcus sciuri		14.25	0.22					
027650	**Acceptable identification**							**027650**
					OXI			
Staphylococcus sciuri		90.05	0.34	BGA 1	99			
Staphylococcus kloosii		9.96	0.31	MNS 89	1			
027651	**Low discrimination**							**027651**
Staphylococcus sciuri		89.89	0.53	BGA 1				
Staphylococcus lentus		10.11	0.35					
027653	**Low discrimination**							**027653**
					OXI	COA		
Staphylococcus lentus		70.31	0.47	GAL 99	99	1		
Staphylococcus sciuri		28.71	0.37		99	1		
Staphylococcus intermedius		0.51	-.02		1	99		
027670	**Good identification**							**027670**
					OXI			
Staphylococcus sciuri		96.47	0.32	BGA 1	99			
Staphylococcus kloosii		3.53	0.19	MNS 89	1			
027671	**Low discrimination**							**027671**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		81.56	0.51	BGA 1	99			
Staphylococcus lentus		12.16	0.35		99			
Staphylococcus gallinarum		5.26	0.20		1			
027673	**Low discrimination**							**027673**
					OXI			
Staphylococcus lentus		58.25	0.47	GAL 99	99			
Staphylococcus sciuri		17.94	0.34		99			
Staphylococcus gallinarum		17.41	0.28		1			
027701	**Low discrimination**							**027701**
					OXI	NOV		
Staphylococcus lentus		47.88	0.29	MAN 99	99	99		
Rothia mucilaginosa		28.43	0.13		1	1		
Staphylococcus sciuri		13.80	0.16		99	99		
027703	**Low discrimination**							**027703**
					OXI	PYR	NOV	
Staphylococcus lentus		71.89	0.40	MAN 99	99	6	99	
Staphylococcus intermedius		25.73	0.27		1	90	1	
Staphylococcus hyicus		1.42	-.05		1	1	1	
027710	**Low discrimination**							**027710**
Staphylococcus sciuri		53.52	0.28	BGA 1				
Staphylococcus lentus		46.48	0.29	MNS 89				
027711	**Good identification**							**027711**
					OXI	COA		
Staphylococcus lentus		90.93	0.70		99	1		
Staphylococcus sciuri		8.56	0.47		99	1		
Staphylococcus intermedius		0.51	0.19		1	99		
027712	**Acceptable identification**							**027712**
Staphylococcus lentus		94.98	0.40	MNS 99				
Staphylococcus sciuri		5.02	0.12					
027713	**Very good identification**							**027713**
					OXI	COA		
Staphylococcus lentus		98.59	0.81		99	1		
Staphylococcus intermedius		0.96	0.36		1	99		
Staphylococcus sciuri		0.43	0.31		99	1		
027721	**Low discrimination**							**027721**
					OXI			
Staphylococcus lentus		76.67	0.29	MAN 99	99			
Staphylococcus sciuri		16.66	0.14		99			
Staphylococcus gallinarum		6.66	-.01		1			
027723	**Acceptable identification**							**027723**
					OXI			
Staphylococcus lentus		94.34	0.40	MAN 99	99			
Staphylococcus gallinarum		5.66	0.08		1			
027730	**Low discrimination**							**027730**
Staphylococcus lentus		53.52	0.29	MNS 99				
Staphylococcus sciuri		46.48	0.26					
027731	**Good identification**							**027731**
					OXI			
Staphylococcus lentus		92.46	0.70		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		6.56	0.45		99			
Staphylococcus gallinarum		0.82	0.20		1			
027732	**Good identification**							**027732**
Staphylococcus lentus		96.17	0.40	MNS 99				
Staphylococcus sciuri		3.84	0.10					
027733	**Very good identification**							**027733**
					OXI			
Staphylococcus lentus		98.95	0.81		99			
Staphylococcus gallinarum		0.61	0.28		1			
Staphylococcus sciuri		0.32	0.28		99			
027741	**Low discrimination**							**027741**
					OXI	NOV		
Staphylococcus lentus		41.46	0.35	MAN 99	99	99		
Rothia mucilaginosa		28.30	0.20		1	1		
Staphylococcus sciuri		22.14	0.28		99	99		
027743	**Low discrimination**							**027743**
					OXI	COA	NOV	
Staphylococcus lentus		75.84	0.47	MAN 99	99	1	99	
Staphylococcus intermedius		20.06	0.30		1	99	1	
Staphylococcus gallinarum		2.24	0.08		1	1	99	
027750	**Low discrimination**							**027750**
					OXI	YEP		
Staphylococcus sciuri		62.44	0.40	BGA 1	99	57		
Staphylococcus lentus		29.27	0.35	MNS 89	99	1		
Staphylococcus kloosii		6.90	0.37		1	99		
027751	**Low discrimination**							**027751**
					OXI			
Staphylococcus lentus		84.54	0.76		99			
Staphylococcus sciuri		14.74	0.59		99			
Staphylococcus gallinarum		0.37	0.20		1			
027752	**Low discrimination**							**027752**
					OXI	YEP		
Staphylococcus lentus		84.94	0.47	MNS 99	99	1		
Staphylococcus sciuri		8.32	0.24		99	57		
Staphylococcus kloosii		6.12	0.38		1	99		
027753	**Very good identification**							**027753**
					OXI	COA		
Staphylococcus lentus		98.20	0.88		99	1		
Staphylococcus sciuri		0.79	0.43		99	1		
Staphylococcus intermedius		0.71	0.39		1	99		
027761	**Low discrimination**							**027761**
					OXI			
Staphylococcus gallinarum		75.13	0.41	URE 92	1			
Staphylococcus lentus		17.73	0.35	MAN 91	99			
Staphylococcus sciuri		7.14	0.25		99			
027763	**Low discrimination**							**027763**
					OXI	COA	NOV	
Staphylococcus gallinarum		72.92	0.49	URE 92	1	1	99	
Staphylococcus lentus		24.93	0.47	MAN 91	99	1	99	
Staphylococcus lutrae		1.69	0.12		1	99	1	
027770	**Low discrimination**							**027770**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI		
Staphylococcus sciuri		51.90	0.38	BGA 1	99		
Staphylococcus lentus		32.25	0.35	MNS 89	99		
Staphylococcus gallinarum		13.95	0.20		1		
027771	**Low discrimination**						**027771**
					OXI		
Staphylococcus lentus		63.57	0.76		99		
Staphylococcus gallinarum		27.51	0.61		1		
Staphylococcus sciuri		8.36	0.56		99		
027772	**Low discrimination**						**027772**
					OXI	YEP	
Staphylococcus lentus		63.65	0.47	MNS 99	99	1	
Staphylococcus gallinarum		19.02	0.28		1	78	
Staphylococcus arlettae		11.49	0.35		1	99	
027773	**Low discrimination**						**027773**
					OXI		
Staphylococcus lentus		76.03	0.88		99		
Staphylococcus gallinarum		22.72	0.69		1		
Staphylococcus equorum		0.47	0.38		1		
030443	**Good identification**						**030443**
Aerococcus viridans		100.0	0.28	ORN 1			
030452	**Good identification**						**030452**
Aerococcus viridans		100.0	0.30	ORN 1			
030453	**Good identification**						**030453**
Aerococcus viridans		100.0	0.38	ORN 1			
030463	**Good identification**						**030463**
Aerococcus viridans		100.0	0.28	ORN 1			
030472	**Good identification**						**030472**
Aerococcus viridans		100.0	0.30	ORN 1			
030473	**Good identification**						**030473**
Aerococcus viridans		100.0	0.38	ORN 1			
030543	**Good identification**						**030543**
Aerococcus viridans		100.0	0.28	ORN 1			
030552	**Low discrimination**						**030552**
					NOV	YEP	
Aerococcus viridans		81.22	0.30	ORN 1	1	1	
Staphylococcus kloosii		18.78	0.18	ESL 90	99	99	
030553	**Good identification**						**030553**
Aerococcus viridans		100.0	0.38	ORN 1			
030563	**Good identification**						**030563**
Aerococcus viridans		100.0	0.28	ORN 1			
030572	**Good identification**						**030572**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.30	ORN 1			
030573	**Good identification**						**030573**
Aerococcus viridans		100.0	0.38	ORN 1			
030641	**Good identification**						**030641**
Kocuria kristinae		100.0	0.30	VPT 90			
030653	**Good identification**						**030653**
Aerococcus viridans		100.0	0.38	ORN 1			
030663	**Good identification**						**030663**
Aerococcus viridans		100.0	0.28	ORN 1			
030672	**Low discrimination**						**030672**
					NOV	YEP	
Aerococcus viridans		59.99	0.30	ORN 1	1	1	
Staphylococcus arlettae		40.01	0.15	ESL 90	99	99	
030673	**Low discrimination**						**030673**
					NOV	YEP	
Aerococcus viridans		72.52	0.38	ORN 1	1	1	
Staphylococcus arlettae		27.48	0.18	ESL 90	99	99	
030752	**Good identification**						**030752**
Aerococcus viridans		100.0	0.30	ORN 1			
030753	**Good identification**						**030753**
Aerococcus viridans		100.0	0.38	ORN 1			
030763	**Good identification**						**030763**
Aerococcus viridans		100.0	0.28	ORN 1			
030772	**Low discrimination**						**030772**
					NOV	YEP	
Staphylococcus arlettae		79.08	0.31	ORN 1	99	99	
Aerococcus viridans		20.93	0.30	GLR 92	1	1	
030773	**Low discrimination**						**030773**
					NOV	YEP	
Staphylococcus arlettae		68.22	0.34	ORN 1	99	99	
Aerococcus viridans		31.78	0.38	GLR 92	1	1	
031053	**Good identification**						**031053**
Aerococcus viridans		100.0	0.31	ORN 1			
031073	**Good identification**						**031073**
Aerococcus viridans		100.0	0.31	ORN 1			
031153	**Good identification**						**031153**
Aerococcus viridans		100.0	0.31	ORN 1			
031173	**Good identification**						**031173**
Aerococcus viridans		100.0	0.31	ORN 1			
031253	**Good identification**						**031253**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.31	ORN 1			
031273	**Good identification**						**031273**
Aerococcus viridans		100.0	0.31	ORN 1			
031353	**Good identification**						**031353**
Aerococcus viridans		100.0	0.31	ORN 1			
031373	**Good identification**						**031373**
Aerococcus viridans		100.0	0.31	ORN 1			
031402	**Good identification**						**031402**
Aerococcus viridans		100.0	0.27	ORN 1			
031403	**Good identification**						**031403**
Aerococcus viridans		100.0	0.35	ORN 1			
031412	**Good identification**						**031412**
Aerococcus viridans		100.0	0.37	ORN 1			
031413	**Good identification**						**031413**
Aerococcus viridans		100.0	0.45	ORN 1			
031422	**Good identification**						**031422**
Aerococcus viridans		100.0	0.27	ORN 1			
031423	**Good identification**						**031423**
Aerococcus viridans		100.0	0.35	ORN 1			
031432	**Good identification**						**031432**
Aerococcus viridans		100.0	0.37	ORN 1			
031433	**Good identification**						**031433**
Aerococcus viridans		100.0	0.45	ORN 1			
031441	**Good identification**						**031441**
Aerococcus viridans		100.0	0.28	ORN 1			
031442	**Good identification**						**031442**
Aerococcus viridans		100.0	0.40	ORN 1			
031443	**Good identification**						**031443**
Aerococcus viridans		100.0	0.47	ORN 1			
031450	**Low discrimination**						**031450**
					OXI	NOV	
Aerococcus viridans		66.64	0.30	ORN 1	1	1	
Micrococcus equipersicus		18.48	0.02	LAC 90	99	99	
Micrococcus bovicus		14.88	-.02		99	99	
031451	**Good identification**						**031451**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Aerococcus viridans		100.0	0.38	ORN 1		
031452	**Good identification**					**031452**
Aerococcus viridans		100.0	0.50	ORN 1		
031453	**Very good identification**					**031453**
Aerococcus viridans		100.0	0.57	ORN 1		
031461	**Good identification**					**031461**
Aerococcus viridans		100.0	0.28	ORN 1		
031462	**Good identification**					**031462**
Aerococcus viridans		100.0	0.40	ORN 1		
031463	**Good identification**					**031463**
Aerococcus viridans		100.0	0.47	ORN 1		
031470	**Good identification**					**031470**
Aerococcus viridans		100.0	0.30	ORN 1		
031471	**Good identification**					**031471**
Aerococcus viridans		100.0	0.38	ORN 1		
031472	**Good identification**					**031472**
Aerococcus viridans		100.0	0.50	ORN 1		
031473	**Very good identification**					**031473**
Aerococcus viridans		100.0	0.57	ORN 1		
031502	**Good identification**					**031502**
Aerococcus viridans		100.0	0.27	ORN 1		
031503	**Good identification**					**031503**
Aerococcus viridans		100.0	0.35	ORN 1		
031512	**Good identification**					**031512**
Aerococcus viridans		100.0	0.37	ORN 1		
031513	**Good identification**					**031513**
Aerococcus viridans		100.0	0.45	ORN 1		
031522	**Good identification**					**031522**
Aerococcus viridans		100.0	0.27	ORN 1		
031523	**Good identification**					**031523**
Aerococcus viridans		100.0	0.35	ORN 1		
031532	**Good identification**					**031532**
Aerococcus viridans		100.0	0.37	ORN 1		
031533	**Good identification**					**031533**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.45	ORN 1			
031541	**Good identification**						**031541**
Aerococcus viridans		100.0	0.28	ORN 1			
031542	**Good identification**						**031542**
Aerococcus viridans		100.0	0.40	ORN 1			
031543	**Good identification**						**031543**
Aerococcus viridans		100.0	0.47	ORN 1			
031550	**Low discrimination**						**031550**
					NOV	YEP	
Aerococcus viridans		82.41	0.30	ORN 1	1	1	
Staphylococcus kloosii		17.59	0.17	LAC 90	99	99	
031551	**Good identification**						**031551**
Aerococcus viridans		100.0	0.38	ORN 1			
031552	**Good identification**						**031552**
					NOV	YEP	
Aerococcus viridans		97.50	0.50	ORN 1	1	1	
Staphylococcus kloosii		2.50	0.18		99	99	
031553	**Very good identification**						**031553**
Aerococcus viridans		100.0	0.57	ORN 1			
031561	**Good identification**						**031561**
Aerococcus viridans		100.0	0.28	ORN 1			
031562	**Good identification**						**031562**
Aerococcus viridans		100.0	0.40	ORN 1			
031563	**Good identification**						**031563**
Aerococcus viridans		100.0	0.47	ORN 1			
031570	**Good identification**						**031570**
Aerococcus viridans		100.0	0.30	ORN 1			
031571	**Good identification**						**031571**
Aerococcus viridans		100.0	0.38	ORN 1			
031572	**Good identification**						**031572**
Aerococcus viridans		100.0	0.50	ORN 1			
031573	**Very good identification**						**031573**
Aerococcus viridans		100.0	0.57	ORN 1			
031602	**Good identification**						**031602**
Aerococcus viridans		100.0	0.27	ORN 1			
031603	**Good identification**						**031603**
Aerococcus viridans		100.0	0.35	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
031612	**Good identification**					**031612**
Aerococcus viridans		100.0	0.37	ORN 1		
031613	**Good identification**					**031613**
Aerococcus viridans		100.0	0.45	ORN 1		
031622	**Good identification**					**031622**
Aerococcus viridans		100.0	0.27	ORN 1		
031623	**Good identification**					**031623**
Aerococcus viridans		100.0	0.35	ORN 1		
031632	**Good identification**					**031632**
Aerococcus viridans		100.0	0.37	ORN 1		
031633	**Good identification**					**031633**
Aerococcus viridans		100.0	0.45	ORN 1		
031642	**Good identification**					**031642**
Aerococcus viridans		100.0	0.40	ORN 1		
031643	**Acceptable identification**					**031643**
					OXI	
Aerococcus viridans		90.99	0.47	ORN 1	1	
Kocuria kristinae		9.01	0.05		99	
031650	**Good identification**					**031650**
Aerococcus viridans		100.0	0.30	ORN 1		
031651	**Good identification**					**031651**
Aerococcus viridans		100.0	0.38	ORN 1		
031652	**Good identification**					**031652**
Aerococcus viridans		100.0	0.50	ORN 1		
031653	**Very good identification**					**031653**
Aerococcus viridans		100.0	0.57	ORN 1		
031661	**Good identification**					**031661**
Aerococcus viridans		100.0	0.28	ORN 1		
031662	**Good identification**					**031662**
Aerococcus viridans		100.0	0.40	ORN 1		
031663	**Good identification**					**031663**
Aerococcus viridans		100.0	0.47	ORN 1		
031670	**Good identification**					**031670**
Aerococcus viridans		100.0	0.30	ORN 1		
031671	**Good identification**					**031671**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.38	ORN 1			
031672	**Acceptable identification**						**031672**
					NOV	YEP	
Aerococcus viridans		93.10	0.50	ORN 1	1	1	
Staphylococcus arlettae		6.90	0.15		99	99	
031673	**Very good identification**						**031673**
					NOV	YEP	
Aerococcus viridans		95.96	0.57	ORN 1	1	1	
Staphylococcus arlettae		4.04	0.18		99	99	
031702	**Good identification**						**031702**
Aerococcus viridans		100.0	0.27	ORN 1			
031703	**Good identification**						**031703**
Aerococcus viridans		100.0	0.35	ORN 1			
031712	**Good identification**						**031712**
Aerococcus viridans		100.0	0.37	ORN 1			
031713	**Low discrimination**						**031713**
					OXI	NOV	
Aerococcus viridans		89.36	0.45	ORN 1	1	1	
Staphylococcus lentus		10.64	0.12		99	99	
031722	**Good identification**						**031722**
Aerococcus viridans		100.0	0.27	ORN 1			
031723	**Good identification**						**031723**
Aerococcus viridans		100.0	0.35	ORN 1			
031732	**Good identification**						**031732**
Aerococcus viridans		100.0	0.37	ORN 1			
031733	**Low discrimination**						**031733**
					OXI	NOV	
Aerococcus viridans		89.36	0.45	ORN 1	1	1	
Staphylococcus lentus		10.64	0.12		99	99	
031742	**Good identification**						**031742**
Aerococcus viridans		100.0	0.40	ORN 1			
031743	**Good identification**						**031743**
					OXI		
Aerococcus viridans		96.80	0.47	ORN 1	1		
Kocuria kristinae		3.20	-.05		99		
031750	**Good identification**						**031750**
Aerococcus viridans		100.0	0.30	ORN 1			
031751	**Low discrimination**						**031751**
					OXI	NOV	
Aerococcus viridans		86.70	0.38	ORN 1	1	1	
Staphylococcus lentus		13.30	0.07	LAC 90	99	99	
031752	**Good identification**						**031752**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.50	ORN 1				
031753	**Good identification**							**031753**
					OXI	NOV		
Aerococcus viridans		94.30	0.57	ORN 1	1	1		
Staphylococcus lentus		5.70	0.18		99	99		
031761	**Good identification**							**031761**
Aerococcus viridans		100.0	0.28	ORN 1				
031762	**Good identification**							**031762**
Aerococcus viridans		100.0	0.40	ORN 1				
031763	**Good identification**							**031763**
Aerococcus viridans		100.0	0.47	ORN 1				
031770	**Good identification**							**031770**
Aerococcus viridans		100.0	0.30	ORN 1				
031771	**Low discrimination**							**031771**
					OXI	NOV		
Aerococcus viridans		86.70	0.38	ORN 1	1	1		
Staphylococcus lentus		13.30	0.07	LAC 90	99	99		
031772	**Low discrimination**							**031772**
					NOV	YEP		
Aerococcus viridans		70.43	0.50	ORN 1	1	1		
Staphylococcus arlettae		29.57	0.31		99	99		
031773	**Low discrimination**							**031773**
					OXI	NOV	YEP	
Aerococcus viridans		76.98	0.57	ORN 1	1	1	1	
Staphylococcus arlettae		18.36	0.34		1	99	99	
Staphylococcus lentus		4.65	0.18		99	99	1	
032000	**Low discrimination**							**032000**
					OXI			
Kocuria varians		53.51	0.33	ORN 1	13			
Kocuria rosea		26.52	0.18		35			
Dermacoccus nishinomiyaensis		12.96	0.13		93			
032100	**Good identification**							**032100**
					OXI			
Kocuria varians		97.02	0.37	ORN 1	13			
Dermacoccus nishinomiyaensis		2.98	-.02		93			
032120	**Good identification**							**032120**
Kocuria varians		100.0	0.27	ORN 1				
032400	**Acceptable identification**							**032400**
Staphylococcus auricularis		91.14	0.30	ORN 1				
Kocuria varians		8.86	0.10	BGA 14				
032440	**Good identification**							**032440**
Staphylococcus auricularis		100.0	0.37	ORN 1				
032600	**Good identification**							**032600**
Staphylococcus auricularis		100.0	0.29	ORN 1				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
032640	**Good identification**						**032640**
Staphylococcus auricularis		96.66	0.36	ORN 1			
Staphylococcus haemolyticus		3.34	0.07	BGA 14			
032643	**Good identification**						**032643**
					OXI		
Staphylococcus lugdunensis		99.52	0.43	VPT 99	1		
Kocuria kristinae		0.48	-.06		99		
032743	**Good identification**						**032743**
					COA		
Staphylococcus lugdunensis		99.41	0.43	VPT 99	1		
Staphylococcus intermedius		0.59	-.01		99		
033711	**Good identification**						**033711**
Staphylococcus lentus		98.64	0.41	ORN 1			
Staphylococcus sciuri		1.36	0.01				
033713	**Good identification**						**033713**
Staphylococcus lentus		100.0	0.53	ORN 1			
033731	**Good identification**						**033731**
Staphylococcus lentus		100.0	0.41	ORN 1			
033733	**Good identification**						**033733**
Staphylococcus lentus		100.0	0.53	ORN 1			
033751	**Good identification**						**033751**
Staphylococcus lentus		97.50	0.48	ORN 1			
Staphylococcus sciuri		2.50	0.13				
033753	**Very good identification**						**033753**
					OXI	NOV	
Staphylococcus lentus		99.83	0.59	ORN 1	99	99	
Aerococcus viridans		0.17	0.16		1	1	
033771	**Good identification**						**033771**
Staphylococcus lentus		98.11	0.48	ORN 1			
Staphylococcus sciuri		1.90	0.11				
033773	**Very good identification**						**033773**
					OXI	NOV	
Staphylococcus lentus		99.83	0.59	ORN 1	99	99	
Aerococcus viridans		0.17	0.16		1	1	
034450	**Good identification**						**034450**
Staphylococcus kloosii		100.0	0.52	ORN 1			
034452	**Good identification**						**034452**
Staphylococcus kloosii		100.0	0.53	ORN 1			
034470	**Good identification**						**034470**
Staphylococcus kloosii		100.0	0.40	ORN 1			
034472	**Good identification**						**034472**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.40	ORN 1		
034510	**Good identification**					**034510**
Staphylococcus kloosii		100.0	0.32	ORN 1		
034512	**Good identification**					**034512**
Staphylococcus kloosii		100.0	0.32	ORN 1		
034550	**Very good identification**					**034550**
Staphylococcus kloosii		100.0	0.58	ORN 1		
034552	**Very good identification**					**034552**
Staphylococcus kloosii		100.0	0.59	ORN 1		
034570	**Good identification**					**034570**
Staphylococcus kloosii		100.0	0.46	ORN 1		
034572	**Good identification**					**034572**
Staphylococcus kloosii		100.0	0.46	ORN 1		
034650	**Good identification**					**034650**
Staphylococcus kloosii		100.0	0.31	ORN 1		
034652	**Good identification**					**034652**
Staphylococcus kloosii		100.0	0.32	ORN 1		
034750	**Good identification**					**034750**
Staphylococcus kloosii		100.0	0.37	ORN 1		
034752	**Good identification**					**034752**
Staphylococcus kloosii		100.0	0.38	ORN 1		
034772	**Acceptable identification**					**034772**
Staphylococcus arlettae		90.92	0.35	ORN 1		
Staphylococcus kloosii		9.08	0.26	GLR 92		
034773	**Good identification**					**034773**
Staphylococcus arlettae		100.0	0.37	ORN 1		
035450	**Good identification**					**035450**
Staphylococcus kloosii		100.0	0.52	ORN 1		
035452	**Good identification**					**035452**
Staphylococcus kloosii		100.0	0.53	ORN 1		
035470	**Good identification**					**035470**
Staphylococcus kloosii		100.0	0.40	ORN 1		
035472	**Good identification**					**035472**
Staphylococcus kloosii		100.0	0.40	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
035510	**Good identification**							**035510**
Staphylococcus kloosii		100.0	0.32	ORN 1				
035512	**Good identification**							**035512**
Staphylococcus kloosii		100.0	0.32	ORN 1				
035550	**Very good identification**							**035550**
Staphylococcus kloosii		100.0	0.58	ORN 1				
035552	**Very good identification**							**035552**
Staphylococcus kloosii		100.0	0.59	ORN 1				
035570	**Good identification**							**035570**
Staphylococcus kloosii		100.0	0.46	ORN 1				
035572	**Good identification**							**035572**
Staphylococcus kloosii		100.0	0.46	ORN 1				
035650	**Good identification**							**035650**
Staphylococcus kloosii		100.0	0.31	ORN 1				
035652	**Good identification**							**035652**
Staphylococcus kloosii		100.0	0.32	ORN 1				
035750	**Good identification**							**035750**
Staphylococcus kloosii		100.0	0.37	ORN 1				
035752	**Good identification**							**035752**
Staphylococcus kloosii		100.0	0.38	ORN 1				
035772	**Acceptable identification**							**035772**
Staphylococcus arlettae		90.92	0.35	ORN 1				
Staphylococcus kloosii		9.08	0.26	GLR 92				
035773	**Acceptable identification**							**035773**
					OXI	NOV	YEP	
Staphylococcus arlettae		93.42	0.37	ORN 1	1	99	99	
Staphylococcus lentus		3.94	0.05	GLR 92	99	99	1	
Aerococcus viridans		2.64	0.16		1	1	1	
036703	**Acceptable identification**							**036703**
					PYR			
Staphylococcus intermedius		94.76	0.27	URE 90	90			
Staphylococcus hyicus		5.24	-.05	ORN 1	1			
036713	**Good identification**							**036713**
					COA			
Staphylococcus intermedius		97.43	0.36	URE 90	99			
Staphylococcus felis		2.57	-.04	ORN 1	1			
036743	**Acceptable identification**							**036743**
					COA			
Staphylococcus intermedius		95.00	0.30	URE 90	99			
Staphylococcus lugdunensis		5.00	0.02	ORN 1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
036753	**Good identification**						**036753**
					OXI	COA	
Staphylococcus intermedius		98.61	0.39	URE 90	1	99	
Staphylococcus lentus		1.39	0.05	ORN 1	99	1	
037711	**Acceptable identification**						**037711**
Staphylococcus lentus		91.40	0.29	ORN 1			
Staphylococcus sciuri		8.60	0.06				
037713	**Good identification**						**037713**
Staphylococcus lentus		100.0	0.40	ORN 1			
037731	**Acceptable identification**						**037731**
Staphylococcus lentus		93.37	0.29	ORN 1			
Staphylococcus sciuri		6.63	0.04				
037733	**Good identification**						**037733**
Staphylococcus lentus		100.0	0.40	ORN 1			
037751	**Low discrimination**						**037751**
Staphylococcus lentus		85.15	0.35	ORN 1			
Staphylococcus sciuri		14.85	0.18				
037753	**Good identification**						**037753**
					OXI	COA	
Staphylococcus lentus		98.50	0.47	ORN 1	99	1	
Staphylococcus sciuri		0.79	0.02		99	1	
Staphylococcus intermedius		0.71	-.02		1	99	
037771	**Low discrimination**						**037771**
					OXI		
Staphylococcus lentus		63.93	0.35	ORN 1	99		
Staphylococcus gallinarum		27.66	0.20		1		
Staphylococcus sciuri		8.41	0.15		99		
037773	**Low discrimination**						**037773**
					OXI		
Staphylococcus lentus		76.99	0.47	ORN 1	99		
Staphylococcus gallinarum		23.01	0.28		1		
040000	**Low discrimination**						**040000**
Kocuria rosea		61.81	0.74	NIT 90			
Micrococcus sp.		27.90	0.59				
Dermacoccus nishinomiyaensis		9.91	0.59				
040001	**Low discrimination**						**040001**
					NOV	OXI	
Kocuria rosea		79.32	0.48	NIT 90	1	35	
Micrococcus sp.		13.89	0.24	MNS 5	1	89	
Staphylococcus cohnii subsp. cohnii		3.76	0.32		99	1	
040002	**Low discrimination**						**040002**
Kocuria rosea		59.65	0.33	NIT 90			
Micrococcus sp.		26.93	0.18	LAC 1			
Dermacoccus nishinomiyaensis		9.57	0.18				
040010	**Low discrimination**						**040010**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					NOV	YEP			
Kocuria rosea		72.16	0.54	NIT 90	1	1			
Macrococcus bovicus		20.09	0.48	MAN 10	99	99			
Micrococcus sp.		2.96	0.18		1	41			
040011	**Low discrimination**								**040011**
					NOV				
Kocuria rosea		49.20	0.28	NIT 90	1				
Staphylococcus cohnii subsp. cohnii		26.74	0.34	MAN 10	99				
Staphylococcus capitis subsp. capitis		21.43	0.23	MNS 5	1				
040020	**Very good identification**								**040020**
Kocuria rosea		98.40	0.67	NIT 90					
Micrococcus sp.		1.05	0.18						
Dermacoccus nishinomiyaensis		0.37	0.18						
040021	**Good identification**								**040021**
Kocuria rosea		100.0	0.40	NIT 90					
040030	**Good identification**								**040030**
					NOV	YEP			
Kocuria rosea		99.35	0.47	NIT 90	1	1			
Macrococcus bovicus		0.65	0.07	MAN 10	99	99			
040040	**Low discrimination**								**040040**
					NOV	OXI			
Micrococcus sp.		60.85	0.36	GLR 1	1	89			
Kocuria rosea		18.09	0.33	MLT 7	1	35			
Staphylococcus cohnii subsp. cohnii		13.07	0.41		99	1			
040041	**Low discrimination**								**040041**
					NOV	OXI			
Staphylococcus cohnii subsp. cohnii		85.30	0.42	TRE 99	99	1			
Micrococcus sp.		7.48	0.01		1	89			
Kocuria rosea		5.74	0.07		1	35			
040043	**Good identification**								**040043**
Aerococcus viridans		100.0	0.29	ESL 90					
040050	**Low discrimination**								**040050**
					OXI	YEP			
Macrococcus bovicus		80.23	0.55	GLR 1	99	99			
Staphylococcus cohnii subsp. cohnii		10.33	0.43		1	12			
Macrococcus equipersicus		7.81	0.36		99	99			
040051	**Acceptable identification**								**040051**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		91.90	0.44	TRE 99	1	99	1	12	
Macrococcus bovicus		6.66	0.14		99	99	1	99	
Aerococcus viridans		1.45	0.19		1	1	95	1	
040053	**Low discrimination**								**040053**
					NOV				
Aerococcus viridans		86.75	0.39	ESL 90	1				
Staphylococcus cohnii subsp. urealyticum		13.26	0.11	TRE 95	99				
040063	**Good identification**								**040063**
Aerococcus viridans		100.0	0.29	ESL 90					
040072	**Good identification**								**040072**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.31	ESL 90				
040073	**Good identification**							**040073**
Aerococcus viridans		100.0	0.39	ESL 90				
040100	**Low discrimination**							**040100**
Dermacoccus nishinomiyaensis		62.07	0.44	GLR 1				
Kocuria rosea		20.53	0.33					
Micrococcus sp.		9.27	0.18					
040110	**Low discrimination**							**040110**
					NOV	YEP		
Macrococcus bovicus		82.20	0.40	GLR 1	99	99		
Kocuria rosea		6.96	0.13		1	1		
Macrococcus equipersicus		6.91	0.20		99	99		
040143	**Good identification**							**040143**
Aerococcus viridans		100.0	0.29	ESL 90				
040150	**Low discrimination**							**040150**
					OXI	YEP		
Macrococcus bovicus		88.13	0.48	GLR 1	99	99		
Macrococcus equipersicus		8.58	0.28		99	99		
Staphylococcus cohnii subsp. cohnii		2.94	0.24		1	12		
040152	**Low discrimination**							**040152**
					OXI	NOV	YEP	
Aerococcus viridans		57.97	0.31	ESL 90	1	1	1	
Macrococcus bovicus		29.62	0.07	TRE 95	99	99	99	
Staphylococcus kloosii		12.41	0.18		1	99	99	
040153	**Good identification**							**040153**
Aerococcus viridans		100.0	0.39	ESL 90				
040163	**Good identification**							**040163**
Aerococcus viridans		100.0	0.29	ESL 90				
040172	**Low discrimination**							**040172**
					NOV	YEP		
Aerococcus viridans		71.91	0.31	ESL 90	1	1		
Staphylococcus arlettae		28.09	0.12	TRE 95	99	99		
040173	**Low discrimination**							**040173**
					NOV	YEP		
Aerococcus viridans		81.84	0.39	ESL 90	1	1		
Staphylococcus arlettae		18.16	0.14	TRE 95	99	99		
040200	**Low discrimination**							**040200**
Kocuria rosea		66.34	0.54	NIT 90				
Micrococcus sp.		26.66	0.38	SUC 10				
Dermacoccus nishinomiyaensis		6.11	0.34					
040201	**Low discrimination**							**040201**
					OXI			
Kocuria rosea		82.19	0.28	NIT 90	35			
Micrococcus sp.		12.81	0.03	SUC 10	89			
Staphylococcus capitis subsp. capitis		5.00	0.06	MNS 5	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
040210	**Low discrimination**							**040210**
					NOV	YEP		
Kocuria rosea		51.19	0.35	NIT 90	1	1		
Macrococcus bovicus		32.07	0.35	SUC 10	99	99		
Staphylococcus vitulinus		12.00	0.21	MAN 10	99	75		
040211	**Low discrimination**							**040211**
Staphylococcus capitis subsp. capitis		87.92	0.29	GLR 1				
Kocuria rosea		12.08	0.08	NIT 89				
040220	**Good identification**							**040220**
					OXI			
Kocuria rosea		98.83	0.47	NIT 90	35			
Micrococcus sp.		0.94	-.03	SUC 10	89			
Kocuria varians		0.23	0.02		13			
040230	**Low discrimination**							**040230**
					NOV			
Staphylococcus vitulinus		59.66	0.29	GLR 1	99			
Kocuria rosea		40.34	0.27	NIT 91	1			
040243	**Low discrimination**							**040243**
					OXI			
Aerococcus viridans		66.63	0.29	ESL 90	1			
Kocuria kristinae		33.37	0.01	TRE 95	99			
040250	**Low discrimination**							**040250**
					OXI	NOV		
Macrococcus bovicus		89.76	0.43	GLR 1	99	99		
Staphylococcus warneri		6.16	0.30		1	1		
Macrococcus equipersicus		3.46	0.15		99	99		
040252	**Low discrimination**							**040252**
					OXI	PYR	NOV	
Aerococcus viridans		50.68	0.31	ESL 90	1	95	1	
Staphylococcus warneri		34.22	0.20	TRE 95	1	8	1	
Macrococcus bovicus		15.11	0.02		99	1	99	
040253	**Good identification**							**040253**
Aerococcus viridans		100.0	0.39	ESL 90				
040263	**Good identification**							**040263**
Aerococcus viridans		100.0	0.29	ESL 90				
040272	**Low discrimination**							**040272**
					NOV	YEP		
Staphylococcus arlettae		87.22	0.37	TRE 99	99	99		
Aerococcus viridans		12.78	0.31		1	1		
040273	**Low discrimination**							**040273**
					NOV	YEP		
Staphylococcus arlettae		79.50	0.40	TRE 99	99	99		
Aerococcus viridans		20.51	0.39		1	1		
040310	**Low discrimination**							**040310**
					NOV	YEP		
Macrococcus bovicus		88.72	0.28	GLR 1	99	99		
Macrococcus carouselicus		4.99	-.02		99	67		
Kocuria rosea		3.34	-.06		1	1		
040343	**Good identification**							**040343**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.29	ESL 90			
040350	**Acceptable identification**						**040350**
					OXI	NOV	
Macrococcus bovicus		91.82	0.35	GLR 1	99	99	
Staphylococcus warneri		4.64	0.20		1	1	
Macrococcus equipersicus		3.54	0.08		99	99	
040352	**Low discrimination**						**040352**
					NOV	PYR	
Aerococcus viridans		62.35	0.31	ESL 90	1	95	
Staphylococcus arlettae		24.36	0.12	TRE 95	99	1	
Staphylococcus warneri		13.29	0.10		1	8	
040353	**Low discrimination**						**040353**
					NOV	YEP	
Aerococcus viridans		81.84	0.39	ESL 90	1	1	
Staphylococcus arlettae		18.16	0.14	TRE 95	99	99	
040363	**Low discrimination**						**040363**
					NOV	YEP	
Aerococcus viridans		60.04	0.29	ESL 90	1	1	
Staphylococcus arlettae		39.97	0.14	TRE 95	99	99	
040372	**Good identification**						**040372**
					NOV	YEP	
Staphylococcus arlettae		97.48	0.53	TRE 99	99	99	
Aerococcus viridans		2.52	0.31		1	1	
040373	**Very good identification**						**040373**
					NOV	YEP	
Staphylococcus arlettae		95.14	0.55	TRE 99	99	99	
Aerococcus viridans		4.33	0.39		1	1	
Staphylococcus xylosus		0.53	-.01		99	40	
040400	**Low discrimination**						**040400**
					NOV		
Staphylococcus cohnii subsp. cohnii		88.61	0.72		99		
Kocuria rosea		3.92	0.33		1		
Staphylococcus auricularis		2.36	0.36		1		
040401	**Very good identification**						**040401**
					NOV		
Staphylococcus cohnii subsp. cohnii		99.41	0.73		99		
Staphylococcus auricularis		0.27	0.17		1		
Kocuria rosea		0.21	0.07		1		
040402	**Low discrimination**						**040402**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		58.36	0.31	LAC 1	99	1	
Aerococcus viridans		24.52	0.35		1	95	
Staphylococcus auricularis		17.12	0.17		1	40	
040403	**Low discrimination**						**040403**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		48.65	0.32	LAC 1	99	1	
Aerococcus viridans		44.02	0.43		1	95	
Staphylococcus cohnii subsp. urealyticum		7.34	0.16		99	53	
040410	**Low discrimination**						**040410**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		84.29	0.74		1	12	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Macrococcus bovicus		13.47	0.51		99	99			
Macrococcus carouselicus		0.98	0.24		99	67			
040411	**Very good identification**								**040411**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		99.49	0.75		1	99	1	12	
Aerococcus viridans		0.24	0.33		1	1	95	1	
Macrococcus bovicus		0.15	0.10		99	99	1	99	
040412	**Low discrimination**								**040412**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		44.06	0.33	LAC 1	1	99	1	12	
Aerococcus viridans		43.63	0.45		1	1	95	1	
Macrococcus bovicus		7.04	0.10		99	99	1	99	
040413	**Low discrimination**								**040413**
					NOV	PYR			
Aerococcus viridans		49.24	0.53	ESL 90	1	95			
Staphylococcus cohnii subsp. urealyticum		27.67	0.36		99	53			
Staphylococcus cohnii subsp. cohnii		23.09	0.34		99	1			
040421	**Low discrimination**								**040421**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		83.91	0.32	XYL 1	99	1			
Aerococcus viridans		8.44	0.23		1	95			
Kocuria rosea		7.66	-.01		1	1			
040422	**Good identification**								**040422**
Aerococcus viridans		100.0	0.35	ESL 90					
040423	**Good identification**								**040423**
Aerococcus viridans		100.0	0.43	ESL 90					
040430	**Low discrimination**								**040430**
					NOV	YEP	OXI		
Staphylococcus cohnii subsp. cohnii		69.75	0.33	XYL 1	99	12	1		
Kocuria rosea		11.44	0.06		1	1	35		
Macrococcus bovicus		11.14	0.10		99	99	99		
040431	**Low discrimination**								**040431**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		80.84	0.34	XYL 1	99	1			
Aerococcus viridans		19.16	0.33		1	95			
040432	**Good identification**								**040432**
Aerococcus viridans		100.0	0.45	ESL 90					
040433	**Good identification**								**040433**
Aerococcus viridans		100.0	0.53	ESL 90					
040440	**Very good identification**								**040440**
					NOV				
Staphylococcus cohnii subsp. cohnii		97.51	0.82		99				
Staphylococcus auricularis		1.92	0.44		1				
Staphylococcus warneri		0.23	0.33		1				
040441	**Very good identification**								**040441**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		99.62	0.83		99	1			
Staphylococcus auricularis		0.20	0.24		1	40			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		0.13	0.35		1	95		
040442	**Low discrimination**							**040442**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		54.80	0.41	LAC 1	99	1		
Aerococcus viridans		29.08	0.48		1	95		
Staphylococcus auricularis		11.85	0.24		1	40		
040443	**Low discrimination**							**040443**
					NOV	PYR		
Aerococcus viridans		46.94	0.55	ESL 90	1	95		
Staphylococcus cohnii subsp. cohnii		41.07	0.42		99	1		
Staphylococcus cohnii subsp. urealyticum		11.09	0.31		99	53		
040450	**Low discrimination**							**040450**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		88.11	0.85		1	12		
Macrococcus bovicus		10.37	0.59		99	99		
Macrococcus equipersicus		0.67	0.36		99	99		
040451	**Very good identification**							**040451**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		99.38	0.85		99	1		
Aerococcus viridans		0.30	0.45		1	95		
Staphylococcus cohnii subsp. urealyticum		0.21	0.31		99	53		
040452	**Low discrimination**							**040452**
					NOV	YEP	PYR	
Aerococcus viridans		39.64	0.57	ESL 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		31.69	0.43		99	12	1	
Staphylococcus kloosii		22.78	0.53		99	99	52	
040453	**Low discrimination**							**040453**
					NOV	PYR		
Aerococcus viridans		46.14	0.65	ESL 90	1	95		
Staphylococcus cohnii subsp. urealyticum		36.74	0.52		99	53		
Staphylococcus cohnii subsp. cohnii		17.13	0.44		99	1		
040460	**Acceptable identification**							**040460**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		92.71	0.41	XYL 1	99	1		
Aerococcus viridans		5.47	0.28		1	95		
Staphylococcus auricularis		1.82	0.03		1	40		
040461	**Low discrimination**							**040461**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		88.73	0.42	XYL 1	99	1		
Aerococcus viridans		11.27	0.35		1	95		
040462	**Good identification**							**040462**
Aerococcus viridans		100.0	0.48	ESL 90				
040463	**Very good identification**							**040463**
Aerococcus viridans		100.0	0.55	ESL 90				
040470	**Low discrimination**							**040470**
					NOV	YEP	PYR	
Staphylococcus cohnii subsp. cohnii		70.30	0.43	XYL 1	99	12	1	
Staphylococcus kloosii		11.66	0.40		99	99	52	
Aerococcus viridans		9.77	0.38		1	1	95	
040471	**Low discrimination**							**040471**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		76.96	0.44	XYL 1	99	1			
Aerococcus viridans		23.04	0.45		1	95			
040472	**Low discrimination**								**040472**
					NOV	YEP			
Aerococcus viridans		66.54	0.57	ESL 90	1	1			
Staphylococcus arlettae		23.90	0.37		99	99			
Staphylococcus kloosii		9.56	0.41		99	99			
040473	**Low discrimination**								**040473**
					NOV	YEP			
Aerococcus viridans		82.50	0.65	ESL 90	1	1			
Staphylococcus arlettae		16.83	0.40		99	99			
Staphylococcus cohnii subsp. urealyticum		0.66	0.11		99	64			
040500	**Low discrimination**								**040500**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		87.27	0.52	GAL 10	1	99			
Macrococcus carouselicus		7.75	0.20		99	99			
Staphylococcus auricularis		2.33	0.17		1	1			
040501	**Very good identification**								**040501**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		99.09	0.53	GAL 10	99	1			
Aerococcus viridans		0.91	0.23		1	95			
040502	**Low discrimination**								**040502**
					NOV	PYR			
Aerococcus viridans		79.08	0.35	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		20.92	0.11		99	1			
040503	**Low discrimination**								**040503**
					NOV	PYR			
Aerococcus viridans		83.73	0.43	ESL 90	1	95			
Staphylococcus cohnii subsp. cohnii		10.28	0.12		99	1			
Staphylococcus cohnii subsp. urealyticum		5.98	0.08		99	53			
040510	**Low discrimination**								**040510**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		57.97	0.55	GAL 10	1	12			
Macrococcus bovicus		35.73	0.44		99	99			
Macrococcus carouselicus		2.59	0.16		99	67			
040511	**Very good identification**								**040511**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		96.91	0.55	GAL 10	1	99	1	12	
Aerococcus viridans		2.09	0.33		1	1	95	1	
Macrococcus bovicus		0.56	0.03		99	99	1	99	
040512	**Low discrimination**								**040512**
					NOV	YEP	PYR		
Aerococcus viridans		70.61	0.45	ESL 90	1	1	95		
Staphylococcus kloosii		16.58	0.33		99	99	52		
Staphylococcus cohnii subsp. cohnii		7.92	0.13		99	12	1		
040513	**Low discrimination**								**040513**
					NOV	PYR			
Aerococcus viridans		77.34	0.53	ESL 90	1	95			
Staphylococcus cohnii subsp. urealyticum		18.63	0.29		99	53			
Staphylococcus cohnii subsp. cohnii		4.03	0.14		99	1			
040522	**Good identification**								**040522**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.35	ESL 90				
040523	**Good identification**							**040523**
Aerococcus viridans		100.0	0.43	ESL 90				
040531	**Low discrimination**							**040531**
					NOV	PYR		
Aerococcus viridans		68.08	0.33	ESL 90	1	95		
Staphylococcus cohnii subsp. cohnii		31.92	0.14	LAC 90	99	1		
040532	**Low discrimination**							**040532**
					NOV	YEP		
Aerococcus viridans		87.65	0.45	ESL 90	1	1		
Staphylococcus arlettae		7.21	0.12		99	99		
Staphylococcus kloosii		5.15	0.20		99	99		
040533	**Good identification**							**040533**
					NOV	YEP		
Aerococcus viridans		95.54	0.53	ESL 90	1	1		
Staphylococcus arlettae		4.46	0.14		99	99		
040540	**Very good identification**							**040540**
					NOV			
Staphylococcus cohnii subsp. cohnii		96.06	0.63	GAL 10	99			
Staphylococcus auricularis		1.89	0.24		1			
Staphylococcus warneri		0.65	0.23		1			
040541	**Very good identification**							**040541**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		98.66	0.63	GAL 10	99	1		
Aerococcus viridans		1.14	0.35		1	95		
Staphylococcus auricularis		0.20	0.05		1	40		
040542	**Low discrimination**							**040542**
					NOV	PYR		
Aerococcus viridans		74.91	0.48	ESL 90	1	95		
Staphylococcus cohnii subsp. cohnii		15.69	0.22		99	1		
Staphylococcus warneri		3.48	0.13		1	8		
040543	**Low discrimination**							**040543**
					NOV	PYR		
Aerococcus viridans		83.44	0.55	ESL 90	1	95		
Staphylococcus cohnii subsp. urealyticum		8.45	0.23		99	53		
Staphylococcus cohnii subsp. cohnii		8.11	0.22		99	1		
040550	**Low discrimination**							**040550**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		61.84	0.65	GAL 10	1	12		
Macrococcus bovicus		28.08	0.51		99	99		
Staphylococcus kloosii		7.24	0.58		1	99		
040551	**Very good identification**							**040551**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		96.08	0.66	GAL 10	99	1		
Aerococcus viridans		2.62	0.45		1	95		
Staphylococcus cohnii subsp. urealyticum		0.79	0.23		99	53		
040552	**Low discrimination**							**040552**
					NOV	YEP	PYR	
Staphylococcus kloosii		49.00	0.59	PHS 99	99	99	52	
Aerococcus viridans		43.93	0.57		1	1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		3.90	0.24		99	12	1	
040553	**Low discrimination**							**040553**
					NOV	PYR		
Aerococcus viridans		71.48	0.65	ESL 90	1	95		
Staphylococcus cohnii subsp. urealyticum		24.39	0.44		99	53		
Staphylococcus cohnii subsp. cohnii		2.95	0.25		99	1		
040561	**Low discrimination**							**040561**
					NOV	PYR		
Aerococcus viridans		53.34	0.35	ESL 90	1	95		
Staphylococcus cohnii subsp. cohnii		46.67	0.22	LAC 90	99	1		
040562	**Acceptable identification**							**040562**
					NOV	YEP		
Aerococcus viridans		94.19	0.48	ESL 90	1	1		
Staphylococcus arlettae		5.81	0.12		99	99		
040563	**Very good identification**							**040563**
					NOV	YEP		
Aerococcus viridans		96.62	0.55	ESL 90	1	1		
Staphylococcus arlettae		3.39	0.14		99	99		
040570	**Low discrimination**							**040570**
					NOV	YEP	PYR	
Staphylococcus kloosii		49.67	0.46	PHS 99	99	99	52	
Aerococcus viridans		21.44	0.38		1	1	95	
Staphylococcus cohnii subsp. cohnii		17.14	0.24		99	12	1	
040571	**Low discrimination**							**040571**
					NOV	YEP	PYR	
Aerococcus viridans		67.74	0.45	ESL 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		25.14	0.25	LAC 90	99	12	1	
Staphylococcus arlettae		7.12	0.14		99	99	1	
040572	**Low discrimination**							**040572**
					NOV	YEP		
Staphylococcus arlettae		61.41	0.53	SUC 99	99	99		
Aerococcus viridans		30.17	0.57		1	1		
Staphylococcus kloosii		8.42	0.47		99	99		
040573	**Low discrimination**							**040573**
					NOV	YEP		
Staphylococcus arlettae		53.42	0.55	SUC 99	99	99		
Aerococcus viridans		46.20	0.65		1	1		
Staphylococcus xylosus		0.38	0.02		99	40		
040600	**Low discrimination**							**040600**
					OXI	NOV		
Macrococcus carouselicus		47.65	0.34	GLR 1	99	99		
Staphylococcus auricularis		25.10	0.36	ESL 94	1	1		
Staphylococcus cohnii subsp. cohnii		10.30	0.31		1	99		
040601	**Low discrimination**							**040601**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		46.38	0.32	SUC 1	1	99		
Kocuria kristinae		37.37	0.20		99	1		
Staphylococcus auricularis		11.60	0.16		1	1		
040602	**Low discrimination**							**040602**
					PYR			
Aerococcus viridans		44.77	0.35	ESL 90	95			
Staphylococcus auricularis		28.87	0.16		40			

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus warneri		26.36	0.23		8				
040603	**Low discrimination**								**040603**
					OXI				
Aerococcus viridans		81.79	0.43	ESL 90	1				
Kocuria kristinae		18.21	0.07		99				
040610	**Low discrimination**								**040610**
					OXI	YEP			
Macrococcus bovicus		45.26	0.39	GLR 1	99	99			
Macrococcus carouselicus		26.61	0.30		99	67			
Staphylococcus cohnii subsp. cohnii		11.44	0.33		1	12			
040611	**Low discrimination**								**040611**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		78.50	0.34	SUC 1	1	99	1	12	
Aerococcus viridans		18.60	0.33		1	1	95	1	
Macrococcus bovicus		2.89	-.02		99	99	1	99	
040612	**Low discrimination**								**040612**
					PYR	YEP			
Aerococcus viridans		78.00	0.45	ESL 90	95	1			
Staphylococcus warneri		11.09	0.20		8	43			
Staphylococcus pasteuri		7.76	0.03		1	86			
040613	**Good identification**								**040613**
					NOV				
Aerococcus viridans		98.29	0.53	ESL 90	1				
Staphylococcus cohnii subsp. urealyticum		1.71	0.05		99				
040622	**Good identification**								**040622**
Aerococcus viridans		100.0	0.35	ESL 90					
040623	**Good identification**								**040623**
Aerococcus viridans		100.0	0.43	ESL 90					
040631	**Good identification**								**040631**
Aerococcus viridans		100.0	0.33	ESL 90					
040632	**Low discrimination**								**040632**
					NOV	YEP			
Staphylococcus arlettae		58.96	0.37	MLT 99	99	99			
Aerococcus viridans		41.04	0.45		1	1			
040633	**Low discrimination**								**040633**
					NOV	YEP			
Aerococcus viridans		55.06	0.53	ESL 90	1	1			
Staphylococcus arlettae		44.94	0.40		99	99			
040640	**Low discrimination**								**040640**
					PYR				
Staphylococcus warneri		39.45	0.59	URE 98	8				
Staphylococcus petrasii subsp. pragensis		33.55	0.33		99				
Staphylococcus auricularis		15.91	0.43		40				
040641	**Low discrimination**								**040641**
					OXI	NOV			
Kocuria kristinae		61.55	0.39	VPT 90	99	1			
Staphylococcus cohnii subsp. cohnii		26.88	0.42	GLR 1	1	99			
Staphylococcus auricularis		4.95	0.23		1	1			
040642	**Low discrimination**								**040642**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR			
Staphylococcus warneri		64.22	0.50	URE 98	8			
Aerococcus viridans		22.96	0.48		95			
Staphylococcus auricularis		8.64	0.23		40			
040643	**Low discrimination**							**040643**
					OXI	PYR		
Aerococcus viridans		64.80	0.55	ESL 90	1	95		
Kocuria kristinae		32.46	0.27		99	93		
Staphylococcus warneri		1.59	0.15		1	8		
040650	**Low discrimination**							**040650**
					OXI	NOV	YEP	
Macrococcus bovicus		40.17	0.47	GLR 1	99	99	99	
Staphylococcus warneri		34.92	0.57		1	1	43	
Staphylococcus cohnii subsp. cohnii		13.79	0.43		1	99	12	
040651	**Low discrimination**							**040651**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		69.00	0.44	SUC 1	1	99	1	
Aerococcus viridans		20.66	0.45		1	1	95	
Kocuria kristinae		5.17	0.11		99	1	93	
040652	**Low discrimination**							**040652**
					NOV	PYR		
Aerococcus viridans		43.74	0.57	ESL 90	1	95		
Staphylococcus warneri		29.53	0.47		1	8		
Staphylococcus arlettae		15.71	0.37		99	1		
040653	**Low discrimination**							**040653**
					NOV	YEP		
Aerococcus viridans		80.55	0.65	ESL 90	1	1		
Staphylococcus arlettae		16.44	0.40		99	99		
Staphylococcus cohnii subsp. urealyticum		1.98	0.21		99	64		
040660	**Low discrimination**							**040660**
					PYR			
Aerococcus viridans		36.78	0.28	ESL 90	95			
Staphylococcus warneri		28.05	0.18	LAC 90	8			
Staphylococcus petrasii subsp. pragensis		23.86	-.09		99			
040661	**Low discrimination**							**040661**
					OXI			
Aerococcus viridans		84.59	0.35	ESL 90	1			
Kocuria kristinae		15.41	-.02	LAC 90	99			
040662	**Low discrimination**							**040662**
					NOV	PYR		
Staphylococcus arlettae		51.17	0.37	MAN 99	99	1		
Aerococcus viridans		47.49	0.48		1	95		
Staphylococcus warneri		1.34	0.09		1	8		
040663	**Low discrimination**							**040663**
					NOV	YEP		
Aerococcus viridans		62.03	0.55	ESL 90	1	1		
Staphylococcus arlettae		37.97	0.40		99	99		
040670	**Low discrimination**							**040670**
					OXI	NOV	YEP	
Staphylococcus arlettae		68.10	0.37	LAC 99	1	99	99	
Aerococcus viridans		21.07	0.38		1	1	1	
Macrococcus bovicus		4.46	0.05		99	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
040671	**Low discrimination**						**040671**
					NOV	YEP	
Staphylococcus arlettae		63.66	0.40	LAC 99	99	99	
Aerococcus viridans		34.67	0.45		1	1	
Staphylococcus xylosus		1.67	-.02		99	40	
040672	**Very good identification**						**040672**
					NOV	YEP	
Staphylococcus arlettae		97.23	0.78		99	99	
Aerococcus viridans		2.73	0.57		1	1	
Staphylococcus kloosii		0.04	0.20		99	99	
040673	**Very good identification**						**040673**
					NOV	YEP	
Staphylococcus arlettae		95.22	0.81		99	99	
Aerococcus viridans		4.71	0.65		1	1	
Staphylococcus xylosus		0.07	0.07		99	40	
040700	**Low discrimination**						**040700**
					OXI	NOV	
Macrococcus carouselicus		76.29	0.27	GLR 1	99	99	
Staphylococcus auricularis		10.42	0.16	ESL 94	1	1	
Staphylococcus warneri		9.01	0.23		1	1	
040702	**Low discrimination**						**040702**
					PYR		
Aerococcus viridans		84.32	0.35	ESL 90	95		
Staphylococcus warneri		15.68	0.13		8		
040703	**Low discrimination**						**040703**
					OXI	PYR	
Aerococcus viridans		74.79	0.43	ESL 90	1	95	
Staphylococcus hyicus		19.66	0.07		1	1	
Kocuria kristinae		5.55	-.03		99	93	
040710	**Low discrimination**						**040710**
					OXI	NOV	
Macrococcus bovicus		54.97	0.31	GLR 1	99	99	
Macrococcus carouselicus		32.32	0.23		99	99	
Staphylococcus warneri		4.33	0.20		1	1	
040711	**Low discrimination**						**040711**
					NOV	PYR	
Aerococcus viridans		68.08	0.33	ESL 90	1	95	
Staphylococcus cohnii subsp. cohnii		31.92	0.14	LAC 90	99	1	
040712	**Low discrimination**						**040712**
					NOV	PYR	
Aerococcus viridans		88.72	0.45	ESL 90	1	95	
Staphylococcus arlettae		7.30	0.12		99	1	
Staphylococcus warneri		3.98	0.10		1	8	
040713	**Good identification**						**040713**
					NOV	YEP	
Aerococcus viridans		95.54	0.53	ESL 90	1	1	
Staphylococcus arlettae		4.46	0.14		99	99	
040722	**Low discrimination**						**040722**
					NOV	YEP	
Aerococcus viridans		80.21	0.35	ESL 90	1	1	
Staphylococcus arlettae		19.79	0.12		99	99	
040723	**Low discrimination**						**040723**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		87.71	0.43	ESL 90	1	1		
Staphylococcus arlettae		12.29	0.14		99	99		
040731	**Low discrimination**							**040731**
					NOV	YEP		
Aerococcus viridans		63.81	0.33	ESL 90	1	1		
Staphylococcus arlettae		26.83	0.14	LAC 90	99	99		
Staphylococcus xylosus		9.37	-.05		99	40		
040732	**Low discrimination**							**040732**
					NOV	YEP		
Staphylococcus arlettae		89.06	0.53	MLT 99	99	99		
Aerococcus viridans		10.94	0.45		1	1		
040733	**Low discrimination**							**040733**
					NOV	YEP		
Staphylococcus arlettae		81.55	0.55	MLT 99	99	99		
Aerococcus viridans		17.63	0.53		1	1		
Staphylococcus xylosus		0.82	0.05		99	40		
040740	**Low discrimination**							**040740**
					NOV			
Staphylococcus warneri		75.28	0.49	URE 98	1			
Staphylococcus auricularis		10.69	0.23		1			
Staphylococcus cohnii subsp. cohnii		5.95	0.22		99			
040741	**Low discrimination**							**040741**
					OXI	NOV	PYR	
Kocuria kristinae		72.81	0.30	VPT 90	99	1	93	
Aerococcus viridans		12.11	0.35	GLR 1	1	1	95	
Staphylococcus cohnii subsp. cohnii		10.60	0.22		1	99	1	
040742	**Low discrimination**							**040742**
					NOV	PYR		
Aerococcus viridans		49.34	0.48	ESL 90	1	95		
Staphylococcus warneri		43.58	0.39		1	8		
Staphylococcus arlettae		3.04	0.12		99	1		
040743	**Low discrimination**							**040743**
					OXI	NOV	YEP	
Aerococcus viridans		83.20	0.55	ESL 90	1	1	1	
Kocuria kristinae		13.89	0.17		99	1	15	
Staphylococcus arlettae		2.92	0.14		1	99	99	
040750	**Low discrimination**							**040750**
					OXI	PYR	NOV	
Macrococcus bovicus		49.60	0.39	GLR 1	99	1	99	
Staphylococcus warneri		31.77	0.46		1	8	1	
Aerococcus viridans		5.52	0.38		1	95	1	
040751	**Low discrimination**							**040751**
					NOV	YEP	PYR	
Aerococcus viridans		57.30	0.45	ESL 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		21.27	0.25	LAC 90	99	12	1	
Staphylococcus arlettae		6.02	0.14		99	99	1	
040752	**Low discrimination**							**040752**
					NOV	PYR		
Staphylococcus arlettae		60.41	0.53	XYL 99	99	1		
Aerococcus viridans		29.68	0.57		1	95		
Staphylococcus warneri		6.33	0.36		1	8		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
040753	**Low discrimination**						**040753**
					NOV	YEP	
Staphylococcus arlettae		53.03	0.55	XYL 99	99	99	
Aerococcus viridans		45.87	0.65		1	1	
Staphylococcus xylosus		0.62	0.06		99	40	
040760	**Low discrimination**						**040760**
					NOV	YEP	
Aerococcus viridans		64.31	0.28	ESL 90	1	1	
Staphylococcus arlettae		35.69	0.12	LAC 90	99	99	
040761	**Low discrimination**						**040761**
					NOV	YEP	
Aerococcus viridans		76.03	0.35	ESL 90	1	1	
Staphylococcus arlettae		23.97	0.14	LAC 90	99	99	
040762	**Low discrimination**						**040762**
					NOV	YEP	
Staphylococcus arlettae		85.93	0.53	MAN 99	99	99	
Aerococcus viridans		14.07	0.48		1	1	
040763	**Low discrimination**						**040763**
					NOV	YEP	
Staphylococcus arlettae		77.11	0.55	MAN 99	99	99	
Aerococcus viridans		22.23	0.55		1	1	
Staphylococcus xylosus		0.66	0.03		99	40	
040770	**Acceptable identification**						**040770**
					NOV	YEP	
Staphylococcus arlettae		93.28	0.53	LAC 99	99	99	
Aerococcus viridans		5.09	0.38		1	1	
Staphylococcus kloosii		1.17	0.25		99	99	
040771	**Low discrimination**						**040771**
					NOV	YEP	
Staphylococcus arlettae		87.20	0.55	LAC 99	99	99	
Aerococcus viridans		8.38	0.45		1	1	
Staphylococcus xylosus		4.09	0.19		99	40	
040772	**Excellent identification**						**040772**
					NOV	YEP	
Staphylococcus arlettae		99.49	0.94		99	99	
Aerococcus viridans		0.49	0.57		1	1	
Staphylococcus kloosii		0.01	0.26		99	99	
040773	**Excellent identification**						**040773**
					NOV	YEP	
Staphylococcus arlettae		99.00	0.96		99	99	
Aerococcus viridans		0.87	0.65		1	1	
Staphylococcus xylosus		0.13	0.28		99	40	
041000	**Low discrimination**						**041000**
					NOV		
Kocuria rosea		46.28	0.33	ESL 1	1		
Macrococcus carouselicus		24.07	0.28	NIT 90	99		
Micrococcus sp.		20.89	0.18		1		
041002	**Good identification**						**041002**
Aerococcus viridans		100.0	0.28	TRE 95			
041003	**Good identification**						**041003**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.36	TRE 95			
041010	**Low discrimination**						**041010**
Macrococcus equipersicus		40.64	0.34	URE 86			
Macrococcus bovicus		25.28	0.28	GLR 1			
Macrococcus carouselicus		24.71	0.24				
041012	**Good identification**						**041012**
Aerococcus viridans		100.0	0.38	TRE 95			
041013	**Good identification**						**041013**
Aerococcus viridans		100.0	0.46	TRE 95			
041022	**Good identification**						**041022**
Aerococcus viridans		100.0	0.28	TRE 95			
041023	**Good identification**						**041023**
Aerococcus viridans		100.0	0.36	TRE 95			
041032	**Good identification**						**041032**
Aerococcus viridans		100.0	0.38	TRE 95			
041033	**Good identification**						**041033**
Aerococcus viridans		100.0	0.46	TRE 95			
041041	**Good identification**						**041041**
Aerococcus viridans		100.0	0.29	TRE 95			
041042	**Good identification**						**041042**
Aerococcus viridans		100.0	0.41	TRE 95			
041043	**Good identification**						**041043**
Aerococcus viridans		100.0	0.48	TRE 95			
041050	**Low discrimination**						**041050**
					OXI	NOV	
Macrococcus equipersicus		63.38	0.43	URE 86	99	99	
Macrococcus bovicus		34.03	0.36	GLR 1	99	99	
Aerococcus viridans		2.60	0.31		1	1	
041051	**Acceptable identification**						**041051**
					OXI	NOV	
Aerococcus viridans		90.44	0.39	TRE 95	1	1	
Macrococcus equipersicus		9.56	0.02	LAC 90	99	99	
041052	**Good identification**						**041052**
					OXI	NOV	
Aerococcus viridans		97.33	0.51	TRE 95	1	1	
Macrococcus equipersicus		2.67	0.02		99	99	
041053	**Very good identification**						**041053**
Aerococcus viridans		100.0	0.58	TRE 95			
041061	**Good identification**						**041061**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.29	TRE 95			
041062	**Good identification**						**041062**
Aerococcus viridans		100.0	0.41	TRE 95			
041063	**Good identification**						**041063**
Aerococcus viridans		100.0	0.48	TRE 95			
041070	**Low discrimination**						**041070**
					OXI	NOV	
Aerococcus viridans		80.21	0.31	TRE 95	1	1	
Macrococcus equipersicus		19.79	0.02	LAC 90	99	99	
041071	**Good identification**						**041071**
Aerococcus viridans		100.0	0.39	TRE 95			
041072	**Good identification**						**041072**
Aerococcus viridans		100.0	0.51	TRE 95			
041073	**Very good identification**						**041073**
Aerococcus viridans		100.0	0.58	TRE 95			
041102	**Good identification**						**041102**
Aerococcus viridans		100.0	0.28	TRE 95			
041103	**Good identification**						**041103**
Aerococcus viridans		100.0	0.36	TRE 95			
041112	**Good identification**						**041112**
Aerococcus viridans		100.0	0.38	TRE 95			
041113	**Good identification**						**041113**
Aerococcus viridans		100.0	0.46	TRE 95			
041122	**Good identification**						**041122**
Aerococcus viridans		100.0	0.28	TRE 95			
041123	**Good identification**						**041123**
Aerococcus viridans		100.0	0.36	TRE 95			
041132	**Good identification**						**041132**
Aerococcus viridans		100.0	0.38	TRE 95			
041133	**Good identification**						**041133**
Aerococcus viridans		100.0	0.46	TRE 95			
041141	**Good identification**						**041141**
Aerococcus viridans		100.0	0.29	TRE 95			
041142	**Good identification**						**041142**
Aerococcus viridans		100.0	0.41	TRE 95			
041143	**Good identification**						**041143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.48	TRE 95			
041150	**Low discrimination**						**041150**
					OXI	NOV	
Macrococcus equipersicus		60.56	0.35	URE 86	99	99	
Macrococcus bovicus		32.51	0.28	GLR 1	99	99	
Aerococcus viridans		5.79	0.31		1	1	
041151	**Good identification**						**041151**
Aerococcus viridans		100.0	0.39	TRE 95			
041152	**Good identification**						**041152**
					NOV	YEP	
Aerococcus viridans		97.68	0.51	TRE 95	1	1	
Staphylococcus kloosii		2.32	0.18		99	99	
041153	**Very good identification**						**041153**
Aerococcus viridans		100.0	0.58	TRE 95			
041161	**Good identification**						**041161**
Aerococcus viridans		100.0	0.29	TRE 95			
041162	**Good identification**						**041162**
Aerococcus viridans		100.0	0.41	TRE 95			
041163	**Good identification**						**041163**
Aerococcus viridans		100.0	0.48	TRE 95			
041170	**Good identification**						**041170**
Aerococcus viridans		100.0	0.31	TRE 95			
041171	**Good identification**						**041171**
Aerococcus viridans		100.0	0.39	TRE 95			
041172	**Good identification**						**041172**
					NOV	YEP	
Aerococcus viridans		95.84	0.51	TRE 95	1	1	
Staphylococcus arlettae		4.16	0.12		99	99	
041173	**Very good identification**						**041173**
					NOV	YEP	
Aerococcus viridans		97.59	0.58	TRE 95	1	1	
Staphylococcus arlettae		2.41	0.14		99	99	
041200	**Low discrimination**						**041200**
					NOV		
Macrococcus carouselicus		87.15	0.34	GLR 1	99		
Kocuria rosea		9.17	0.13	TRE 94	1		
Micrococcus sp.		3.69	-.03		1		
041202	**Good identification**						**041202**
Aerococcus viridans		100.0	0.28	TRE 95			
041203	**Good identification**						**041203**
Aerococcus viridans		100.0	0.36	TRE 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
041210	**Low discrimination**						**041210**
Staphylococcus vitulinus		47.93	0.29	GLR 1			
Macrococcus carouselicus		41.77	0.30	NIT 91			
Macrococcus bovicus		5.26	0.16				
041212	**Good identification**						**041212**
Aerococcus viridans		100.0	0.38	TRE 95			
041213	**Good identification**						**041213**
Aerococcus viridans		100.0	0.46	TRE 95			
041222	**Good identification**						**041222**
Aerococcus viridans		100.0	0.28	TRE 95			
041223	**Good identification**						**041223**
Aerococcus viridans		100.0	0.36	TRE 95			
041230	**Good identification**						**041230**
					OXI	NOV	
Staphylococcus vitulinus		99.28	0.38	GLR 1	99	99	
Aerococcus viridans		0.72	0.19	NIT 91	1	1	
041232	**Low discrimination**						**041232**
					OXI	NOV	
Aerococcus viridans		86.56	0.38	TRE 95	1	1	
Staphylococcus vitulinus		13.44	-.03		99	99	
041233	**Good identification**						**041233**
Aerococcus viridans		100.0	0.46	TRE 95			
041241	**Low discrimination**						**041241**
					OXI		
Aerococcus viridans		66.63	0.29	TRE 95	1		
Kocuria kristinae		33.37	0.01	LAC 90	99		
041242	**Good identification**						**041242**
Aerococcus viridans		100.0	0.41	TRE 95			
041243	**Good identification**						**041243**
Aerococcus viridans		100.0	0.48	TRE 95			
041251	**Low discrimination**						**041251**
					OXI	NOV	
Aerococcus viridans		84.79	0.39	TRE 95	1	1	
Staphylococcus sciuri		15.21	0.07	LAC 90	99	99	
041252	**Good identification**						**041252**
Aerococcus viridans		100.0	0.51	TRE 95			
041253	**Very good identification**						**041253**
Aerococcus viridans		100.0	0.58	TRE 95			
041261	**Good identification**						**041261**
Aerococcus viridans		100.0	0.29	TRE 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
041262	**Good identification**						**041262**
Aerococcus viridans		100.0	0.41	TRE 95			
041263	**Good identification**						**041263**
Aerococcus viridans		100.0	0.48	TRE 95			
041270	**Low discrimination**						**041270**
					OXI	NOV	
Aerococcus viridans		74.10	0.31	TRE 95	1	1	
Staphylococcus vitulinus		25.90	-.03	LAC 90	99	99	
041271	**Low discrimination**						**041271**
					OXI	NOV	
Aerococcus viridans		88.08	0.39	TRE 95	1	1	
Staphylococcus sciuri		11.92	0.05	LAC 90	99	99	
041272	**Low discrimination**						**041272**
					NOV	YEP	
Aerococcus viridans		56.88	0.51	TRE 95	1	1	
Staphylococcus arlettae		43.12	0.37		99	99	
041273	**Low discrimination**						**041273**
					NOV	YEP	
Aerococcus viridans		69.89	0.58	TRE 95	1	1	
Staphylococcus arlettae		30.11	0.40		99	99	
041300	**Good identification**						**041300**
Macrococcus carouselicus		100.0	0.27	GLR 1			
041302	**Good identification**						**041302**
Aerococcus viridans		100.0	0.28	TRE 95			
041303	**Good identification**						**041303**
Aerococcus viridans		100.0	0.36	TRE 95			
041312	**Good identification**						**041312**
Aerococcus viridans		100.0	0.38	TRE 95			
041313	**Good identification**						**041313**
Aerococcus viridans		100.0	0.46	TRE 95			
041322	**Good identification**						**041322**
Aerococcus viridans		100.0	0.28	TRE 95			
041323	**Good identification**						**041323**
Aerococcus viridans		100.0	0.36	TRE 95			
041332	**Low discrimination**						**041332**
					NOV	YEP	
Aerococcus viridans		85.21	0.38	TRE 95	1	1	
Staphylococcus arlettae		14.79	0.12		99	99	
041333	**Acceptable identification**						**041333**
					NOV	YEP	
Aerococcus viridans		91.02	0.46	TRE 95	1	1	
Staphylococcus arlettae		8.98	0.14		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
041341	**Good identification**							**041341**
Aerococcus viridans		100.0	0.29	TRE 95				
041342	**Good identification**							**041342**
Aerococcus viridans		100.0	0.41	TRE 95				
041343	**Good identification**							**041343**
Aerococcus viridans		100.0	0.48	TRE 95				
041351	**Low discrimination**							**041351**
					OXI	NOV		
Aerococcus viridans		74.17	0.39	TRE 95	1	1		
Staphylococcus sciuri		25.84	0.13	LAC 90	99	99		
041352	**Good identification**							**041352**
					NOV	YEP		
Aerococcus viridans		95.84	0.51	TRE 95	1	1		
Staphylococcus arlettae		4.16	0.12		99	99		
041353	**Very good identification**							**041353**
					NOV	YEP		
Aerococcus viridans		97.59	0.58	TRE 95	1	1		
Staphylococcus arlettae		2.41	0.14		99	99		
041361	**Good identification**							**041361**
Aerococcus viridans		100.0	0.29	TRE 95				
041362	**Low discrimination**							**041362**
					NOV	YEP		
Aerococcus viridans		88.48	0.41	TRE 95	1	1		
Staphylococcus arlettae		11.52	0.12		99	99		
041363	**Acceptable identification**							**041363**
					NOV	YEP		
Aerococcus viridans		93.11	0.48	TRE 95	1	1		
Staphylococcus arlettae		6.89	0.14		99	99		
041370	**Low discrimination**							**041370**
					NOV	YEP		
Aerococcus viridans		71.91	0.31	TRE 95	1	1		
Staphylococcus arlettae		28.09	0.12	LAC 90	99	99		
041371	**Low discrimination**							**041371**
					OXI	NOV	YEP	
Aerococcus viridans		67.36	0.39	TRE 95	1	1	1	
Staphylococcus sciuri		17.70	0.11	LAC 90	99	99	57	
Staphylococcus arlettae		14.95	0.14		1	99	99	
041372	**Low discrimination**							**041372**
					NOV	YEP		
Staphylococcus arlettae		81.12	0.53	TRE 99	99	99		
Aerococcus viridans		18.88	0.51		1	1		
041373	**Low discrimination**							**041373**
					NOV	YEP		
Staphylococcus arlettae		70.94	0.55	TRE 99	99	99		
Aerococcus viridans		29.06	0.58		1	1		
041400	**Good identification**							**041400**
					OXI	NOV	PYR	
Macrococcus carouselicus		95.85	0.53	GLR 1	99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		2.68	0.31		1	99	1	
Aerococcus viridans		1.13	0.35		1	1	95	
041401	**Low discrimination**							**041401**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		44.69	0.32	ESL 1	1	99	1	
Aerococcus viridans		40.43	0.43		1	1	95	
Macrococcus carouselicus		14.88	0.12		99	99	1	
041402	**Good identification**							**041402**
					OXI	NOV		
Aerococcus viridans		91.29	0.55		1	1		
Macrococcus carouselicus		8.71	0.12		99	99		
041403	**Very good identification**							**041403**
					NOV			
Aerococcus viridans		99.29	0.62		1			
Staphylococcus cohnii subsp. urealyticum		0.72	0.07		99			
041410	**Low discrimination**							**041410**
Macrococcus carouselicus		75.75	0.49	GLR 1				
Macrococcus equipersicus		7.95	0.34					
Macrococcus bovicus		7.42	0.32					
041411	**Low discrimination**							**041411**
					OXI	NOV	PYR	
Aerococcus viridans		63.74	0.53	LAC 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		29.89	0.34		1	99	1	
Macrococcus carouselicus		5.00	0.08		99	99	1	
041412	**Very good identification**							**041412**
					OXI	NOV	YEP	
Aerococcus viridans		96.73	0.65		1	1	1	
Macrococcus carouselicus		1.97	0.08		99	99	67	
Staphylococcus kloosii		1.30	0.27		1	99	99	
041413	**Very good identification**							**041413**
					NOV			
Aerococcus viridans		97.63	0.72		1			
Staphylococcus cohnii subsp. urealyticum		2.37	0.28		99			
041420	**Low discrimination**							**041420**
					OXI	NOV		
Aerococcus viridans		53.80	0.35	LAC 90	1	1		
Macrococcus carouselicus		46.20	0.12		99	99		
041421	**Good identification**							**041421**
Aerococcus viridans		100.0	0.43	LAC 90				
041422	**Very good identification**							**041422**
Aerococcus viridans		100.0	0.55					
041423	**Very good identification**							**041423**
Aerococcus viridans		100.0	0.62					
041430	**Low discrimination**							**041430**
					OXI	NOV		
Aerococcus viridans		81.68	0.45	LAC 90	1	1		
Macrococcus carouselicus		14.95	0.08		99	99		
Staphylococcus vitulinus		3.38	-.08		99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
041431	**Good identification**							**041431**
Aerococcus viridans		100.0	0.53	LAC 90				
041432	**Very good identification**							**041432**
Aerococcus viridans		100.0	0.65					
041433	**Very good identification**							**041433**
Aerococcus viridans		100.0	0.72					
041440	**Low discrimination**							**041440**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		55.91	0.41	ESL 1	1	99	1	
Aerococcus viridans		29.67	0.48		1	1	95	
Macrococcus carouselicus		6.37	0.12		99	99	1	
041441	**Low discrimination**							**041441**
					NOV	PYR		
Aerococcus viridans		53.34	0.55	LAC 90	1	95		
Staphylococcus cohnii subsp. cohnii		46.67	0.42		99	1		
041442	**Very good identification**							**041442**
Aerococcus viridans		100.0	0.67					
041443	**Very good identification**							**041443**
					NOV			
Aerococcus viridans		98.99	0.75		1			
Staphylococcus cohnii subsp. urealyticum		1.01	0.23		99			
041450	**Low discrimination**							**041450**
					OXI	NOV		
Macrococcus equipersicus		27.37	0.43	URE 86	99	99		
Macrococcus bovicus		22.04	0.39	GLR 1	99	99		
Aerococcus viridans		21.29	0.57		1	1		
041451	**Low discrimination**							**041451**
					NOV	PYR		
Aerococcus viridans		70.83	0.65	LAC 90	1	95		
Staphylococcus cohnii subsp. cohnii		26.29	0.44		99	1		
Staphylococcus cohnii subsp. urealyticum		2.17	0.23		99	53		
041452	**Good identification**							**041452**
					OXI	NOV	YEP	
Aerococcus viridans		93.77	0.77		1	1	1	
Staphylococcus kloosii		5.99	0.53		1	99	99	
Macrococcus equipersicus		0.14	0.02		99	99	99	
041453	**Very good identification**							**041453**
					NOV			
Aerococcus viridans		96.67	0.85		1			
Staphylococcus cohnii subsp. urealyticum		3.33	0.43		99			
041460	**Good identification**							**041460**
Aerococcus viridans		100.0	0.48	LAC 90				
041461	**Very good identification**							**041461**
Aerococcus viridans		100.0	0.55	LAC 90				
041462	**Very good identification**							**041462**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.67					
041463	**Very good identification**							**041463**
Aerococcus viridans		100.0	0.75					
041470	**Low discrimination**							**041470**
					OXI	NOV	YEP	
Aerococcus viridans		86.50	0.57	LAC 90	1	1	1	
Staphylococcus kloosii		11.47	0.40		1	99	99	
Macrococcus equipersicus		1.12	0.02		99	99	99	
041471	**Very good identification**							**041471**
Aerococcus viridans		100.0	0.65	LAC 90				
041472	**Good identification**							**041472**
					NOV	YEP		
Aerococcus viridans		94.71	0.77		1	1		
Staphylococcus arlettae		3.78	0.37		99	99		
Staphylococcus kloosii		1.51	0.41		99	99		
041473	**Very good identification**							**041473**
					NOV	YEP		
Aerococcus viridans		97.78	0.85		1	1		
Staphylococcus arlettae		2.22	0.40		99	99		
041500	**Good identification**							**041500**
					OXI	NOV	PYR	
Macrococcus carouselicus		96.32	0.45	GLR 1	99	99	1	
Aerococcus viridans		2.64	0.35		1	1	95	
Staphylococcus cohnii subsp. cohnii		0.70	0.11		1	99	1	
041501	**Low discrimination**							**041501**
					OXI	NOV	PYR	
Aerococcus viridans		78.09	0.43	LAC 90	1	1	95	
Macrococcus carouselicus		12.32	0.04		99	99	1	
Staphylococcus cohnii subsp. cohnii		9.59	0.12		1	99	1	
041502	**Very good identification**							**041502**
					OXI	NOV		
Aerococcus viridans		96.07	0.55		1	1		
Macrococcus carouselicus		3.93	0.04		99	99		
041503	**Very good identification**							**041503**
Aerococcus viridans		100.0	0.62					
041510	**Low discrimination**							**041510**
					OXI	NOV		
Macrococcus carouselicus		72.77	0.41	GLR 1	99	99		
Aerococcus viridans		9.37	0.45		1	1		
Macrococcus equipersicus		7.64	0.26		99	99		
041511	**Acceptable identification**							**041511**
					OXI	NOV	PYR	
Aerococcus viridans		92.11	0.53	LAC 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		4.80	0.14		1	99	1	
Macrococcus carouselicus		3.10	0.00		99	99	1	
041512	**Very good identification**							**041512**
					OXI	NOV	YEP	
Aerococcus viridans		96.64	0.65		1	1	1	
Staphylococcus kloosii		2.52	0.33		1	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus carouselicus		0.84	0.00		99	99	67	
041513	**Very good identification**							**041513**
					NOV			
Aerococcus viridans		98.97	0.72		1			
Staphylococcus cohnii subsp. urealyticum		1.03	0.20		99			
041520	**Low discrimination**							**041520**
					OXI	NOV		
Aerococcus viridans		73.10	0.35	LAC 90	1	1		
Macrococcus carouselicus		26.90	0.04		99	99		
041521	**Good identification**							**041521**
Aerococcus viridans		100.0	0.43	LAC 90				
041522	**Very good identification**							**041522**
Aerococcus viridans		100.0	0.55					
041523	**Very good identification**							**041523**
Aerococcus viridans		100.0	0.62					
041530	**Low discrimination**							**041530**
					OXI	NOV	YEP	
Aerococcus viridans		88.29	0.45	LAC 90	1	1	1	
Macrococcus carouselicus		6.92	0.00		99	99	67	
Staphylococcus kloosii		4.79	0.20		1	99	99	
041531	**Good identification**							**041531**
Aerococcus viridans		100.0	0.53	LAC 90				
041532	**Very good identification**							**041532**
					NOV	YEP		
Aerococcus viridans		98.46	0.65		1	1		
Staphylococcus arlettae		0.90	0.12		99	99		
Staphylococcus kloosii		0.64	0.20		99	99		
041533	**Very good identification**							**041533**
					NOV	YEP		
Aerococcus viridans		99.48	0.72		1	1		
Staphylococcus arlettae		0.52	0.14		99	99		
041540	**Low discrimination**							**041540**
					OXI	NOV	PYR	
Aerococcus viridans		70.44	0.48	LAC 90	1	1	95	
Staphylococcus cohnii subsp. cohnii		14.75	0.22		1	99	1	
Macrococcus carouselicus		6.48	0.04		99	99	1	
041541	**Good identification**							**041541**
					NOV	PYR		
Aerococcus viridans		91.14	0.55	LAC 90	1	95		
Staphylococcus cohnii subsp. cohnii		8.86	0.22		99	1		
041542	**Very good identification**							**041542**
					NOV	YEP		
Aerococcus viridans		99.63	0.67		1	1		
Staphylococcus kloosii		0.37	0.18		99	99		
041543	**Very good identification**							**041543**
					NOV			
Aerococcus viridans		99.56	0.75		1			
Staphylococcus cohnii subsp. urealyticum		0.44	0.15		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
041550	**Low discrimination**							**041550**
					OXI	NOV	YEP	
Staphylococcus kloosii		32.87	0.58	PHS 99	1	99	99	
Aerococcus viridans		31.92	0.57		1	1	1	
Macrococcus equipersicus		17.59	0.35		99	99	99	
041551	**Good identification**							**041551**
					NOV	PYR		
Aerococcus viridans		93.91	0.65	LAC 90	1	95		
Staphylococcus cohnii subsp. cohnii		3.87	0.25		99	1		
Staphylococcus cohnii subsp. urealyticum		1.23	0.15		99	53		
041552	**Low discrimination**							**041552**
					NOV	YEP		
Aerococcus viridans		88.79	0.77		1	1		
Staphylococcus kloosii		11.01	0.59		99	99		
Staphylococcus arlettae		0.20	0.12		99	99		
041553	**Very good identification**							**041553**
					NOV	YEP		
Aerococcus viridans		98.37	0.85		1	1		
Staphylococcus cohnii subsp. urealyticum		1.45	0.36		99	64		
Staphylococcus arlettae		0.13	0.14		99	99		
041560	**Good identification**							**041560**
Aerococcus viridans		100.0	0.48	LAC 90				
041561	**Very good identification**							**041561**
Aerococcus viridans		100.0	0.55	LAC 90				
041562	**Very good identification**							**041562**
					NOV	YEP		
Aerococcus viridans		99.32	0.67		1	1		
Staphylococcus arlettae		0.68	0.12		99	99		
041563	**Very good identification**							**041563**
					NOV	YEP		
Aerococcus viridans		99.61	0.75		1	1		
Staphylococcus arlettae		0.39	0.14		99	99		
041570	**Low discrimination**							**041570**
					NOV	YEP		
Aerococcus viridans		78.25	0.57	LAC 90	1	1		
Staphylococcus kloosii		20.14	0.46		99	99		
Staphylococcus arlettae		1.61	0.12		99	99		
041571	**Very good identification**							**041571**
					OXI	NOV	YEP	
Aerococcus viridans		98.41	0.65	LAC 90	1	1	1	
Staphylococcus arlettae		1.15	0.14		1	99	99	
Staphylococcus sciuri		0.45	0.01		99	99	57	
041572	**Low discrimination**							**041572**
					NOV	YEP		
Aerococcus viridans		79.55	0.77		1	1		
Staphylococcus arlettae		17.99	0.53		99	99		
Staphylococcus kloosii		2.47	0.47		99	99		
041573	**Low discrimination**							**041573**
					NOV	YEP		
Aerococcus viridans		88.62	0.85		1	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		11.38	0.55		99	99		
041600	**Very good identification**							**041600**
					OXI	NOV		
Macrococcus carouselicus		99.42	0.59	GLR 1	99	99		
Aerococcus viridans		0.58	0.35		1	1		
041601	**Low discrimination**							**041601**
					OXI	NOV	PYR	
Aerococcus viridans		50.77	0.43	LAC 90	1	1	95	
Macrococcus carouselicus		37.93	0.18		99	99	1	
Kocuria kristinae		11.30	0.07		99	1	93	
041602	**Low discrimination**							**041602**
					OXI	NOV		
Aerococcus viridans		83.77	0.55		1	1		
Macrococcus carouselicus		16.23	0.18		99	99		
041603	**Very good identification**							**041603**
					OXI			
Aerococcus viridans		99.39	0.62		1			
Kocuria kristinae		0.62	-.05		99			
041610	**Good identification**							**041610**
					OXI	NOV		
Macrococcus carouselicus		91.90	0.55	GLR 1	99	99		
Staphylococcus vitulinus		3.79	0.24		99	99		
Aerococcus viridans		2.50	0.45		1	1		
041611	**Low discrimination**							**041611**
					OXI	NOV		
Aerococcus viridans		67.39	0.53	LAC 90	1	1		
Staphylococcus sciuri		21.88	0.26		99	99		
Macrococcus carouselicus		10.73	0.14		99	99		
041612	**Very good identification**							**041612**
					OXI	NOV		
Aerococcus viridans		96.04	0.65		1	1		
Macrococcus carouselicus		3.96	0.14		99	99		
041613	**Very good identification**							**041613**
					OXI	NOV		
Aerococcus viridans		99.42	0.72		1	1		
Staphylococcus sciuri		0.58	0.10		99	99		
041621	**Good identification**							**041621**
Aerococcus viridans		100.0	0.43	LAC 90				
041622	**Very good identification**							**041622**
Aerococcus viridans		100.0	0.55					
041623	**Very good identification**							**041623**
Aerococcus viridans		100.0	0.62					
041630	**Low discrimination**							**041630**
					OXI	NOV		
Staphylococcus vitulinus		73.96	0.33	GLR 1	99	99		
Aerococcus viridans		18.05	0.45	NIT 91	1	1		
Macrococcus carouselicus		6.71	0.14		99	99		
041631	**Low discrimination**							**041631**
					OXI	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		79.20	0.53	LAC 90	1	1		
Staphylococcus sciuri		19.40	0.24		99	99		
Staphylococcus vitulinus		1.41	-.08		99	99		
041632	**Low discrimination**							**041632**
					OXI	NOV	YEP	
Aerococcus viridans		85.89	0.65		1	1	1	
Staphylococcus arlettae		13.71	0.37		1	99	99	
Staphylococcus vitulinus		0.40	-.08		99	99	75	
041633	**Good identification**							**041633**
					OXI	NOV	YEP	
Aerococcus viridans		91.32	0.72		1	1	1	
Staphylococcus arlettae		8.28	0.40		1	99	99	
Staphylococcus sciuri		0.41	0.08		99	99	57	
041640	**Low discrimination**							**041640**
					OXI	PYR	NOV	
Aerococcus viridans		61.47	0.48	LAC 90	1	95	1	
Macrococcus carouselicus		26.79	0.18		99	1	99	
Staphylococcus warneri		5.21	0.18		1	8	1	
041641	**Low discrimination**							**041641**
					OXI	NOV	PYR	
Aerococcus viridans		65.39	0.55	LAC 90	1	1	95	
Kocuria kristinae		32.76	0.27		99	1	93	
Staphylococcus sciuri		1.85	0.07		99	99	1	
041642	**Very good identification**							**041642**
					PYR			
Aerococcus viridans		99.69	0.67		95			
Staphylococcus warneri		0.31	0.09		8			
041643	**Very good identification**							**041643**
					OXI			
Aerococcus viridans		98.63	0.75		1			
Kocuria kristinae		1.37	0.15		99			
041650	**Low discrimination**							**041650**
					OXI	NOV		
Aerococcus viridans		60.98	0.57	LAC 90	1	1		
Macrococcus bovicus		15.78	0.27		99	99		
Macrococcus equipersicus		7.75	0.22		99	99		
041651	**Low discrimination**							**041651**
					OXI	NOV	PYR	
Aerococcus viridans		76.20	0.65	LAC 90	1	1	95	
Staphylococcus sciuri		23.27	0.38		99	99	1	
Kocuria kristinae		0.53	-.02		99	1	93	
041652	**Very good identification**							**041652**
					NOV	YEP		
Aerococcus viridans		95.44	0.77		1	1		
Staphylococcus arlettae		3.81	0.37		99	99		
Staphylococcus kloosii		0.60	0.32		99	99		
041653	**Very good identification**							**041653**
					OXI	NOV	YEP	
Aerococcus viridans		97.16	0.85		1	1	1	
Staphylococcus arlettae		2.20	0.40		1	99	99	
Staphylococcus sciuri		0.54	0.22		99	99	57	
041660	**Good identification**							**041660**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.48	LAC 90				
041661	**Very good identification**							**041661**
					OXI	NOV		
Aerococcus viridans		97.91	0.55	LAC 90	1	1		
Staphylococcus sciuri		2.09	0.05		99	99		
041662	**Low discrimination**							**041662**
					NOV	YEP		
Aerococcus viridans		89.31	0.67		1	1		
Staphylococcus arlettae		10.69	0.37		99	99		
041663	**Good identification**							**041663**
					NOV	YEP		
Aerococcus viridans		93.63	0.75		1	1		
Staphylococcus arlettae		6.37	0.40		99	99		
041670	**Low discrimination**							**041670**
					OXI	NOV	YEP	
Aerococcus viridans		69.01	0.57	LAC 90	1	1	1	
Staphylococcus arlettae		24.79	0.37		1	99	99	
Staphylococcus sciuri		4.59	0.17		99	99	57	
041671	**Low discrimination**							**041671**
					OXI	NOV	YEP	
Aerococcus viridans		69.26	0.65	LAC 90	1	1	1	
Staphylococcus sciuri		15.95	0.36		99	99	57	
Staphylococcus arlettae		14.13	0.40		1	99	99	
041672	**Low discrimination**							**041672**
					NOV	YEP		
Staphylococcus arlettae		79.76	0.78		99	99		
Aerococcus viridans		20.19	0.77		1	1		
Staphylococcus kloosii		0.03	0.20		99	99		
041673	**Low discrimination**							**041673**
					OXI	NOV	YEP	
Staphylococcus arlettae		69.03	0.81		1	99	99	
Aerococcus viridans		30.76	0.85		1	1	1	
Staphylococcus sciuri		0.13	0.20		99	99	57	
041700	**Good identification**							**041700**
					OXI	NOV		
Macrococcus carouselicus		98.67	0.51	GLR 1	99	99		
Aerococcus viridans		1.33	0.35		1	1		
041701	**Low discrimination**							**041701**
					OXI	NOV	PYR	
Aerococcus viridans		68.83	0.43	LAC 90	1	1	95	
Macrococcus carouselicus		22.04	0.10		99	99	1	
Kocuria kristinae		5.11	-.03		99	1	93	
041702	**Good identification**							**041702**
					OXI	NOV		
Aerococcus viridans		92.34	0.55		1	1		
Macrococcus carouselicus		7.67	0.10		99	99		
041703	**Very good identification**							**041703**
Aerococcus viridans		100.0	0.62					
041710	**Acceptable identification**							**041710**
					OXI	NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus carouselicus		91.56	0.47	GLR 1	99	99		
Aerococcus viridans		5.81	0.45		1	1		
Macrococcus bovicus		1.11	0.12		99	99		
041711	**Low discrimination**							**041711**
					OXI	NOV		
Aerococcus viridans		58.87	0.53	LAC 90	1	1		
Staphylococcus sciuri		37.11	0.32		99	99		
Macrococcus carouselicus		4.02	0.06		99	99		
041712	**Very good identification**							**041712**
					OXI	NOV	YEP	
Aerococcus viridans		97.39	0.65		1	1	1	
Macrococcus carouselicus		1.72	0.06		99	99	67	
Staphylococcus arlettae		0.89	0.12		1	99	99	
041713	**Very good identification**							**041713**
					OXI	NOV		
Aerococcus viridans		97.83	0.72		1	1		
Staphylococcus sciuri		1.12	0.16		99	99		
Staphylococcus lentus		0.55	0.12		99	99		
041720	**Low discrimination**							**041720**
					OXI	NOV		
Aerococcus viridans		57.24	0.35	LAC 90	1	1		
Macrococcus carouselicus		42.76	0.10		99	99		
041721	**Good identification**							**041721**
Aerococcus viridans		100.0	0.43	LAC 90				
041722	**Very good identification**							**041722**
					NOV	YEP		
Aerococcus viridans		97.33	0.55		1	1		
Staphylococcus arlettae		2.67	0.12		99	99		
041723	**Very good identification**							**041723**
					NOV	YEP		
Aerococcus viridans		98.47	0.62		1	1		
Staphylococcus arlettae		1.53	0.14		99	99		
041730	**Low discrimination**							**041730**
					OXI	NOV		
Aerococcus viridans		70.42	0.45	LAC 90	1	1		
Macrococcus carouselicus		11.21	0.06		99	99		
Staphylococcus sciuri		9.66	0.11		99	99		
041731	**Low discrimination**							**041731**
					OXI	NOV	YEP	
Aerococcus viridans		64.09	0.53	LAC 90	1	1	1	
Staphylococcus sciuri		30.48	0.30		99	99	57	
Staphylococcus arlettae		2.99	0.14		1	99	99	
041732	**Low discrimination**							**041732**
					NOV	YEP		
Aerococcus viridans		52.51	0.65		1	1		
Staphylococcus arlettae		47.49	0.53		99	99		
041733	**Low discrimination**							**041733**
					NOV	YEP	PYR	
Aerococcus viridans		64.98	0.72		1	1	95	
Staphylococcus arlettae		33.39	0.55		99	99	1	
Staphylococcus equorum		0.71	0.13		99	1	5	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
041740	**Low discrimination**							**041740**
					OXI	NOV		
Aerococcus viridans		84.26	0.48	LAC 90	1	1		
Macrococcus carouselicus		15.74	0.10		99	99		
041741	**Low discrimination**							**041741**
					OXI	NOV	PYR	
Aerococcus viridans		80.86	0.55	LAC 90	1	1	95	
Kocuria kristinae		13.50	0.17		99	1	93	
Staphylococcus sciuri		4.45	0.13		99	99	1	
041742	**Very good identification**							**041742**
					NOV	YEP		
Aerococcus viridans		99.32	0.67		1	1		
Staphylococcus arlettae		0.68	0.12		99	99		
041743	**Very good identification**							**041743**
					OXI	NOV	YEP	
Aerococcus viridans		99.15	0.75		1	1	1	
Kocuria kristinae		0.46	0.05		99	1	15	
Staphylococcus arlettae		0.39	0.14		1	99	99	
041750	**Low discrimination**							**041750**
					OXI	NOV		
Aerococcus viridans		66.73	0.57	LAC 90	1	1		
Staphylococcus sciuri		11.41	0.26		99	99		
Macrococcus bovicus		7.40	0.19		99	99		
041751	**Low discrimination**							**041751**
					OXI	NOV	YEP	
Aerococcus viridans		62.05	0.65	LAC 90	1	1	1	
Staphylococcus sciuri		36.78	0.44		99	99	57	
Staphylococcus arlettae		0.73	0.14		1	99	99	
041752	**Low discrimination**							**041752**
					NOV	YEP		
Aerococcus viridans		80.55	0.77		1	1		
Staphylococcus arlettae		18.22	0.53		99	99		
Staphylococcus kloosii		0.99	0.38		99	99		
041753	**Low discrimination**							**041753**
					OXI	NOV	YEP	
Aerococcus viridans		87.53	0.85		1	1	1	
Staphylococcus arlettae		11.25	0.55		1	99	99	
Staphylococcus sciuri		0.94	0.28		99	99	57	
041760	**Acceptable identification**							**041760**
					NOV	YEP		
Aerococcus viridans		94.19	0.48	LAC 90	1	1		
Staphylococcus arlettae		5.81	0.12		99	99		
041761	**Good identification**							**041761**
					OXI	NOV	YEP	
Aerococcus viridans		92.89	0.55	LAC 90	1	1	1	
Staphylococcus sciuri		3.85	0.11		99	99	57	
Staphylococcus arlettae		3.26	0.14		1	99	99	
041762	**Low discrimination**							**041762**
					NOV	YEP		
Aerococcus viridans		59.58	0.67		1	1		
Staphylococcus arlettae		40.42	0.53		99	99		
041763	**Low discrimination**							**041763**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		72.18	0.75		1	1		
Staphylococcus arlettae		27.82	0.55		99	99		
041770	**Low discrimination**							**041770**
					OXI	NOV	YEP	
Staphylococcus arlettae		63.81	0.53	LAC 99	1	99	99	
Aerococcus viridans		31.35	0.57		1	1	1	
Staphylococcus sciuri		4.04	0.23		99	99	57	
041771	**Low discrimination**							**041771**
					OXI	NOV	YEP	
Staphylococcus arlettae		42.50	0.55	LAC 99	1	99	99	
Aerococcus viridans		36.76	0.65		1	1	1	
Staphylococcus sciuri		16.44	0.42		99	99	57	
041772	**Very good identification**							**041772**
					NOV	YEP		
Staphylococcus arlettae		95.70	0.94		99	99		
Aerococcus viridans		4.28	0.77		1	1		
Staphylococcus kloosii		0.01	0.26		99	99		
041773	**Good identification**							**041773**
					NOV	YEP	PYR	
Staphylococcus arlettae		92.42	0.96		99	99	1	
Aerococcus viridans		7.27	0.85		1	1	95	
Staphylococcus equorum		0.20	0.34		99	1	5	
042000	**Very good identification**							**042000**
Kocuria rosea		98.56	0.94					
Dermacoccus nishinomiyaensis		0.99	0.54					
Micrococcus sp.		0.26	0.33					
042001	**Very good identification**							**042001**
Kocuria rosea		99.03	0.67	MNS 5				
Staphylococcus capitis subsp. capitis		0.32	0.19					
Kocuria varians		0.21	0.21					
042002	**Acceptable identification**							**042002**
					OXI			
Kocuria rosea		94.92	0.53	LAC 1	35			
Kocuria varians		3.32	0.32		13			
Dermacoccus nishinomiyaensis		0.95	0.13		93			
042010	**Very good identification**							**042010**
					YEP	NOV		
Kocuria rosea		99.46	0.74	MAN 10	1	1		
Macrococcus bovicus		0.34	0.28		99	99		
Dermacoccus nishinomiyaensis		0.09	0.13		1	80		
042011	**Low discrimination**							**042011**
					NOV			
Kocuria rosea		71.63	0.48	MAN 10	1			
Staphylococcus capitis subsp. capitis		28.05	0.42	MNS 5	1			
Staphylococcus cohnii subsp. cohnii		0.33	0.11		99			
042012	**Good identification**							**042012**
Kocuria rosea		100.0	0.33	MAN 10				
042020	**Very good identification**							**042020**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Kocuria rosea		99.87	0.86		35			
Kocuria varians		0.10	0.34		13			
Dermacoccus nishinomiyaensis		0.02	0.13		93			
042021	**Very good identification**							**042021**
Kocuria rosea		99.83	0.60	MNS 5				
Kocuria varians		0.17	0.12					
042022	**Good identification**							**042022**
Kocuria rosea		97.21	0.45	LAC 1				
Kocuria varians		2.79	0.22					
042030	**Very good identification**							**042030**
					NOV			
Kocuria rosea		99.81	0.67	MAN 10	1			
Staphylococcus vitulinus		0.19	0.09		99			
042031	**Good identification**							**042031**
Kocuria rosea		99.09	0.40	MAN 10				
Staphylococcus capitis subsp. capitis		0.91	0.01	MNS 5				
042040	**Low discrimination**							**042040**
Kocuria rosea		84.37	0.53	MLT 1				
Staphylococcus auricularis		10.32	0.42					
Kocuria varians		2.30	0.29					
042050	**Low discrimination**							**042050**
					NOV	YEP	OXI	
Kocuria rosea		53.36	0.33	MAN 10	1	1	35	
Macrococcus bovicus		42.36	0.36	MLT 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		3.69	0.20		99	12	1	
042060	**Good identification**							**042060**
Kocuria rosea		97.54	0.45	MLT 1				
Kocuria varians		2.18	0.20					
Staphylococcus auricularis		0.28	0.01					
042100	**Low discrimination**							**042100**
					OXI			
Kocuria rosea		72.27	0.53	GAL 1	35			
Dermacoccus nishinomiyaensis		13.66	0.39		93			
Kocuria varians		13.46	0.47		13			
042102	**Low discrimination**							**042102**
					OXI			
Kocuria varians		81.39	0.35	GLR 1	13			
Kocuria rosea		15.65	0.12		35			
Dermacoccus nishinomiyaensis		2.96	-.02		93			
042110	**Low discrimination**							**042110**
					YEP	NOV		
Kocuria rosea		84.75	0.33	GAL 1	1	1		
Macrococcus bovicus		12.36	0.21	MAN 10	99	99		
Dermacoccus nishinomiyaensis		1.46	-.02		1	80		
042120	**Low discrimination**							**042120**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		86.42	0.45	GAL 1	35		
Kocuria varians		13.19	0.37		13		
Dermacoccus nishinomiyaensis		0.39	-.02		93		
042140	**Low discrimination**						**042140**
Kocuria varians		61.50	0.33	GLR 1			
Staphylococcus auricularis		20.44	0.22				
Kocuria rosea		15.20	0.12				
042150	**Good identification**						**042150**
Macrococcus bovicus		100.0	0.28	GLR 1			
042200	**Very good identification**						**042200**
					OXI		
Kocuria rosea		98.59	0.74	SUC 10	35		
Dermacoccus nishinomiyaensis		0.57	0.29		93		
Staphylococcus auricularis		0.43	0.33		1		
042201	**Acceptable identification**						**042201**
Kocuria rosea		93.76	0.48	SUC 10			
Staphylococcus capitis subsp. capitis		5.13	0.24	MNS 5			
Staphylococcus auricularis		0.87	0.14				
042202	**Acceptable identification**						**042202**
Kocuria rosea		91.63	0.33	SUC 10			
Staphylococcus auricularis		4.43	0.14	LAC 1			
Kocuria varians		3.94	0.14				
042210	**Low discrimination**						**042210**
					NOV	YEP	
Kocuria rosea		78.46	0.54	SUC 10	1	1	
Staphylococcus vitulinus		20.66	0.41	MAN 10	99	75	
Macrococcus bovicus		0.61	0.16		99	99	
042211	**Low discrimination**						**042211**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		86.17	0.47	GLR 1	1	1	
Kocuria rosea		13.17	0.28		1	35	
Staphylococcus vitulinus		0.67	0.00		99	99	
042220	**Very good identification**						**042220**
					NOV	OXI	
Kocuria rosea		99.53	0.67	SUC 10	1	35	
Staphylococcus vitulinus		0.19	0.09		99	99	
Staphylococcus muscae		0.16	-.02		1	1	
042221	**Good identification**						**042221**
Kocuria rosea		100.0	0.40	SUC 10			
042230	**Low discrimination**						**042230**
					NOV		
Staphylococcus vitulinus		62.43	0.50	GLR 1	99		
Kocuria rosea		37.57	0.47		1		
042240	**Low discrimination**						**042240**
					PYR		
Staphylococcus auricularis		43.54	0.41	GLR 3	40		
Kocuria rosea		42.87	0.33	TRE 92	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus haemolyticus		7.72	0.26		98			
042300	**Low discrimination**							**042300**
					OXI			
Kocuria rosea		72.18	0.33	GAL 1	35			
Kocuria varians		16.50	0.29	SUC 10	13			
Dermacoccus nishinomiyaensis		7.84	0.14		93			
042340	**Low discrimination**							**042340**
					PYR			
Staphylococcus haemolyticus		47.67	0.28	VPT 86	98			
Staphylococcus auricularis		26.48	0.21	ARG 92	40			
Kocuria varians		11.77	0.15	TRE 93	10			
042400	**Low discrimination**							**042400**
					NOV			
Kocuria rosea		53.47	0.53	TRE 1	1			
Staphylococcus auricularis		32.22	0.56		1			
Staphylococcus cohnii subsp. cohnii		10.10	0.49		99			
042401	**Low discrimination**							**042401**
					NOV			
Staphylococcus cohnii subsp. cohnii		63.12	0.50	NIT 7	99			
Staphylococcus auricularis		20.65	0.36		1			
Kocuria rosea		16.23	0.26		1			
042402	**Low discrimination**							**042402**
Staphylococcus auricularis		82.05	0.36	GLR 3				
Kocuria rosea		12.38	0.12	LAC 10				
Kocuria varians		3.23	0.09					
042410	**Low discrimination**							**042410**
					NOV	YEP	OXI	
Staphylococcus cohnii subsp. cohnii		57.45	0.51	NIT 7	99	12	1	
Kocuria rosea		26.55	0.33		1	1	35	
Macrococcus bovicus		13.55	0.32		99	99	99	
042411	**Good identification**							**042411**
					NOV			
Staphylococcus cohnii subsp. cohnii		96.13	0.52	NIT 7	99			
Kocuria rosea		2.16	0.07		1			
Staphylococcus capitis subsp. capitis		1.71	0.07		1			
042420	**Good identification**							**042420**
Kocuria rosea		95.82	0.45	TRE 1				
Staphylococcus muscae		1.66	-.02					
Staphylococcus auricularis		1.36	0.15					
042440	**Low discrimination**							**042440**
					NOV			
Staphylococcus auricularis		69.31	0.63	GLR 3	1			
Staphylococcus cohnii subsp. cohnii		29.49	0.59		99			
Kocuria rosea		0.50	0.12		1			
042441	**Low discrimination**							**042441**
					NOV			
Staphylococcus cohnii subsp. cohnii		80.58	0.60	NIT 7	99			
Staphylococcus auricularis		19.42	0.44		1			
042442	**Acceptable identification**							**042442**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		93.85	0.44	GLR 3	1			
Staphylococcus cohnii subsp. cohnii		3.63	0.18	LAC 10	99			
Staphylococcus warneri		1.62	0.16		1			
042450	**Low discrimination**							**042450**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		82.25	0.61	NIT 7	1	99	12	
Macrococcus bovicus		14.30	0.39		99	99	99	
Staphylococcus auricularis		1.53	0.22		1	1	1	
042451	**Very good identification**							**042451**
					PYR			
Staphylococcus cohnii subsp. cohnii		98.84	0.62	NIT 7	1			
Staphylococcus cohnii subsp. urealyticum		0.50	0.16		53			
Staphylococcus saprophyticus subsp. bovis		0.31	0.06		99			
042453	**Low discrimination**							**042453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		78.69	0.36	URE 99	99	53		
Staphylococcus cohnii subsp. cohnii		15.65	0.21		99	1		
Aerococcus viridans		5.66	0.24		1	95		
042500	**Low discrimination**							**042500**
					NOV			
Staphylococcus auricularis		57.94	0.36	GLR 3	1			
Staphylococcus cohnii subsp. cohnii		18.17	0.29	GAL 10	99			
Kocuria varians		12.13	0.23		1			
042501	**Low discrimination**							**042501**
					NOV			
Staphylococcus cohnii subsp. cohnii		72.43	0.30	NIT 7	99			
Staphylococcus auricularis		23.69	0.17	GAL 10	1			
Kocuria varians		3.88	0.02		1			
042510	**Low discrimination**							**042510**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		52.36	0.31	NIT 7	1	12		
Macrococcus bovicus		47.64	0.24	GAL 10	99	99		
042511	**Good identification**							**042511**
Staphylococcus cohnii subsp. cohnii		100.0	0.32	NIT 7				
042540	**Low discrimination**							**042540**
					OXI	NOV	YEP	
Staphylococcus auricularis		62.65	0.44	GLR 3	1	1	1	
Staphylococcus cohnii subsp. cohnii		26.66	0.40	GAL 10	1	99	12	
Macrococcus caseolyticus		7.22	0.14		99	99	99	
042541	**Low discrimination**							**042541**
					NOV			
Staphylococcus cohnii subsp. cohnii		80.58	0.40	NIT 7	99			
Staphylococcus auricularis		19.42	0.24	GAL 10	1			
042550	**Low discrimination**							**042550**
					OXI	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		44.71	0.42	NIT 7	1	1	12	
Macrococcus bovicus		29.97	0.32	GAL 10	99	1	99	
Staphylococcus saprophyticus subsp. bovis		21.87	0.31		1	99	80	
042551	**Low discrimination**							**042551**
					PYR			
Staphylococcus cohnii subsp. cohnii		88.32	0.42	NIT 7	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		9.97	0.18	GAL 10	99		
Staphylococcus cohnii subsp. urealyticum		1.71	0.08		53		
042553	**Low discrimination**						**042553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		71.09	0.29	URE 99	99		
Aerococcus viridans		11.93	0.24		1		
Staphylococcus saprophyticus subsp. bovis		10.24	0.06		99		
042600	**Low discrimination**						**042600**
Staphylococcus auricularis		79.33	0.55	GLR 3			
Kocuria rosea		15.85	0.33				
Staphylococcus warneri		1.15	0.26				
042601	**Low discrimination**						**042601**
Staphylococcus auricularis		81.03	0.36	GLR 3			
Kocuria rosea		7.67	0.07	MNS 10			
Staphylococcus hyicus		6.41	0.04				
042602	**Low discrimination**						**042602**
					PYR		
Staphylococcus auricularis		89.93	0.36	GLR 3	40		
Staphylococcus warneri		3.91	0.16	LAC 10	8		
Staphylococcus haemolyticus		2.20	0.03		98		
042610	**Low discrimination**						**042610**
					OXI	NOV	
Staphylococcus vitulinus		72.90	0.36	GLR 1	99	99	
Staphylococcus pasteuri		7.67	0.19		1	1	
Macrococcus bovicus		5.71	0.19		99	99	
042620	**Acceptable identification**						**042620**
Staphylococcus muscae		92.67	0.39	GLR 10			
Kocuria rosea		6.00	0.25	PHS 99			
Staphylococcus auricularis		0.71	0.14				
042630	**Good identification**						**042630**
					NOV	OXI	
Staphylococcus vitulinus		97.47	0.45	GLR 1	99	99	
Staphylococcus muscae		1.48	-.02		1	1	
Kocuria rosea		1.05	0.06		1	35	
042640	**Low discrimination**						**042640**
					PYR		
Staphylococcus auricularis		72.89	0.63	GLR 3	40		
Staphylococcus haemolyticus		14.94	0.50		98		
Staphylococcus warneri		8.60	0.52		8		
042641	**Low discrimination**						**042641**
					OXI		
Staphylococcus auricularis		80.79	0.43	GLR 3	1		
Kocuria kristinae		8.40	0.16	MNS 10	99		
Staphylococcus haemolyticus		4.61	0.19		1		
042642	**Low discrimination**						**042642**
					PYR		
Staphylococcus haemolyticus		41.77	0.44	VPT 86	98		
Staphylococcus auricularis		40.26	0.43	ARG 92	40		
Staphylococcus warneri		14.26	0.42		8		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
042650	**Low discrimination**							**042650**
					PYR			
Staphylococcus haemolyticus		32.08	0.45	VPT 86	98			
Staphylococcus pasteuri		24.05	0.35	ARG 92	1			
Staphylococcus warneri		22.79	0.49		8			
042651	**Low discrimination**							**042651**
					OXI	PYR		
Staphylococcus sciuri		62.34	0.32	GLR 14	99	1		
Staphylococcus saprophyticus subsp. bovis		14.31	0.18	ESL 99	1	99		
Staphylococcus cohnii subsp. cohnii		11.52	0.21		1	1		
042652	**Low discrimination**							**042652**
					PYR			
Staphylococcus haemolyticus		46.64	0.40	VPT 86	98			
Staphylococcus pasteuri		25.39	0.27	ARG 92	1			
Staphylococcus warneri		19.63	0.39		8			
042671	**Low discrimination**							**042671**
					OXI			
Staphylococcus sciuri		72.46	0.30	GLR 14	99			
Staphylococcus xylosus		27.55	0.17	ESL 99	1			
042672	**Good identification**							**042672**
					NOV	YEP		
Staphylococcus arlettae		97.27	0.37	NIT 1	99	99		
Aerococcus viridans		2.74	0.16		1	1		
042673	**Low discrimination**							**042673**
					OXI			
Staphylococcus arlettae		57.30	0.40	NIT 1	1			
Staphylococcus xylosus		34.65	0.26		1			
Staphylococcus sciuri		5.21	0.13		99			
042700	**Low discrimination**							**042700**
					OXI	NOV		
Staphylococcus auricularis		82.63	0.36	GLR 3	1	1		
Macrococcus carouselicus		4.29	0.02	GAL 10	99	99		
Staphylococcus haemolyticus		4.05	0.10		1	2		
042703	**Good identification**							**042703**
Staphylococcus hyicus		100.0	0.48	ARG 96				
042740	**Low discrimination**							**042740**
					PYR			
Staphylococcus haemolyticus		58.06	0.51	VPT 86	98			
Staphylococcus auricularis		27.91	0.43	ARG 92	40			
Staphylococcus warneri		9.37	0.42		8			
042742	**Low discrimination**							**042742**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		75.04	0.46	VPT 86	1	98	2	
Macrococcus caseolyticus		8.01	0.14	ARG 92	99	80	99	
Staphylococcus warneri		7.17	0.32		1	8	1	
042750	**Low discrimination**							**042750**
					NOV	PYR		
Staphylococcus haemolyticus		43.69	0.46	VPT 86	2	98		
Staphylococcus saprophyticus subsp. bovis		41.06	0.43	ARG 92	99	99		
Staphylococcus warneri		8.69	0.39		1	8		
042751	**Low discrimination**							**042751**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	PYR	
Staphylococcus sciuri		59.19	0.38	GLR 14	99	1	
Staphylococcus saprophyticus subsp. bovis		27.99	0.31	ESL 99	1	99	
Staphylococcus xylosus		7.69	0.15		1	65	
042752	**Low discrimination**						**042752**
					NOV	PYR	
Staphylococcus haemolyticus		62.78	0.41	VPT 86	2	98	
Staphylococcus saprophyticus subsp. bovis		26.22	0.31	ARG 92	99	99	
Staphylococcus warneri		7.40	0.29		1	8	
042771	**Low discrimination**						**042771**
					OXI		
Staphylococcus xylosus		63.55	0.37	URE 99	1		
Staphylococcus sciuri		32.09	0.35		99		
Staphylococcus arlettae		1.69	0.14		1		
042772	**Good identification**						**042772**
					OXI		
Staphylococcus arlettae		97.61	0.53	NIT 1	1		
Staphylococcus xylosus		1.41	0.05		1		
Staphylococcus sciuri		0.50	0.01		99		
042773	**Low discrimination**						**042773**
					OXI		
Staphylococcus xylosus		49.72	0.47	URE 99	1		
Staphylococcus arlettae		46.08	0.55		1		
Staphylococcus sciuri		1.44	0.19		99		
043000	**Good identification**						**043000**
					NOV		
Kocuria rosea		98.27	0.53	ESL 1	1		
Dermacoccus nishinomiyaensis		0.99	0.13		80		
Macrococcus carouselicus		0.36	0.03		99		
043010	**Low discrimination**						**043010**
					NOV	YEP	
Kocuria rosea		87.86	0.33	ESL 1	1	1	
Staphylococcus vitulinus		6.95	0.09	MAN 10	99	75	
Macrococcus bovicus		3.32	0.08		99	99	
043020	**Good identification**						**043020**
Kocuria rosea		100.0	0.45	ESL 1			
043200	**Low discrimination**						**043200**
					NOV		
Kocuria rosea		87.22	0.33	ESL 1	1		
Staphylococcus vitulinus		6.90	0.09	SUC 10	99		
Macrococcus carouselicus		5.88	0.10		99		
043210	**Good identification**						**043210**
					NOV		
Staphylococcus vitulinus		97.57	0.50	GLR 1	99		
Kocuria rosea		1.38	0.13		1		
Macrococcus carouselicus		0.54	0.06		99		
043211	**Low discrimination**						**043211**
					OXI	NOV	
Staphylococcus sciuri		73.77	0.30	GLR 14	99	99	
Staphylococcus vitulinus		17.68	0.09	TRE 97	99	99	
Staphylococcus capitis subsp. capitis		8.55	0.06		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
043230	**Very good identification**					**043230**
					NOV	
Staphylococcus vitulinus		99.63	0.59	GLR 1	99	
Kocuria rosea		0.22	0.06		1	
Staphylococcus sciuri		0.15	0.09		99	
043231	**Low discrimination**					**043231**
Staphylococcus sciuri		53.80	0.28	GLR 14		
Staphylococcus vitulinus		46.20	0.18	TRE 97		
043251	**Good identification**					**043251**
Staphylococcus sciuri		100.0	0.42	GLR 14		
043271	**Good identification**					**043271**
Staphylococcus sciuri		100.0	0.40	GLR 14		
043311	**Good identification**					**043311**
Staphylococcus sciuri		100.0	0.36	GLR 14		
043331	**Good identification**					**043331**
Staphylococcus sciuri		100.0	0.34	GLR 14		

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
043350	**Good identification**							**043350**
Staphylococcus sciuri		100.0	0.29	GLR 14				
043351	**Good identification**							**043351**
Staphylococcus sciuri		99.21	0.48	GLR 14				
Staphylococcus lentus		0.79	0.07	TRE 97				
043353	**Low discrimination**							**043353**
					OXI	NOV		
Staphylococcus sciuri		82.86	0.32	GLR 14	99	99		
Staphylococcus lentus		14.43	0.18	TRE 97	99	99		
Aerococcus viridans		2.71	0.17	LAC 14	1	1		
043370	**Good identification**							**043370**
Staphylococcus sciuri		100.0	0.27	GLR 14				
043371	**Good identification**							**043371**
					OXI			
Staphylococcus sciuri		96.73	0.45	GLR 14	99			
Staphylococcus equorum		1.60	0.06	TRE 97	1			
Staphylococcus lentus		1.03	0.07		99			
043373	**Low discrimination**							**043373**
					OXI			
Staphylococcus sciuri		55.41	0.29	GLR 14	99			
Staphylococcus equorum		14.44	0.14	TRE 97	1			
Staphylococcus lentus		12.79	0.18	LAC 14	99			
043400	**Low discrimination**							**043400**
					NOV	OXI		
Macrococcus carouselicus		76.15	0.28	GLR 1	99	99		
Kocuria rosea		13.31	0.12	NIT 6	1	35		
Staphylococcus auricularis		8.02	0.15		1	1		
043413	**Low discrimination**							**043413**
					OXI	NOV		
Aerococcus viridans		58.41	0.31	NIT 1	1	1		
Staphylococcus cohnii subsp. urealyticum		24.78	0.12		1	99		
Staphylococcus sciuri		16.81	0.04		99	99		
043433	**Low discrimination**							**043433**
					OXI	NOV		
Aerococcus viridans		82.16	0.31	NIT 1	1	1		
Staphylococcus sciuri		17.84	0.01		99	99		
043443	**Low discrimination**							**043443**
					NOV			
Aerococcus viridans		84.87	0.34	NIT 1	1			
Staphylococcus cohnii subsp. urealyticum		15.13	0.07		99			
043451	**Low discrimination**							**043451**
					OXI	NOV	PYR	
Staphylococcus sciuri		77.67	0.32	GLR 14	99	99	1	
Staphylococcus cohnii subsp. cohnii		14.36	0.21	SUC 99	1	99	1	
Aerococcus viridans		5.19	0.24		1	1	95	
043452	**Good identification**							**043452**
Aerococcus viridans		100.0	0.36	NIT 1				
043453	**Low discrimination**							**043453**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Aerococcus viridans		53.43	0.43	NIT 1	1	1		
Staphylococcus cohnii subsp. urealyticum		32.11	0.28		1	99		
Staphylococcus sciuri		14.46	0.16		99	99		
043463	**Good identification**							**043463**
Aerococcus viridans		100.0	0.34	NIT 1				
043471	**Acceptable identification**							**043471**
					OXI	NOV		
Staphylococcus sciuri		91.86	0.30	GLR 14	99	99		
Aerococcus viridans		8.14	0.24	SUC 99	1	1		
043472	**Good identification**							**043472**
Aerococcus viridans		100.0	0.36	NIT 1				
043473	**Low discrimination**							**043473**
					OXI	NOV		
Aerococcus viridans		83.04	0.43	NIT 1	1	1		
Staphylococcus sciuri		16.96	0.13		99	99		
043513	**Low discrimination**							**043513**
					OXI	NOV		
Aerococcus viridans		43.62	0.31	NIT 1	1	1		
Staphylococcus sciuri		24.37	0.10		99	99		
Staphylococcus lentus		24.08	0.12		99	99		
043533	**Low discrimination**							**043533**
					OXI	NOV		
Aerococcus viridans		50.68	0.31	NIT 1	1	1		
Staphylococcus lentus		27.97	0.12		99	99		
Staphylococcus sciuri		21.36	0.07		99	99		
043543	**Good identification**							**043543**
Aerococcus viridans		100.0	0.34	NIT 1				
043551	**Acceptable identification**							**043551**
					OXI	NOV		
Staphylococcus sciuri		94.44	0.38	GLR 14	99	99		
Aerococcus viridans		3.25	0.24	SUC 99	1	1		
Staphylococcus lentus		2.31	0.07		99	99		
043552	**Low discrimination**							**043552**
					OXI	NOV	YEP	
Aerococcus viridans		78.40	0.36	NIT 1	1	1	1	
Staphylococcus sciuri		11.88	0.03		99	99	57	
Staphylococcus kloosii		9.72	0.18		1	99	99	
043553	**Low discrimination**							**043553**
					OXI	NOV		
Aerococcus viridans		48.47	0.43	NIT 1	1	1		
Staphylococcus sciuri		25.47	0.22		99	99		
Staphylococcus lentus		13.58	0.18		99	99		
043563	**Good identification**							**043563**
Aerococcus viridans		100.0	0.34	NIT 1				
043571	**Low discrimination**							**043571**
					OXI	NOV	PYR	
Staphylococcus sciuri		86.61	0.35	GLR 14	99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus equorum		4.37	0.06	SUC 99	1	99	5	
Aerococcus viridans		3.95	0.24		1	1	95	
043572	**Low discrimination**							**043572**
					OXI	NOV	YEP	
Aerococcus viridans		74.60	0.36	NIT 1	1	1	1	
Staphylococcus arlettae		16.87	0.12		1	99	99	
Staphylococcus sciuri		8.53	0.01		99	99	57	
043573	**Low discrimination**							**043573**
					OXI	NOV	PYR	
Aerococcus viridans		42.41	0.43	NIT 1	1	1	95	
Staphylococcus sciuri		16.81	0.19		99	99	1	
Staphylococcus equorum		13.41	0.14		1	99	5	
043600	**Low discrimination**							**043600**
					OXI	NOV		
Macrococcus carouselicus		88.60	0.34	GLR 1	99	99		
Staphylococcus auricularis		4.24	0.14	NIT 6	1	1		
Staphylococcus vitulinus		3.73	0.04		99	99		
043601	**Good identification**							**043601**
Staphylococcus sciuri		100.0	0.30	GLR 14				
043610	**Low discrimination**							**043610**
Staphylococcus vitulinus		68.69	0.45	GLR 1				
Staphylococcus sciuri		20.57	0.43					
Macrococcus carouselicus		10.53	0.30					
043611	**Very good identification**							**043611**
Staphylococcus sciuri		99.59	0.61	GLR 14				
Staphylococcus vitulinus		0.42	0.04					
043613	**Acceptable identification**							**043613**
					OXI	NOV		
Staphylococcus sciuri		94.83	0.45	GLR 14	99	99		
Aerococcus viridans		3.33	0.31	LAC 14	1	1		
Staphylococcus lentus		1.84	0.12		99	99		
043621	**Good identification**							**043621**
Staphylococcus sciuri		100.0	0.28	GLR 14				
043630	**Good identification**							**043630**
Staphylococcus vitulinus		92.29	0.54	GLR 1				
Staphylococcus sciuri		7.71	0.40					
043631	**Very good identification**							**043631**
					OXI			
Staphylococcus sciuri		98.18	0.59	GLR 14	99			
Staphylococcus vitulinus		1.47	0.13		99			
Staphylococcus equorum		0.35	0.06		1			
043633	**Low discrimination**							**043633**
					OXI	NOV	PYR	
Staphylococcus sciuri		88.56	0.43	GLR 14	99	99	1	
Staphylococcus equorum		5.05	0.14	LAC 14	1	99	5	
Aerococcus viridans		4.12	0.31		1	1	95	
043641	**Good identification**							**043641**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	PYR	
Staphylococcus sciuri		95.39	0.42	GLR 14	99	99	1	
Kocuria kristinae		2.59	0.04	MAN 97	99	1	93	
Staphylococcus auricularis		1.01	0.02		1	1	40	
043650	**Very good identification**							**043650**
Staphylococcus sciuri		97.96	0.54	GLR 14				
Staphylococcus vitulinus		0.88	0.04	MNS 89				
Macrococcus bovicus		0.65	0.07					
043651	**Very good identification**							**043651**
					OXI	NOV		
Staphylococcus sciuri		99.89	0.73	GLR 14	99	99		
Aerococcus viridans		0.07	0.24		1	1		
Staphylococcus lentus		0.05	0.07		99	99		
043652	**Low discrimination**							**043652**
					OXI	NOV		
Staphylococcus sciuri		88.54	0.38	GLR 14	99	99		
Aerococcus viridans		11.46	0.36	MNS 89	1	1		
043653	**Very good identification**							**043653**
					OXI	NOV		
Staphylococcus sciuri		95.44	0.57	GLR 14	99	99		
Aerococcus viridans		3.56	0.43	LAC 14	1	1		
Staphylococcus lentus		1.00	0.18		99	99		
043661	**Good identification**							**043661**
Staphylococcus sciuri		100.0	0.40	GLR 14				
043670	**Good identification**							**043670**
					OXI	NOV		
Staphylococcus sciuri		96.60	0.52	GLR 14	99	99		
Staphylococcus vitulinus		3.10	0.13	MNS 89	99	99		
Aerococcus viridans		0.30	0.16		1	1		
043671	**Very good identification**							**043671**
					OXI	NOV	PYR	
Staphylococcus sciuri		98.80	0.71	GLR 14	99	99	1	
Staphylococcus equorum		0.96	0.26		1	99	5	
Aerococcus viridans		0.09	0.24		1	1	95	
043672	**Low discrimination**							**043672**
					OXI	NOV	YEP	
Staphylococcus sciuri		54.08	0.36	GLR 14	99	99	57	
Staphylococcus arlettae		36.65	0.37	MNS 89	1	99	99	
Aerococcus viridans		9.28	0.36	LAC 14	1	1	1	
043673	**Low discrimination**							**043673**
					OXI	YEP		
Staphylococcus sciuri		74.23	0.54	GLR 14	99	57		
Staphylococcus equorum		11.37	0.35	LAC 14	1	1		
Staphylococcus arlettae		8.24	0.40		1	99		
043700	**Low discrimination**							**043700**
					OXI	NOV		
Macrococcus carouselicus		63.04	0.27	GLR 1	99	99		
Rothia mucilaginosa		25.92	0.15	NIT 6	1	1		
Staphylococcus sciuri		11.04	0.17		99	99		
043701	**Low discrimination**							**043701**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Staphylococcus sciuri		53.45	0.36	GLR 14	99	99	
Rothia mucilaginosa		46.55	0.25	MAN 97	1	1	
043710	**Low discrimination**						**043710**
Staphylococcus sciuri		88.47	0.49	GLR 14			
Macrococcus carouselicus		9.99	0.23	MNS 89			
Staphylococcus vitulinus		1.54	0.04				
043711	**Very good identification**						**043711**
Staphylococcus sciuri		95.66	0.67	GLR 14			
Staphylococcus lentus		4.34	0.41				
043712	**Low discrimination**						**043712**
					OXI	NOV	
Staphylococcus sciuri		87.46	0.32	GLR 14	99	99	
Staphylococcus lentus		7.06	0.12	MNS 89	99	99	
Aerococcus viridans		5.48	0.23	LAC 14	1	1	
043713	**Low discrimination**						**043713**
					OXI	NOV	
Staphylococcus sciuri		49.85	0.51	GLR 14	99	99	
Staphylococcus lentus		49.25	0.53	LAC 14	99	99	
Aerococcus viridans		0.90	0.31		1	1	
043721	**Good identification**						**043721**
Staphylococcus sciuri		100.0	0.34	GLR 14			
043730	**Acceptable identification**						**043730**
Staphylococcus sciuri		94.14	0.46	GLR 14			
Staphylococcus vitulinus		5.86	0.13	MNS 89			
043731	**Good identification**						**043731**
					OXI		
Staphylococcus sciuri		92.56	0.65	GLR 14	99		
Staphylococcus lentus		5.57	0.41		99		
Staphylococcus equorum		1.74	0.26		1		
043732	**Low discrimination**						**043732**
					OXI	NOV	
Staphylococcus sciuri		79.03	0.30	GLR 14	99	99	
Staphylococcus lentus		8.46	0.12	MNS 89	99	99	
Aerococcus viridans		6.57	0.23	LAC 14	1	1	
043733	**Low discrimination**						**043733**
					OXI		
Staphylococcus lentus		49.00	0.53	GLR 1	99		
Staphylococcus sciuri		37.41	0.48		99		
Staphylococcus equorum		11.10	0.35		1		
043741	**Low discrimination**						**043741**
					OXI	NOV	
Staphylococcus sciuri		64.39	0.48	GLR 14	99	99	
Rothia mucilaginosa		34.79	0.33	MAN 97	1	1	
Staphylococcus lentus		0.52	0.07		99	99	
043743	**Low discrimination**						**043743**
					OXI	NOV	
Staphylococcus sciuri		70.68	0.32	GLR 14	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		14.65	0.34	MAN 97	1	1		
Staphylococcus lentus		12.31	0.18	LAC 14	99	99		
043750	**Very good identification**							**043750**
					OXI	NOV		
Staphylococcus sciuri		99.22	0.60	GLR 14	99	99		
Staphylococcus lentus		0.20	0.07	MNS 89	99	99		
Staphylococcus haemolyticus		0.17	0.05		1	2		
043751	**Very good identification**							**043751**
					OXI			
Staphylococcus sciuri		97.49	0.79	GLR 14	99			
Staphylococcus lentus		2.39	0.48		99			
Staphylococcus equorum		0.04	0.06		1			
043752	**Low discrimination**							**043752**
					OXI	NOV		
Staphylococcus sciuri		88.86	0.44	GLR 14	99	99		
Aerococcus viridans		5.92	0.36	MNS 89	1	1		
Staphylococcus lentus		3.87	0.18	LAC 14	99	99		
043753	**Low discrimination**							**043753**
					OXI	NOV		
Staphylococcus sciuri		63.78	0.63	GLR 14	99	99		
Staphylococcus lentus		34.01	0.59	LAC 14	99	99		
Aerococcus viridans		1.23	0.43		1	1		
043760	**Good identification**							**043760**
Staphylococcus sciuri		100.0	0.27	GLR 14				
043761	**Acceptable identification**							**043761**
					OXI			
Staphylococcus sciuri		93.04	0.45	GLR 14	99			
Staphylococcus gallinarum		2.81	0.08	MAN 97	1			
Staphylococcus equorum		1.54	0.06		1			
043763	**Low discrimination**							**043763**
					OXI	NOV	PYR	
Staphylococcus sciuri		39.89	0.29	GLR 14	99	99	1	
Staphylococcus gallinarum		18.13	0.16	MAN 97	1	99	12	
Aerococcus viridans		10.96	0.34	LAC 14	1	1	95	
043770	**Very good identification**							**043770**
					OXI	YEP		
Staphylococcus sciuri		98.85	0.58	GLR 14	99	57		
Staphylococcus equorum		0.41	0.06	MNS 89	1	1		
Staphylococcus arlettae		0.32	0.12		1	99		
043771	**Good identification**							**043771**
					OXI			
Staphylococcus sciuri		91.04	0.77	GLR 14	99			
Staphylococcus equorum		4.60	0.47		1			
Staphylococcus lentus		2.96	0.48		99			
043772	**Low discrimination**							**043772**
					OXI	NOV	YEP	
Staphylococcus arlettae		61.80	0.53	NIT 1	1	99	99	
Staphylococcus sciuri		31.24	0.42		99	99	57	
Aerococcus viridans		2.76	0.36		1	1	1	
043773	**Low discrimination**							**043773**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		32.43	0.60	GLR 14	99		
Staphylococcus equorum		25.87	0.55	LAC 14	1		
Staphylococcus lentus		22.92	0.59		99		
044000	**Low discrimination**						**044000**
Kocuria rosea		43.30	0.33	NIT 90			
Dermacoccus nishinomiyaensis		36.18	0.33	PHS 1			
Micrococcus sp.		19.55	0.18				
044012	**Good identification**						**044012**
Staphylococcus kloosii		100.0	0.27	TRE 99			
044050	**Low discrimination**						**044050**
					OXI	YEP	
Staphylococcus kloosii		78.91	0.52	TRE 99	1	99	
Staphylococcus cohnii subsp. cohnii		11.76	0.23		1	12	
Macrococcus bovicus		9.33	0.14		99	99	
044052	**Very good identification**						**044052**
Staphylococcus kloosii		100.0	0.53	TRE 99			
044070	**Good identification**						**044070**
Staphylococcus kloosii		100.0	0.40	TRE 99			
044072	**Good identification**						**044072**
Staphylococcus kloosii		100.0	0.41	TRE 99			
044110	**Low discrimination**						**044110**
					OXI		
Staphylococcus kloosii		82.46	0.32	TRE 99	1		
Macrococcus bovicus		17.54	-.01	MLT 95	99		
044112	**Good identification**						**044112**
Staphylococcus kloosii		100.0	0.33	TRE 99			
044150	**Very good identification**						**044150**
					OXI		
Staphylococcus kloosii		97.46	0.58	TRE 99	1		
Macrococcus bovicus		2.55	0.07		99		
044152	**Very good identification**						**044152**
Staphylococcus kloosii		100.0	0.59	TRE 99			
044170	**Good identification**						**044170**
Staphylococcus kloosii		100.0	0.46	TRE 99			
044172	**Acceptable identification**						**044172**
Staphylococcus kloosii		90.04	0.47	TRE 99			
Staphylococcus arlettae		9.96	0.15				
044240	**Very good identification**						**044240**
Staphylococcus aureus subsp. anaerobius		100.0	0.57	GLR 1			
044252	**Good identification**						**044252**
Staphylococcus kloosii		100.0	0.32	SUC 9			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
044272	**Good identification**						**044272**
	Staphylococcus arlettae	97.43	0.41	TRE 99			
	Staphylococcus kloosii	2.57	0.20				
044273	**Good identification**						**044273**
	Staphylococcus arlettae	100.0	0.43	TRE 99			
044350	**Good identification**						**044350**
	Staphylococcus kloosii	100.0	0.38	SUC 9			
044352	**Low discrimination**						**044352**
	Staphylococcus kloosii	78.16	0.38	SUC 9			
	Staphylococcus arlettae	21.85	0.15	TRE 99			
044372	**Very good identification**						**044372**
	Staphylococcus arlettae	99.11	0.56	TRE 99			
	Staphylococcus kloosii	0.90	0.26				
044373	**Very good identification**						**044373**
	Staphylococcus arlettae	100.0	0.59	TRE 99			
044400	**Good identification**						**044400**
					YEP		
	Staphylococcus cohnii subsp. cohnii	98.58	0.51	PHS 9	12		
	Staphylococcus kloosii	1.42	0.26		99		
044401	**Good identification**						**044401**
	Staphylococcus cohnii subsp. cohnii	100.0	0.52	PHS 9			
044402	**Low discrimination**						**044402**
					YEP		
	Staphylococcus kloosii	60.66	0.27	MAN 99	99		
	Staphylococcus cohnii subsp. cohnii	39.34	0.10	MLT 95	12		
044410	**Low discrimination**						**044410**
					OXI	YEP	
	Staphylococcus kloosii	52.39	0.67	MLT 95	1	99	
	Staphylococcus cohnii subsp. cohnii	46.85	0.54		1	12	
	Macrococcus bovicus	0.77	0.10		99	99	
044411	**Very good identification**						**044411**
					YEP		
	Staphylococcus cohnii subsp. cohnii	97.22	0.54	PHS 9	12		
	Staphylococcus cohnii subsp. urealyticum	1.77	0.19		64		
	Staphylococcus kloosii	1.01	0.26		99		
044412	**Very good identification**						**044412**
					YEP		
	Staphylococcus kloosii	99.17	0.68	MLT 95	99		
	Staphylococcus cohnii subsp. cohnii	0.83	0.12		12		
044413	**Low discrimination**						**044413**
					YEP		
	Staphylococcus cohnii subsp. urealyticum	89.57	0.40	URE 99	64		
	Staphylococcus kloosii	5.51	0.27		99		
	Staphylococcus cohnii subsp. cohnii	4.93	0.13		12		
044430	**Very good identification**						**044430**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					YEP		
Staphylococcus kloosii		96.51	0.55	MLT 95	99		
Staphylococcus cohnii subsp. cohnii		3.49	0.12		12		
044432	**Very good identification**						**044432**
Staphylococcus kloosii		100.0	0.55	MLT 95			
044440	**Good identification**						**044440**
					NOV	YEP	
Staphylococcus cohnii subsp. cohnii		91.89	0.62	PHS 9	99	12	
Staphylococcus kloosii		7.93	0.52		99	99	
Staphylococcus auricularis		0.18	0.03		1	1	
044441	**Very good identification**						**044441**
Staphylococcus cohnii subsp. cohnii		99.59	0.62	PHS 9			
Staphylococcus cohnii subsp. urealyticum		0.41	0.14				
044442	**Good identification**						**044442**
					YEP		
Staphylococcus kloosii		90.24	0.53	MAN 99	99		
Staphylococcus cohnii subsp. cohnii		9.76	0.21		12		
044443	**Low discrimination**						**044443**
Staphylococcus cohnii subsp. urealyticum		80.37	0.35	URE 99			
Staphylococcus cohnii subsp. cohnii		19.63	0.21	MAN 91			
044450	**Low discrimination**						**044450**
					OXI	YEP	
Staphylococcus kloosii		86.89	0.93		1	99	
Staphylococcus cohnii subsp. cohnii		12.95	0.64		1	12	
Macrococcus bovicus		0.16	0.18		99	99	
044451	**Good identification**						**044451**
					YEP		
Staphylococcus cohnii subsp. cohnii		91.32	0.65	PHS 9	12		
Staphylococcus kloosii		5.71	0.52		99		
Staphylococcus cohnii subsp. urealyticum		2.97	0.35		64		
044452	**Excellent identification**						**044452**
					YEP		
Staphylococcus kloosii		99.79	0.94		99		
Staphylococcus cohnii subsp. cohnii		0.14	0.23		12		
Staphylococcus cohnii subsp. urealyticum		0.05	0.14		64		
044453	**Low discrimination**						**044453**
					YEP		
Staphylococcus cohnii subsp. urealyticum		80.29	0.55	URE 99	64		
Staphylococcus kloosii		16.56	0.53		99		
Staphylococcus cohnii subsp. cohnii		2.47	0.23		12		
044460	**Low discrimination**						**044460**
					YEP		
Staphylococcus kloosii		68.10	0.40	MAN 99	99		
Staphylococcus cohnii subsp. cohnii		31.90	0.21		12		
044462	**Good identification**						**044462**
Staphylococcus kloosii		100.0	0.41	MAN 99			
044470	**Very good identification**						**044470**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		99.40	0.81		99		
Staphylococcus cohnii subsp. cohnii		0.60	0.23		12		
044471	**Low discrimination**						**044471**
					YEP		
Staphylococcus kloosii		60.76	0.40	MNS 1	99		
Staphylococcus cohnii subsp. cohnii		39.24	0.23		12		
044472	**Very good identification**						**044472**
					NOV	YEP	
Staphylococcus kloosii		96.28	0.82		99	99	
Staphylococcus arlettae		3.65	0.41		99	99	
Aerococcus viridans		0.07	0.16		1	1	
044473	**Low discrimination**						**044473**
Staphylococcus arlettae		78.52	0.43	SUC 99			
Staphylococcus kloosii		15.80	0.41				
Staphylococcus cohnii subsp. urealyticum		3.10	0.14				
044500	**Low discrimination**						**044500**
					YEP		
Staphylococcus cohnii subsp. cohnii		79.93	0.32	PHS 9	12		
Staphylococcus kloosii		20.07	0.32	GAL 10	99		
044501	**Good identification**						**044501**
Staphylococcus cohnii subsp. cohnii		100.0	0.32	PHS 9			
044502	**Good identification**						**044502**
Staphylococcus kloosii		100.0	0.33	MAN 99			
044510	**Good identification**						**044510**
					OXI	YEP	
Staphylococcus kloosii		94.84	0.73	MLT 95	1	99	
Staphylococcus cohnii subsp. cohnii		4.86	0.34		1	12	
Macrococcus bovicus		0.31	0.03		99	99	
044511	**Low discrimination**						**044511**
					YEP		
Staphylococcus cohnii subsp. cohnii		79.86	0.35	PHS 9	12		
Staphylococcus kloosii		14.55	0.32	GAL 10	99		
Staphylococcus cohnii subsp. urealyticum		5.59	0.12		64		
044512	**Very good identification**						**044512**
Staphylococcus kloosii		100.0	0.74	MLT 95			
044513	**Low discrimination**						**044513**
Staphylococcus cohnii subsp. urealyticum		78.21	0.32	URE 99			
Staphylococcus kloosii		21.79	0.33				
044530	**Very good identification**						**044530**
Staphylococcus kloosii		100.0	0.61	MLT 95			
044532	**Very good identification**						**044532**
Staphylococcus kloosii		97.92	0.61	MLT 95			
Staphylococcus arlettae		2.08	0.15				
044540	**Low discrimination**						**044540**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		60.11	0.58	MAN 99	99		
Staphylococcus cohnii subsp. cohnii		39.89	0.42		12		
044541	**Good identification**						**044541**
					YEP		
Staphylococcus cohnii subsp. cohnii		97.10	0.43	PHS 9	12		
Staphylococcus cohnii subsp. urealyticum		1.53	0.06	GAL 10	64		
Staphylococcus kloosii		1.36	0.17		99		
044542	**Very good identification**						**044542**
Staphylococcus kloosii		100.0	0.59	MAN 99			
044543	**Acceptable identification**						**044543**
Staphylococcus cohnii subsp. urealyticum		91.29	0.27	URE 99			
Staphylococcus kloosii		8.71	0.18	MAN 91			
044550	**Excellent identification**						**044550**
					OXI	YEP	
Staphylococcus kloosii		99.12	0.99		1	99	
Staphylococcus cohnii subsp. cohnii		0.85	0.44		1	12	
Macrococcus bovicus		0.04	0.10		99	99	
044551	**Low discrimination**						**044551**
					YEP		
Staphylococcus kloosii		49.27	0.58	MNS 1	99		
Staphylococcus cohnii subsp. cohnii		45.08	0.45		12		
Staphylococcus cohnii subsp. urealyticum		5.65	0.27		64		
044552	**Excellent identification**						**044552**
					NOV	YEP	
Staphylococcus kloosii		99.95	1.00		99	99	
Staphylococcus arlettae		0.03	0.15		99	99	
Aerococcus viridans		0.01	0.16		1	1	
044553	**Low discrimination**						**044553**
Staphylococcus cohnii subsp. urealyticum		50.10	0.48	URE 99			
Staphylococcus kloosii		46.80	0.59				
Staphylococcus arlettae		1.72	0.18				
044560	**Good identification**						**044560**
Staphylococcus kloosii		100.0	0.46	MAN 99			
044562	**Acceptable identification**						**044562**
Staphylococcus kloosii		90.04	0.47	MAN 99			
Staphylococcus arlettae		9.96	0.15				
044570	**Very good identification**						**044570**
Staphylococcus kloosii		99.88	0.87				
Staphylococcus arlettae		0.12	0.15				
044571	**Low discrimination**						**044571**
					YEP		
Staphylococcus kloosii		83.63	0.46	MNS 1	99		
Staphylococcus arlettae		13.28	0.18		99		
Staphylococcus cohnii subsp. cohnii		3.09	0.04		12		
044572	**Good identification**						**044572**
					NOV	YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		90.01	0.88		99	99	
Staphylococcus arlettae		9.95	0.56		99	99	
Aerococcus viridans		0.03	0.16		1	1	
044573	**Good identification**						**044573**
					NOV	YEP	
Staphylococcus arlettae		92.79	0.59	SUC 99	99	99	
Staphylococcus kloosii		6.39	0.47		99	99	
Aerococcus viridans		0.54	0.24		1	1	
044610	**Low discrimination**						**044610**
					OXI	YEP	
Staphylococcus kloosii		88.63	0.46	SUC 9	1	99	
Staphylococcus cohnii subsp. cohnii		8.10	0.12	MLT 95	1	12	
Macrococcus bovicus		3.27	-.02		99	99	
044612	**Good identification**						**044612**
Staphylococcus kloosii		100.0	0.47	SUC 9			
044620	**Good identification**						**044620**
Staphylococcus muscae		100.0	0.39	GLR 10			
044630	**Low discrimination**						**044630**
					NOV	YEP	
Staphylococcus kloosii		54.03	0.34	SUC 9	99	99	
Staphylococcus muscae		45.97	-.02	MLT 95	1	1	
044632	**Low discrimination**						**044632**
Staphylococcus arlettae		87.92	0.41	MLT 99			
Staphylococcus kloosii		12.08	0.35				
044633	**Good identification**						**044633**
Staphylococcus arlettae		100.0	0.43	MLT 99			
044642	**Low discrimination**						**044642**
					NOV		
Staphylococcus kloosii		85.79	0.32	SUC 9	99		
Staphylococcus warneri		14.21	0.09	MAN 99	1		
044650	**Very good identification**						**044650**
					OXI	YEP	
Staphylococcus kloosii		97.69	0.73	SUC 9	1	99	
Staphylococcus cohnii subsp. cohnii		1.49	0.23		1	12	
Macrococcus bovicus		0.44	0.05		99	99	
044652	**Good identification**						**044652**
					NOV	YEP	
Staphylococcus kloosii		91.11	0.73	SUC 9	99	99	
Staphylococcus arlettae		8.72	0.41		99	99	
Aerococcus viridans		0.16	0.16		1	1	
044653	**Low discrimination**						**044653**
Staphylococcus arlettae		81.09	0.43	XYL 99			
Staphylococcus cohnii subsp. urealyticum		9.79	0.24				
Staphylococcus kloosii		6.45	0.32				
044662	**Good identification**						**044662**
Staphylococcus arlettae		97.43	0.41	MAN 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		2.57	0.20				
044663	**Good identification**						**044663**
Staphylococcus arlettae		100.0	0.43	MAN 99			
044670	**Low discrimination**						**044670**
Staphylococcus kloosii		70.68	0.60	SUC 9			
Staphylococcus arlettae		29.33	0.41				
044671	**Good identification**						**044671**
Staphylococcus arlettae		98.20	0.43	LAC 99			
Staphylococcus kloosii		1.80	0.19				
044672	**Very good identification**						**044672**
					NOV	YEP	
Staphylococcus arlettae		97.41	0.82		99	99	
Staphylococcus kloosii		2.57	0.61		99	99	
Aerococcus viridans		0.02	0.16		1	1	
044673	**Very good identification**						**044673**
					NOV	YEP	
Staphylococcus arlettae		99.93	0.85		99	99	
Aerococcus viridans		0.03	0.24		1	1	
Staphylococcus kloosii		0.02	0.20		99	99	
044703	**Good identification**						**044703**
Staphylococcus hyicus		100.0	0.29	ARG 96			
044710	**Good identification**						**044710**
Staphylococcus kloosii		100.0	0.52	SUC 9			
044712	**Good identification**						**044712**
Staphylococcus kloosii		94.91	0.53	SUC 9			
Staphylococcus arlettae		5.09	0.15	MLT 95			
044730	**Low discrimination**						**044730**
Staphylococcus kloosii		81.14	0.40	SUC 9			
Staphylococcus arlettae		18.86	0.15	MLT 95			
044732	**Very good identification**						**044732**
Staphylococcus arlettae		95.50	0.56	MLT 99			
Staphylococcus kloosii		4.50	0.41				
044733	**Very good identification**						**044733**
Staphylococcus arlettae		100.0	0.59	MLT 99			
044740	**Good identification**						**044740**
Staphylococcus kloosii		100.0	0.38	SUC 9			
044742	**Low discrimination**						**044742**
Staphylococcus kloosii		78.16	0.38	SUC 9			
Staphylococcus arlettae		21.85	0.15	MAN 99			
044750	**Very good identification**						**044750**
					OXI		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		99.60	0.79	SUC 9	1		
Staphylococcus arlettae		0.31	0.15		1		
Macrococcus bovicus		0.10	-.02		99		
044751	**Low discrimination**						**044751**
					OXI		
Staphylococcus kloosii		57.86	0.38	SUC 9	1		
Staphylococcus arlettae		23.22	0.18	MNS 1	1		
Staphylococcus sciuri		13.52	0.08		99		
044752	**Low discrimination**						**044752**
					NOV	YEP	
Staphylococcus kloosii		78.10	0.79	SUC 9	99	99	
Staphylococcus arlettae		21.83	0.56		99	99	
Aerococcus viridans		0.07	0.16		1	1	
044753	**Very good identification**						**044753**
Staphylococcus arlettae		95.95	0.59	XYL 99			
Staphylococcus kloosii		2.62	0.38				
Staphylococcus cohnii subsp. urealyticum		0.88	0.17				
044762	**Very good identification**						**044762**
Staphylococcus arlettae		99.11	0.56	MAN 99			
Staphylococcus kloosii		0.90	0.26				
044763	**Very good identification**						**044763**
Staphylococcus arlettae		100.0	0.59	MAN 99			
044770	**Low discrimination**						**044770**
Staphylococcus arlettae		54.78	0.56	LAC 99			
Staphylococcus kloosii		45.22	0.66				
044771	**Very good identification**						**044771**
Staphylococcus arlettae		98.08	0.59	LAC 99			
Staphylococcus xylosus		0.86	0.07				
Staphylococcus kloosii		0.62	0.25				
044772	**Excellent identification**						**044772**
					NOV	YEP	
Staphylococcus arlettae		99.10	0.98		99	99	
Staphylococcus kloosii		0.90	0.67		99	99	
Aerococcus viridans		0.00	0.16		1	1	
044773	**Excellent identification**						**044773**
Staphylococcus arlettae		99.96	1.00				
Staphylococcus xylosus		0.03	0.17				
Staphylococcus kloosii		0.01	0.26				
045012	**Good identification**						**045012**
Staphylococcus kloosii		100.0	0.27	TRE 99			
045050	**Good identification**						**045050**
					OXI		
Staphylococcus kloosii		97.61	0.52	TRE 99	1		
Macrococcus equipersicus		2.39	0.02		99		
045052	**Very good identification**						**045052**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.53	TRE 99			
045070	**Good identification**						**045070**
Staphylococcus kloosii		100.0	0.40	TRE 99			
045072	**Good identification**						**045072**
Staphylococcus kloosii		100.0	0.41	TRE 99			
045110	**Good identification**						**045110**
Staphylococcus kloosii		100.0	0.32	TRE 99			
045112	**Good identification**						**045112**
Staphylococcus kloosii		100.0	0.33	TRE 99			
045150	**Very good identification**						**045150**
Staphylococcus kloosii		100.0	0.58	TRE 99			
045152	**Very good identification**						**045152**
Staphylococcus kloosii		100.0	0.59	TRE 99			
045170	**Good identification**						**045170**
Staphylococcus kloosii		100.0	0.46	TRE 99			
045172	**Acceptable identification**						**045172**
Staphylococcus kloosii		90.04	0.47	TRE 99			
Staphylococcus arlettae		9.96	0.15				
045250	**Good identification**						**045250**
Staphylococcus kloosii		100.0	0.32	SUC 9			
045252	**Good identification**						**045252**
Staphylococcus kloosii		100.0	0.32	SUC 9			
045272	**Good identification**						**045272**
Staphylococcus arlettae		97.43	0.41	TRE 99			
Staphylococcus kloosii		2.57	0.20				
045273	**Good identification**						**045273**
					NOV	YEP	
Staphylococcus arlettae		98.46	0.43	TRE 99	99	99	
Aerococcus viridans		1.54	0.17		1	1	
045350	**Good identification**						**045350**
Staphylococcus kloosii		100.0	0.38	SUC 9			
045352	**Low discrimination**						**045352**
Staphylococcus kloosii		78.16	0.38	SUC 9			
Staphylococcus arlettae		21.85	0.15	TRE 99			
045372	**Very good identification**						**045372**
Staphylococcus arlettae		99.11	0.56	TRE 99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		0.90	0.26					
045373	**Very good identification**							**045373**
					NOV	PYR		
Staphylococcus arlettae		99.43	0.59	TRE 99	99	1		
Staphylococcus gallinarum		0.29	-.04		99	12		
Aerococcus viridans		0.27	0.17		1	95		
045402	**Good identification**							**045402**
Staphylococcus kloosii		100.0	0.27	MAN 99				
045410	**Very good identification**							**045410**
					OXI	YEP		
Staphylococcus kloosii		97.51	0.67	MLT 95	1	99		
Macrococcus carouselicus		1.61	0.08		99	67		
Staphylococcus cohnii subsp. cohnii		0.88	0.12		1	12		
045412	**Very good identification**							**045412**
					NOV	YEP		
Staphylococcus kloosii		99.25	0.68	MLT 95	99	99		
Aerococcus viridans		0.75	0.23		1	1		
045413	**Low discrimination**							**045413**
					NOV	YEP		
Staphylococcus cohnii subsp. urealyticum		69.66	0.31	URE 99	99	64		
Aerococcus viridans		19.32	0.31		1	1		
Staphylococcus kloosii		11.02	0.27		99	99		
045430	**Very good identification**							**045430**
Staphylococcus kloosii		100.0	0.55	MLT 95				
045432	**Very good identification**							**045432**
					NOV	YEP		
Staphylococcus kloosii		97.05	0.55	MLT 95	99	99		
Aerococcus viridans		2.95	0.23		1	1		
045433	**Good identification**							**045433**
Aerococcus viridans		100.0	0.31	PHS 1				
045440	**Low discrimination**							**045440**
					YEP			
Staphylococcus kloosii		89.52	0.52	MAN 99	99			
Staphylococcus cohnii subsp. cohnii		10.48	0.21		12			
045442	**Good identification**							**045442**
					NOV	YEP		
Staphylococcus kloosii		94.99	0.53	MAN 99	99	99		
Aerococcus viridans		5.01	0.26		1	1		
045450	**Very good identification**							**045450**
					NOV	YEP	PYR	
Staphylococcus kloosii		99.79	0.93		99	99	52	
Staphylococcus cohnii subsp. cohnii		0.15	0.23		99	12	1	
Aerococcus viridans		0.02	0.16		1	1	95	
045451	**Low discrimination**							**045451**
					YEP			
Staphylococcus kloosii		69.88	0.52	MNS 1	99			
Staphylococcus cohnii subsp. urealyticum		14.12	0.26		64			
Staphylococcus cohnii subsp. cohnii		11.28	0.23		12			
045452	**Excellent identification**							**045452**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus kloosii		99.82	0.94		99	99	
Aerococcus viridans		0.16	0.36		1	1	
Staphylococcus cohnii subsp. urealyticum		0.02	0.06		99	64	
045453	**Low discrimination**						**045453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		57.94	0.47	URE 99	99	64	
Staphylococcus kloosii		30.72	0.53		99	99	
Aerococcus viridans		11.34	0.43		1	1	
045460	**Good identification**						**045460**
Staphylococcus kloosii		100.0	0.40	MAN 99			
045462	**Low discrimination**						**045462**
					NOV	YEP	
Staphylococcus kloosii		82.58	0.41	MAN 99	99	99	
Aerococcus viridans		17.42	0.26		1	1	
045463	**Good identification**						**045463**
Aerococcus viridans		100.0	0.34	PHS 1			
045470	**Very good identification**						**045470**
					NOV	YEP	
Staphylococcus kloosii		99.92	0.81		99	99	
Aerococcus viridans		0.08	0.16		1	1	
045471	**Low discrimination**						**045471**
					NOV	YEP	
Staphylococcus kloosii		84.91	0.40	MNS 1	99	99	
Aerococcus viridans		15.09	0.24		1	1	
045472	**Very good identification**						**045472**
					NOV	YEP	
Staphylococcus kloosii		95.76	0.82		99	99	
Staphylococcus arlettae		3.63	0.41		99	99	
Aerococcus viridans		0.61	0.36		1	1	
045473	**Low discrimination**						**045473**
					NOV	YEP	
Staphylococcus arlettae		66.07	0.43	SUC 99	99	99	
Aerococcus viridans		19.63	0.43		1	1	
Staphylococcus kloosii		13.29	0.41		99	99	
045500	**Low discrimination**						**045500**
					OXI		
Staphylococcus kloosii		63.86	0.32	MAN 99	1		
Macrococcus carouselicus		36.14	0.04	MLT 95	99		
045502	**Good identification**						**045502**
Staphylococcus kloosii		100.0	0.33	MAN 99			
045510	**Very good identification**						**045510**
					OXI		
Staphylococcus kloosii		99.64	0.73	MLT 95	1		
Macrococcus carouselicus		0.36	0.00		99		
045511	**Good identification**						**045511**
Staphylococcus kloosii		100.0	0.32	MLT 95			
045512	**Very good identification**						**045512**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus kloosii		99.61	0.74	MLT 95	99	99	
Aerococcus viridans		0.39	0.23		1	1	
045530	**Very good identification**						**045530**
Staphylococcus kloosii		100.0	0.61	MLT 95			
045532	**Very good identification**						**045532**
					NOV	YEP	
Staphylococcus kloosii		96.45	0.61	MLT 95	99	99	
Staphylococcus arlettae		2.05	0.15		99	99	
Aerococcus viridans		1.51	0.23		1	1	
045533	**Low discrimination**						**045533**
					NOV	YEP	
Aerococcus viridans		48.84	0.31	PHS 1	1	1	
Staphylococcus arlettae		37.65	0.18		99	99	
Staphylococcus kloosii		13.52	0.20		99	99	
045540	**Very good identification**						**045540**
Staphylococcus kloosii		100.0	0.58	MAN 99			
045542	**Very good identification**						**045542**
					NOV	YEP	
Staphylococcus kloosii		97.36	0.59	MAN 99	99	99	
Aerococcus viridans		2.65	0.26		1	1	
045543	**Low discrimination**						**045543**
					NOV	YEP	
Aerococcus viridans		55.28	0.34	PHS 1	1	1	
Staphylococcus cohnii subsp. urealyticum		35.91	0.19		99	64	
Staphylococcus kloosii		8.81	0.18		99	99	
045550	**Excellent identification**						**045550**
					NOV	YEP	
Staphylococcus kloosii		99.99	0.99		99	99	
Aerococcus viridans		0.01	0.16		1	1	
045551	**Good identification**						**045551**
					OXI		
Staphylococcus kloosii		90.92	0.58	MNS 1	1		
Staphylococcus cohnii subsp. urealyticum		4.06	0.19		1		
Staphylococcus sciuri		2.10	0.08		99		
045552	**Excellent identification**						**045552**
					NOV	YEP	
Staphylococcus kloosii		99.89	1.00		99	99	
Aerococcus viridans		0.08	0.36		1	1	
Staphylococcus arlettae		0.03	0.15		99	99	
045553	**Low discrimination**						**045553**
					NOV	YEP	
Staphylococcus kloosii		60.86	0.59	MNS 1	99	99	
Staphylococcus cohnii subsp. urealyticum		25.34	0.39		99	64	
Aerococcus viridans		11.57	0.43		1	1	
045560	**Good identification**						**045560**
Staphylococcus kloosii		100.0	0.46	MAN 99			
045562	**Low discrimination**						**045562**
					NOV	YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		82.02	0.47	MAN 99	99	99	
Staphylococcus arlettae		9.07	0.15		99	99	
Aerococcus viridans		8.91	0.26		1	1	
045563	**Low discrimination**						**045563**
					NOV	YEP	
Aerococcus viridans		63.37	0.34	PHS 1	1	1	
Staphylococcus arlettae		36.63	0.18		99	99	
045570	**Very good identification**						**045570**
					NOV	YEP	
Staphylococcus kloosii		99.84	0.87		99	99	
Staphylococcus arlettae		0.12	0.15		99	99	
Aerococcus viridans		0.04	0.16		1	1	
045571	**Low discrimination**						**045571**
					NOV	YEP	
Staphylococcus kloosii		75.75	0.46	MNS 1	99	99	
Staphylococcus arlettae		12.03	0.18		99	99	
Aerococcus viridans		6.94	0.24		1	1	
045572	**Low discrimination**						**045572**
					NOV	YEP	
Staphylococcus kloosii		89.78	0.88		99	99	
Staphylococcus arlettae		9.93	0.56		99	99	
Aerococcus viridans		0.30	0.36		1	1	
045573	**Low discrimination**						**045573**
					NOV	YEP	
Staphylococcus arlettae		88.95	0.59	SUC 99	99	99	
Staphylococcus kloosii		6.13	0.47		99	99	
Aerococcus viridans		4.66	0.43		1	1	
045610	**Low discrimination**						**045610**
					OXI		
Staphylococcus kloosii		67.31	0.46	SUC 9	1		
Macrococcus carouselicus		22.87	0.14	MLT 95	99		
Staphylococcus sciuri		9.82	0.13		99		
045611	**Good identification**						**045611**
Staphylococcus sciuri		100.0	0.31	GLR 14			
045612	**Acceptable identification**						**045612**
					NOV	YEP	
Staphylococcus kloosii		92.87	0.47	SUC 9	99	99	
Aerococcus viridans		7.13	0.23	MLT 95	1	1	
045613	**Low discrimination**						**045613**
					OXI	NOV	
Aerococcus viridans		50.25	0.31	PHS 1	1	1	
Staphylococcus sciuri		49.75	0.15		99	99	
045630	**Low discrimination**						**045630**
					OXI		
Staphylococcus kloosii		62.84	0.34	SUC 9	1		
Staphylococcus sciuri		27.65	0.10	MLT 95	99		
Staphylococcus vitulinus		9.51	-.08		99		
045631	**Good identification**						**045631**
Staphylococcus sciuri		100.0	0.29	GLR 14			
045632	**Low discrimination**						**045632**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus arlettae		84.77	0.41	MLT 99	99	99		
Staphylococcus kloosii		11.65	0.35		99	99		
Aerococcus viridans		3.58	0.23		1	1		
045633	**Low discrimination**							**045633**
					OXI	NOV	YEP	
Staphylococcus arlettae		88.52	0.43	MLT 99	1	99	99	
Aerococcus viridans		6.57	0.31		1	1	1	
Staphylococcus sciuri		4.91	0.13		99	99	57	
045640	**Good identification**							**045640**
Staphylococcus kloosii		100.0	0.32	SUC 9				
045642	**Low discrimination**							**045642**
					NOV	YEP		
Staphylococcus kloosii		65.22	0.32	SUC 9	99	99		
Aerococcus viridans		34.78	0.26	MAN 99	1	1		
045643	**Good identification**							**045643**
Aerococcus viridans		100.0	0.34	PHS 1				
045650	**Very good identification**							**045650**
					OXI	NOV	YEP	
Staphylococcus kloosii		97.01	0.73	SUC 9	1	99	99	
Staphylococcus sciuri		2.80	0.24		99	99	57	
Aerococcus viridans		0.19	0.16		1	1	1	
045651	**Acceptable identification**							**045651**
					OXI	NOV	YEP	
Staphylococcus sciuri		94.10	0.43	GLR 14	99	99	57	
Staphylococcus kloosii		4.07	0.32	NIT 98	1	99	99	
Aerococcus viridans		1.83	0.24		1	1	1	
045652	**Low discrimination**							**045652**
					NOV	YEP		
Staphylococcus kloosii		89.59	0.73	SUC 9	99	99		
Staphylococcus arlettae		8.58	0.41		99	99		
Aerococcus viridans		1.45	0.36		1	1		
045653	**Low discrimination**							**045653**
					OXI	NOV	YEP	
Staphylococcus arlettae		58.82	0.43	XYL 99	1	99	99	
Aerococcus viridans		17.47	0.43		1	1	1	
Staphylococcus sciuri		16.27	0.27		99	99	57	
045662	**Acceptable identification**							**045662**
					NOV	YEP		
Staphylococcus arlettae		92.37	0.41	MAN 99	99	99		
Aerococcus viridans		5.20	0.26		1	1		
Staphylococcus kloosii		2.44	0.20		99	99		
045663	**Acceptable identification**							**045663**
					NOV	YEP		
Staphylococcus arlettae		90.99	0.43	MAN 99	99	99		
Aerococcus viridans		9.01	0.34		1	1		
045670	**Low discrimination**							**045670**
					OXI			
Staphylococcus kloosii		66.23	0.60	SUC 9	1			
Staphylococcus arlettae		27.48	0.41		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		5.77	0.22		99			
045671	**Low discrimination**							**045671**
					OXI	YEP		
Staphylococcus sciuri		53.86	0.41	GLR 14	99	57		
Staphylococcus arlettae		42.02	0.43	NIT 98	1	99		
Staphylococcus equorum		1.96	0.08		1	1		
045672	**Very good identification**							**045672**
					NOV	YEP		
Staphylococcus arlettae		97.24	0.82		99	99		
Staphylococcus kloosii		2.56	0.61		99	99		
Aerococcus viridans		0.17	0.36		1	1		
045673	**Very good identification**							**045673**
					OXI	NOV	YEP	
Staphylococcus arlettae		99.34	0.85		1	99	99	
Aerococcus viridans		0.30	0.43		1	1	1	
Staphylococcus sciuri		0.21	0.24		99	99	57	
045710	**Low discrimination**							**045710**
					OXI			
Staphylococcus kloosii		81.91	0.52	SUC 9	1			
Staphylococcus sciuri		11.94	0.18	MLT 95	99			
Macrococcus carouselicus		6.15	0.06		99			
045711	**Good identification**							**045711**
Staphylococcus sciuri		100.0	0.37	GLR 14				
045712	**Low discrimination**							**045712**
					NOV	YEP		
Staphylococcus kloosii		89.68	0.53	SUC 9	99	99		
Staphylococcus arlettae		4.81	0.15	MLT 95	99	99		
Aerococcus viridans		3.55	0.23		1	1		
045730	**Low discrimination**							**045730**
					OXI			
Staphylococcus kloosii		59.79	0.40	SUC 9	1			
Staphylococcus sciuri		26.31	0.16	MLT 95	99			
Staphylococcus arlettae		13.90	0.15		1			
045731	**Low discrimination**							**045731**
					OXI	YEP		
Staphylococcus sciuri		86.42	0.35	GLR 14	99	57		
Staphylococcus arlettae		7.48	0.18	NIT 98	1	99		
Staphylococcus equorum		6.10	0.08		1	1		
045732	**Good identification**							**045732**
					NOV	YEP		
Staphylococcus arlettae		94.55	0.56	MLT 99	99	99		
Staphylococcus kloosii		4.45	0.41		99	99		
Aerococcus viridans		0.70	0.23		1	1		
045733	**Good identification**							**045733**
					OXI	YEP		
Staphylococcus arlettae		94.68	0.59	MLT 99	1	99		
Staphylococcus equorum		2.00	0.16		1	1		
Staphylococcus sciuri		1.80	0.18		99	57		
045740	**Good identification**							**045740**
Staphylococcus kloosii		100.0	0.38	SUC 9				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
045742	**Low discrimination**							**045742**
					NOV	YEP		
Staphylococcus kloosii		64.34	0.38	SUC 9	99	99		
Staphylococcus arlettae		17.99	0.15	MAN 99	99	99		
Aerococcus viridans		17.67	0.26		1	1		
045743	**Low discrimination**							**045743**
					OXI	NOV	YEP	
Aerococcus viridans		57.28	0.34	PHS 1	1	1	1	
Staphylococcus arlettae		33.11	0.18		1	99	99	
Staphylococcus sciuri		9.61	0.02		99	99	57	
045750	**Very good identification**							**045750**
					OXI			
Staphylococcus kloosii		96.81	0.79	SUC 9	1			
Staphylococcus sciuri		2.80	0.30		99			
Staphylococcus arlettae		0.30	0.15		1			
045751	**Acceptable identification**							**045751**
					OXI			
Staphylococcus sciuri		93.41	0.49	GLR 14	99			
Staphylococcus kloosii		4.04	0.38	NIT 98	1			
Staphylococcus arlettae		1.62	0.18		1			
045752	**Low discrimination**							**045752**
					NOV	YEP		
Staphylococcus kloosii		77.39	0.79	SUC 9	99	99		
Staphylococcus arlettae		21.63	0.56		99	99		
Aerococcus viridans		0.64	0.36		1	1		
045753	**Low discrimination**							**045753**
					OXI	NOV	YEP	
Staphylococcus arlettae		84.42	0.59	XYL 99	1	99	99	
Staphylococcus sciuri		8.00	0.33		99	99	57	
Aerococcus viridans		4.43	0.43		1	1	1	
045762	**Very good identification**							**045762**
					NOV	YEP		
Staphylococcus arlettae		98.14	0.56	MAN 99	99	99		
Aerococcus viridans		0.97	0.26		1	1		
Staphylococcus kloosii		0.89	0.26		99	99		
045763	**Very good identification**							**045763**
					NOV	PYR		
Staphylococcus arlettae		95.57	0.59	MAN 99	99	1		
Staphylococcus gallinarum		2.76	0.16		99	12		
Aerococcus viridans		1.67	0.34		1	95		
045770	**Low discrimination**							**045770**
					OXI			
Staphylococcus arlettae		52.61	0.56	LAC 99	1			
Staphylococcus kloosii		43.43	0.66		1			
Staphylococcus sciuri		3.78	0.28		99			
045771	**Low discrimination**							**045771**
					OXI			
Staphylococcus arlettae		60.47	0.59	LAC 99	1			
Staphylococcus sciuri		26.55	0.47		99			
Staphylococcus gallinarum		7.22	0.29		1			
045772	**Excellent identification**							**045772**
					NOV	YEP		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		99.06	0.98		99	99	
Staphylococcus kloosii		0.90	0.67		99	99	
Aerococcus viridans		0.03	0.36		1	1	
045773	**Excellent identification**						**045773**
					YEP		
Staphylococcus arlettae		99.35	1.00		99		
Staphylococcus gallinarum		0.29	0.37		78		
Staphylococcus equorum		0.22	0.37		1		
046000	**Acceptable identification**						**046000**
Kocuria rosea		94.69	0.53	PHS 1			
Dermacoccus nishinomiyaensis		4.95	0.28				
Micrococcus sp.		0.25	-.09				
046010	**Good identification**						**046010**
Kocuria rosea		100.0	0.33	PHS 1			
046020	**Good identification**						**046020**
Kocuria rosea		98.30	0.45	PHS 1			
Staphylococcus muscae		1.70	-.02				
046200	**Acceptable identification**						**046200**
					OXI		
Kocuria rosea		91.17	0.33	PHS 1	35		
Staphylococcus muscae		6.10	-.02	SUC 10	1		
Dermacoccus nishinomiyaensis		2.74	0.03		93		
046211	**Good identification**						**046211**
Staphylococcus capitis subsp. capitis		100.0	0.29	GLR 1			
046220	**Acceptable identification**						**046220**
Staphylococcus muscae		93.92	0.39	GLR 10			
Kocuria rosea		6.08	0.25	TRE 99			
046303	**Good identification**						**046303**
Staphylococcus hyicus		100.0	0.48	ARG 96			
046400	**Low discrimination**						**046400**
					NOV		
Staphylococcus cohnii subsp. cohnii		44.17	0.28	NIT 7	99		
Kocuria rosea		23.88	0.12	PHS 9	1		
Staphylococcus muscae		17.57	-.02		1		
046401	**Acceptable identification**						**046401**
					NOV		
Staphylococcus cohnii subsp. cohnii		93.26	0.29	NIT 7	99		
Staphylococcus schleiferi subsp. schleiferi		6.74	-.02	PHS 9	1		
046410	**Low discrimination**						**046410**
					YEP		
Staphylococcus cohnii subsp. cohnii		86.95	0.30	NIT 7	12		
Staphylococcus kloosii		13.05	0.26	PHS 9	99		
046411	**Good identification**						**046411**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	NIT 7			
046412	**Good identification**						**046412**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.27	NIT 1			
046420	**Good identification**						**046420**
Staphylococcus muscae		99.42	0.39	GLR 10			
Kocuria rosea		0.59	0.04	SUC 99			
046440	**Low discrimination**						**046440**
					NOV		
Staphylococcus cohnii subsp. cohnii		80.64	0.39	NIT 7	99		
Staphylococcus auricularis		19.36	0.22	PHS 9	1		
046441	**Good identification**						**046441**
					NOV		
Staphylococcus cohnii subsp. cohnii		97.60	0.39	NIT 7	99		
Staphylococcus auricularis		2.40	0.03	PHS 9	1		
046450	**Low discrimination**						**046450**
					PYR	YEP	
Staphylococcus cohnii subsp. cohnii		51.22	0.41	NIT 7	1	12	
Staphylococcus kloosii		46.11	0.52	PHS 9	52	99	
Staphylococcus saprophyticus subsp. bovis		1.76	0.06		99	80	
046451	**Acceptable identification**						**046451**
Staphylococcus cohnii subsp. cohnii		92.92	0.41	NIT 7			
Staphylococcus cohnii subsp. urealyticum		7.08	0.19	PHS 9			
046452	**Very good identification**						**046452**
Staphylococcus kloosii		100.0	0.53	NIT 1			
046453	**Good identification**						**046453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	URE 99			
046470	**Good identification**						**046470**
Staphylococcus kloosii		100.0	0.40	NIT 1			
046472	**Good identification**						**046472**
Staphylococcus kloosii		100.0	0.41	NIT 1			
046503	**Good identification**						**046503**
Staphylococcus hyicus		100.0	0.29	ARG 96			
046510	**Low discrimination**						**046510**
					YEP		
Staphylococcus kloosii		72.39	0.32	NIT 1	99		
Staphylococcus cohnii subsp. cohnii		27.61	0.11	MLT 95	12		
046512	**Good identification**						**046512**
Staphylococcus kloosii		100.0	0.33	NIT 1			
046550	**Low discrimination**						**046550**
					PYR	YEP	
Staphylococcus kloosii		87.55	0.58	NIT 1	52	99	
Staphylococcus saprophyticus subsp. bovis		6.88	0.18		99	80	
Staphylococcus cohnii subsp. cohnii		5.57	0.21		1	12	
046552	**Very good identification**						**046552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		98.22	0.59	NIT 1			
Staphylococcus saprophyticus subsp. bovis		1.78	0.06				
046553	**Acceptable identification**						**046553**
Staphylococcus cohnii subsp. urealyticum		94.93	0.32	URE 99			
Staphylococcus kloosii		5.08	0.18				
046570	**Good identification**						**046570**
Staphylococcus kloosii		100.0	0.46	NIT 1			
046572	**Acceptable identification**						**046572**
Staphylococcus kloosii		90.04	0.47	NIT 1			
Staphylococcus arlettae		9.96	0.15				
046600	**Good identification**						**046600**
Staphylococcus muscae		99.24	0.39	GLR 10			
Staphylococcus auricularis		0.76	0.14	XYL 99			
046603	**Good identification**						**046603**
Staphylococcus hyicus		99.75	0.48	ARG 96			
Staphylococcus simulans		0.25	0.02	GAL 92			
046620	**Very good identification**						**046620**
Staphylococcus muscae		100.0	0.80	GLR 10			
046621	**Good identification**						**046621**
Staphylococcus muscae		100.0	0.39	GLR 10			
046622	**Good identification**						**046622**
Staphylococcus muscae		100.0	0.39	GLR 10			
046630	**Good identification**						**046630**
					OXI	NOV	
Staphylococcus muscae		99.08	0.39	GLR 10	1	1	
Staphylococcus vitulinus		0.67	0.04	MAN 1	99	99	
Staphylococcus sciuri		0.26	0.04		99	99	
046651	**Good identification**						**046651**
					OXI	PYR	
Staphylococcus sciuri		96.74	0.37	GLR 14	99	1	
Staphylococcus saprophyticus subsp. bovis		3.26	0.06	ESL 99	1	99	
046652	**Low discrimination**						**046652**
					OXI	PYR	
Staphylococcus kloosii		63.75	0.32	NIT 1	1	52	
Staphylococcus saprophyticus subsp. bovis		22.70	0.06	SUC 9	1	99	
Staphylococcus sciuri		13.55	0.02		99	1	
046660	**Good identification**						**046660**
Staphylococcus muscae		100.0	0.39	GLR 10			
046671	**Acceptable identification**						**046671**
					OXI		
Staphylococcus sciuri		94.08	0.34	GLR 14	99		
Staphylococcus xylosus		5.92	0.05	ESL 99	1		
046672	**Good identification**						**046672**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		97.43	0.41	NIT 1				
Staphylococcus kloosii		2.57	0.20					
046673	**Low discrimination**							**046673**
					OXI			
Staphylococcus arlettae		80.15	0.43	NIT 1	1			
Staphylococcus xylosus		9.11	0.15		1			
Staphylococcus sciuri		8.28	0.18		99			
046701	**Good identification**							**046701**
					OXI	NOV		
Staphylococcus hyicus		99.81	0.48	ARG 96	1	1		
Staphylococcus sciuri		0.19	0.00	LAC 92	99	99		
046702	**Good identification**							**046702**
Staphylococcus hyicus		100.0	0.29	ARG 96				
046703	**Very good identification**							**046703**
					YEP			
Staphylococcus hyicus		99.98	0.70	ARG 96	1			
Staphylococcus chromogenes		0.02	-.02		95			
046711	**Low discrimination**							**046711**
					OXI	NOV		
Staphylococcus sciuri		85.84	0.31	GLR 14	99	99		
Staphylococcus hyicus		14.16	0.07	ESL 99	1	1		
046713	**Low discrimination**							**046713**
					OXI	PYR	NOV	
Staphylococcus hyicus		89.07	0.29	ARG 96	1	1	1	
Staphylococcus sciuri		7.65	0.15	MAN 1	99	1	99	
Staphylococcus intermedius		3.29	0.05		1	90	1	
046720	**Good identification**							**046720**
Staphylococcus muscae		100.0	0.39	GLR 10				
046723	**Good identification**							**046723**
Staphylococcus hyicus		100.0	0.29	ARG 96				
046731	**Acceptable identification**							**046731**
					OXI			
Staphylococcus sciuri		91.50	0.28	GLR 14	99			
Staphylococcus xylosus		8.50	0.03	ESL 99	1			
046743	**Good identification**							**046743**
					YEP	PYR	COA	
Staphylococcus hyicus		95.98	0.29	ARG 96	1	1	24	
Staphylococcus chromogenes		2.05	-.02	MLT 1	95	45	1	
Staphylococcus intermedius		1.97	-.01		1	90	99	
046750	**Low discrimination**							**046750**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		55.48	0.31	URE 99	1	99		
Staphylococcus sciuri		24.69	0.24		99	1		
Staphylococcus kloosii		17.45	0.38		1	52		
046751	**Acceptable identification**							**046751**
					OXI	PYR		
Staphylococcus sciuri		91.66	0.43	GLR 14	99	1		
Staphylococcus saprophyticus subsp. bovis		6.37	0.18	ESL 99	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		1.97	0.04		1 65	
046752	**Low discrimination**					**046752**
					PYR	
Staphylococcus kloosii		44.93	0.38	NIT 1	52	
Staphylococcus saprophyticus subsp. bovis		32.96	0.18	SUC 9	99	
Staphylococcus arlettae		12.56	0.15		1	
046753	**Low discrimination**					**046753**
					OXI	
Staphylococcus sciuri		49.73	0.27	GLR 14	99	
Staphylococcus xylosus		18.71	0.13	ESL 99	1	
Staphylococcus arlettae		10.71	0.18	LAC 14	1	
046771	**Low discrimination**					**046771**
					OXI	
Staphylococcus sciuri		50.80	0.40	GLR 14	99	
Staphylococcus gallinarum		27.91	0.29	ESL 99	1	
Staphylococcus xylosus		16.66	0.26		1	
046772	**Very good identification**					**046772**
					OXI	
Staphylococcus arlettae		98.16	0.56	NIT 1	1	
Staphylococcus kloosii		0.89	0.26		1	
Staphylococcus sciuri		0.57	0.05		99	
046773	**Low discrimination**					**046773**
Staphylococcus arlettae		64.14	0.59	NIT 1		
Staphylococcus gallinarum		18.75	0.37			
Staphylococcus xylosus		13.01	0.35			
047211	**Good identification**					**047211**
Staphylococcus sciuri		100.0	0.35	GLR 14		
047231	**Good identification**					**047231**
Staphylococcus sciuri		100.0	0.32	GLR 14		
047250	**Good identification**					**047250**
Staphylococcus sciuri		100.0	0.28	GLR 14		
047251	**Good identification**					**047251**
Staphylococcus sciuri		100.0	0.47	GLR 14		
047253	**Good identification**					**047253**
Staphylococcus sciuri		100.0	0.31	GLR 14		
047271	**Good identification**					**047271**
Staphylococcus sciuri		100.0	0.44	GLR 14		
047273	**Acceptable identification**					**047273**
					OXI	
Staphylococcus sciuri		91.16	0.28	GLR 14	99	
Staphylococcus gallinarum		4.82	-.04	TRE 97	1	
Staphylococcus equorum		4.02	-.03	LAC 14	1	
047311	**Good identification**					**047311**
Staphylococcus sciuri		100.0	0.41	GLR 14		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
047331	**Good identification**					**047331**
Staphylococcus sciuri		100.0	0.38	GLR 14		
047350	**Good identification**					**047350**
Staphylococcus sciuri		100.0	0.34	GLR 14		
047351	**Good identification**					**047351**
Staphylococcus sciuri		100.0	0.53	GLR 14		
047353	**Good identification**					**047353**
					OXI	
Staphylococcus sciuri		95.59	0.37	GLR 14	99	
Staphylococcus lentus		2.44	0.05	TRE 97	99	
Staphylococcus gallinarum		1.97	-.04	LAC 14	1	
047370	**Good identification**					**047370**
Staphylococcus sciuri		100.0	0.32	GLR 14		
047371	**Low discrimination**					**047371**
					OXI	
Staphylococcus sciuri		83.76	0.50	GLR 14	99	
Staphylococcus gallinarum		15.03	0.29	TRE 97	1	
Staphylococcus equorum		1.22	0.09		1	
047373	**Low discrimination**					**047373**
					OXI	
Staphylococcus gallinarum		66.55	0.37	URE 92	1	
Staphylococcus sciuri		24.66	0.34	TRE 99	99	
Staphylococcus equorum		5.66	0.18		1	
047412	**Good identification**					**047412**
Staphylococcus kloosii		100.0	0.27	NIT 1		
047450	**Low discrimination**					**047450**
					OXI	
Staphylococcus kloosii		87.73	0.52	NIT 1	1	
Staphylococcus sciuri		12.27	0.18		99	
047451	**Good identification**					**047451**
					OXI	
Staphylococcus sciuri		96.95	0.37	GLR 14	99	
Staphylococcus cohnii subsp. urealyticum		3.05	0.11	SUC 99	1	
047452	**Very good identification**					**047452**
					OXI	
Staphylococcus kloosii		97.94	0.53	NIT 1	1	
Staphylococcus sciuri		2.06	0.02		99	
047453	**Low discrimination**					**047453**
					OXI	
Staphylococcus cohnii subsp. urealyticum		66.17	0.31	URE 99	1	
Staphylococcus sciuri		33.83	0.21		99	
047470	**Low discrimination**					**047470**
					OXI	
Staphylococcus kloosii		70.31	0.40	NIT 1	1	
Staphylococcus sciuri		29.69	0.16		99	
047471	**Good identification**					**047471**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus sciuri		100.0	0.34	GLR 14		
047472	**Good identification**					**047472**
Staphylococcus kloosii		100.0	0.41	NIT 1		
047510	**Low discrimination**					**047510**
					OXI	
Staphylococcus kloosii		58.59	0.32	NIT 1	1	
Staphylococcus sciuri		41.41	0.12	MLT 95	99	
047511	**Good identification**					**047511**
Staphylococcus sciuri		100.0	0.31	GLR 14		
047512	**Good identification**					**047512**
Staphylococcus kloosii		100.0	0.33	NIT 1		
047531	**Good identification**					**047531**
Staphylococcus sciuri		100.0	0.28	GLR 14		
047550	**Low discrimination**					**047550**
					OXI	
Staphylococcus kloosii		87.73	0.58	NIT 1	1	
Staphylococcus sciuri		12.27	0.24		99	
047551	**Good identification**					**047551**
					OXI	
Staphylococcus sciuri		99.12	0.43	GLR 14	99	
Staphylococcus kloosii		0.88	0.17	SUC 99	1	
047552	**Very good identification**					**047552**
					OXI	
Staphylococcus kloosii		97.94	0.59	NIT 1	1	
Staphylococcus sciuri		2.06	0.08		99	
047553	**Low discrimination**					**047553**
					OXI	
Staphylococcus sciuri		61.25	0.27	GLR 14	99	
Staphylococcus cohnii subsp. urealyticum		26.46	0.24	SUC 99	1	
Staphylococcus lentus		4.80	0.05	LAC 14	99	
047570	**Low discrimination**					**047570**
					OXI	
Staphylococcus kloosii		70.31	0.46	NIT 1	1	
Staphylococcus sciuri		29.69	0.22		99	
047571	**Low discrimination**					**047571**
					OXI	
Staphylococcus sciuri		62.74	0.40	GLR 14	99	
Staphylococcus gallinarum		34.47	0.29	SUC 99	1	
Staphylococcus equorum		2.79	0.09		1	
047572	**Low discrimination**					**047572**
					OXI	
Staphylococcus kloosii		82.18	0.47	NIT 1	1	
Staphylococcus arlettae		9.09	0.15		1	
Staphylococcus sciuri		5.21	0.05		99	
047573	**Low discrimination**					**047573**
					OXI	
Staphylococcus gallinarum		79.22	0.37	URE 92	1	
Staphylococcus sciuri		9.59	0.24	SUC 99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus equorum		6.74	0.18		1		
047601	**Good identification**						**047601**
Staphylococcus sciuri		100.0	0.35	GLR 14			
047610	**Good identification**						**047610**
Staphylococcus sciuri		98.06	0.47	GLR 14			
Staphylococcus vitulinus		1.94	0.04	MNS 89			
047611	**Very good identification**						**047611**
Staphylococcus sciuri		100.0	0.66	GLR 14			
047612	**Good identification**						**047612**
Staphylococcus sciuri		100.0	0.31	GLR 14			
047613	**Good identification**						**047613**
Staphylococcus sciuri		100.0	0.50	GLR 14			
047620	**Good identification**						**047620**
					OXI	NOV	
Staphylococcus muscae		99.22	0.39	GLR 10	1	1	
Staphylococcus sciuri		0.79	0.14	ESL 1	99	99	
047621	**Low discrimination**						**047621**
					OXI	NOV	
Staphylococcus sciuri		86.38	0.32	GLR 14	99	99	
Staphylococcus muscae		13.62	-.02	MAN 97	1	1	
047630	**Acceptable identification**						**047630**
					OXI	NOV	
Staphylococcus sciuri		90.05	0.45	GLR 14	99	99	
Staphylococcus vitulinus		6.39	0.13	MNS 89	99	99	
Staphylococcus muscae		3.55	-.02		1	1	
047631	**Very good identification**						**047631**
					OXI		
Staphylococcus sciuri		99.68	0.64	GLR 14	99		
Staphylococcus equorum		0.32	0.09		1		
047632	**Good identification**						**047632**
Staphylococcus sciuri		100.0	0.29	GLR 14			
047633	**Acceptable identification**						**047633**
					OXI		
Staphylococcus sciuri		94.67	0.47	GLR 14	99		
Staphylococcus equorum		4.75	0.18	LAC 14	1		
Staphylococcus gallinarum		0.58	-.04		1		
047640	**Good identification**						**047640**
Staphylococcus sciuri		100.0	0.28	GLR 14			
047641	**Good identification**						**047641**
Staphylococcus sciuri		100.0	0.47	GLR 14			
047643	**Good identification**						**047643**
Staphylococcus sciuri		100.0	0.31	GLR 14			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
047650	**Very good identification**						**047650**
					OXI		
Staphylococcus sciuri		99.29	0.59	GLR 14	99		
Staphylococcus kloosii		0.71	0.32	MNS 89	1		
047651	**Very good identification**						**047651**
Staphylococcus sciuri		100.0	0.78	GLR 14			
047652	**Good identification**						**047652**
					OXI		
Staphylococcus sciuri		95.46	0.43	GLR 14	99		
Staphylococcus kloosii		4.54	0.32	MNS 89	1		
047653	**Very good identification**						**047653**
					OXI		
Staphylococcus sciuri		99.62	0.62	GLR 14	99		
Staphylococcus lentus		0.15	0.05	LAC 14	99		
Staphylococcus gallinarum		0.12	-.04		1		
047661	**Good identification**						**047661**
					OXI		
Staphylococcus sciuri		96.67	0.44	GLR 14	99		
Staphylococcus gallinarum		3.33	0.08	MAN 97	1		
047663	**Low discrimination**						**047663**
					OXI		
Staphylococcus sciuri		64.01	0.28	GLR 14	99		
Staphylococcus gallinarum		33.17	0.16	MAN 97	1		
Staphylococcus equorum		2.82	-.03	LAC 14	1		
047670	**Very good identification**						**047670**
					OXI		
Staphylococcus sciuri		99.76	0.57	GLR 14	99		
Staphylococcus kloosii		0.24	0.19	MNS 89	1		
047671	**Very good identification**						**047671**
					OXI		
Staphylococcus sciuri		98.11	0.75	GLR 14	99		
Staphylococcus gallinarum		1.06	0.29		1		
Staphylococcus equorum		0.84	0.30		1		
047672	**Low discrimination**						**047672**
					OXI		
Staphylococcus sciuri		61.09	0.40	GLR 14	99		
Staphylococcus arlettae		36.47	0.41	MNS 89	1		
Staphylococcus kloosii		0.96	0.20	LAC 14	1		
047673	**Low discrimination**						**047673**
					OXI		
Staphylococcus sciuri		71.45	0.59	GLR 14	99		
Staphylococcus gallinarum		11.58	0.37	LAC 14	1		
Staphylococcus equorum		9.64	0.38		1		
047701	**Acceptable identification**						**047701**
					OXI	NOV	
Staphylococcus sciuri		94.89	0.41	GLR 14	99	99	
Staphylococcus hyicus		5.11	0.07	MAN 97	1	1	
047703	**Low discrimination**						**047703**
					OXI	NOV	
Staphylococcus hyicus		79.19	0.29	ARG 96	1	1	
Staphylococcus sciuri		20.81	0.25	ESL 1	99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
047710	**Very good identification**						**047710**
Staphylococcus sciuri		100.0	0.53	GLR 14			
047711	**Very good identification**						**047711**
Staphylococcus sciuri		99.34	0.72	GLR 14			
Staphylococcus lentus		0.66	0.29				
047712	**Good identification**						**047712**
Staphylococcus sciuri		100.0	0.37	GLR 14			
047713	**Low discrimination**						**047713**
					OXI		
Staphylococcus sciuri		87.00	0.56	GLR 14	99		
Staphylococcus lentus		12.62	0.40	LAC 14	99		
Staphylococcus gallinarum		0.21	-.04		1		
047721	**Acceptable identification**						**047721**
					OXI		
Staphylococcus sciuri		93.74	0.38	GLR 14	99		
Staphylococcus gallinarum		6.26	0.08	MAN 97	1		
047730	**Good identification**						**047730**
Staphylococcus sciuri		100.0	0.51	GLR 14			
047731	**Very good identification**						**047731**
					OXI		
Staphylococcus sciuri		95.58	0.69	GLR 14	99		
Staphylococcus gallinarum		2.00	0.29		1		
Staphylococcus equorum		1.58	0.30		1		
047732	**Low discrimination**						**047732**
					OXI		
Staphylococcus sciuri		89.83	0.35	GLR 14	99		
Staphylococcus arlettae		5.95	0.15	MNS 89	1		
Staphylococcus gallinarum		2.31	-.04	LAC 14	1		
047733	**Low discrimination**						**047733**
					OXI		
Staphylococcus sciuri		56.09	0.53	GLR 14	99		
Staphylococcus gallinarum		17.61	0.37	LAC 14	1		
Staphylococcus equorum		14.67	0.38		1		
047740	**Good identification**						**047740**
Staphylococcus sciuri		100.0	0.34	GLR 14			
047741	**Good identification**						**047741**
					OXI	NOV	
Staphylococcus sciuri		98.37	0.53	GLR 14	99	99	
Staphylococcus gallinarum		1.32	0.08	MAN 97	1	99	
Rothia mucilaginosa		0.32	-.09		1	1	
047743	**Low discrimination**						**047743**
					OXI		
Staphylococcus sciuri		81.51	0.37	GLR 14	99		
Staphylococcus gallinarum		16.41	0.16	MAN 97	1		
Staphylococcus lentus		2.08	0.05	LAC 14	99		
047750	**Very good identification**						**047750**
					OXI		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus sciuri		99.29	0.65	GLR 14	99	
Staphylococcus kloosii		0.71	0.38	MNS 89	1	
047751	**Very good identification**					**047751**
					OXI	
Staphylococcus sciuri		99.20	0.84	GLR 14	99	
Staphylococcus gallinarum		0.42	0.29		1	
Staphylococcus lentus		0.36	0.35		99	
047752	**Acceptable identification**					**047752**
					OXI	
Staphylococcus sciuri		93.25	0.49	GLR 14	99	
Staphylococcus kloosii		4.43	0.38	MNS 89	1	
Staphylococcus arlettae		1.24	0.15	LAC 14	1	
047753	**Low discrimination**					**047753**
					OXI	
Staphylococcus sciuri		86.98	0.68	GLR 14	99	
Staphylococcus lentus		6.81	0.47	LAC 14	99	
Staphylococcus gallinarum		5.48	0.37		1	
047760	**Low discrimination**					**047760**
					OXI	
Staphylococcus sciuri		87.44	0.32	GLR 14	99	
Staphylococcus gallinarum		12.56	0.08	MAN 97	1	
047761	**Low discrimination**					**047761**
					OXI	
Staphylococcus gallinarum		63.39	0.50	URE 92	1	
Staphylococcus sciuri		36.08	0.50	MAN 91	99	
Staphylococcus equorum		0.52	0.09		1	
047763	**Very good identification**					**047763**
					OXI	
Staphylococcus gallinarum		95.01	0.58	URE 92	1	
Staphylococcus sciuri		3.60	0.34	MAN 91	99	
Staphylococcus equorum		0.83	0.18		1	
047770	**Good identification**					**047770**
					OXI	
Staphylococcus sciuri		94.90	0.63	GLR 14	99	
Staphylococcus gallinarum		4.26	0.29	MNS 89	1	
Staphylococcus equorum		0.35	0.09		1	
047771	**Low discrimination**					**047771**
					OXI	
Staphylococcus sciuri		62.50	0.81	GLR 14	99	
Staphylococcus gallinarum		34.34	0.70		1	
Staphylococcus equorum		2.78	0.51		1	
047772	**Low discrimination**					**047772**
					OXI	
Staphylococcus arlettae		49.80	0.56	NIT 1	1	
Staphylococcus sciuri		28.57	0.46		99	
Staphylococcus gallinarum		19.29	0.37		1	
047773	**Low discrimination**					**047773**
					OXI	
Staphylococcus gallinarum		79.68	0.78	URE 92	1	
Staphylococcus sciuri		9.64	0.65		99	
Staphylococcus equorum		6.78	0.59		1	
050000	**Low discrimination**					**050000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		62.04	0.33	ORN 1			
Micrococcus sp.		28.01	0.18	NIT 90			
Dermacoccus nishinomiyaensis		9.95	0.18				
050400	**Good identification**						**050400**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	ORN 1			
050401	**Good identification**						**050401**
Staphylococcus cohnii subsp. cohnii		100.0	0.32	ORN 1			
050410	**Low discrimination**						**050410**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		86.22	0.33	ORN 1	1	12	
Macrococcus bovicus		13.78	0.10		99	99	
050411	**Good identification**						**050411**
Staphylococcus cohnii subsp. cohnii		100.0	0.34	ORN 1			
050440	**Good identification**						**050440**
					NOV		
Staphylococcus cohnii subsp. cohnii		98.07	0.41	ORN 1	99		
Staphylococcus auricularis		1.93	0.03		1		
050441	**Good identification**						**050441**
Staphylococcus cohnii subsp. cohnii		100.0	0.42	ORN 1			
050450	**Low discrimination**						**050450**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		89.47	0.43	ORN 1	1	12	
Macrococcus bovicus		10.53	0.18		99	99	
050451	**Good identification**						**050451**
Staphylococcus cohnii subsp. cohnii		100.0	0.44	ORN 1			
050672	**Good identification**						**050672**
					NOV	YEP	
Staphylococcus arlettae		97.27	0.37	ORN 1	99	99	
Aerococcus viridans		2.74	0.16		1	1	
050673	**Good identification**						**050673**
					NOV	YEP	
Staphylococcus arlettae		95.28	0.40	ORN 1	99	99	
Aerococcus viridans		4.72	0.24		1	1	
050772	**Good identification**						**050772**
					NOV	YEP	
Staphylococcus arlettae		99.51	0.53	ORN 1	99	99	
Aerococcus viridans		0.49	0.16		1	1	
050773	**Very good identification**						**050773**
					NOV	YEP	
Staphylococcus arlettae		99.13	0.55	ORN 1	99	99	
Aerococcus viridans		0.87	0.24		1	1	
051413	**Good identification**						**051413**
Aerococcus viridans		100.0	0.31	ORN 1			
051433	**Good identification**						**051433**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.31	ORN 1			
051443	**Good identification**						**051443**
Aerococcus viridans		100.0	0.34	ORN 1			
051452	**Good identification**						**051452**
Aerococcus viridans		100.0	0.36	ORN 1			
051453	**Good identification**						**051453**
Aerococcus viridans		100.0	0.43	ORN 1			
051463	**Good identification**						**051463**
Aerococcus viridans		100.0	0.34	ORN 1			
051472	**Good identification**						**051472**
Aerococcus viridans		100.0	0.36	ORN 1			
051473	**Good identification**						**051473**
Aerococcus viridans		100.0	0.43	ORN 1			
051513	**Good identification**						**051513**
Aerococcus viridans		100.0	0.31	ORN 1			
051533	**Good identification**						**051533**
Aerococcus viridans		100.0	0.31	ORN 1			
051543	**Good identification**						**051543**
Aerococcus viridans		100.0	0.34	ORN 1			
051552	**Low discrimination**						**051552**
					NOV	YEP	
Aerococcus viridans		88.97	0.36	ORN 1	1	1	
Staphylococcus kloosii		11.03	0.18		99	99	
051553	**Good identification**						**051553**
Aerococcus viridans		100.0	0.43	ORN 1			
051563	**Good identification**						**051563**
Aerococcus viridans		100.0	0.34	ORN 1			
051572	**Low discrimination**						**051572**
					NOV	YEP	
Aerococcus viridans		81.56	0.36	ORN 1	1	1	
Staphylococcus arlettae		18.44	0.12		99	99	
051573	**Low discrimination**						**051573**
					NOV	YEP	
Aerococcus viridans		88.62	0.43	ORN 1	1	1	
Staphylococcus arlettae		11.38	0.14		99	99	
051613	**Good identification**						**051613**
Aerococcus viridans		100.0	0.31	ORN 1			
051633	**Good identification**						**051633**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.31	ORN 1			
051643	**Good identification**						**051643**
Aerococcus viridans		100.0	0.34	ORN 1			
051652	**Good identification**						**051652**
Aerococcus viridans		100.0	0.36	ORN 1			
051653	**Good identification**						**051653**
Aerococcus viridans		100.0	0.43	ORN 1			
051663	**Good identification**						**051663**
Aerococcus viridans		100.0	0.34	ORN 1			
051672	**Low discrimination**						**051672**
					NOV	YEP	
Staphylococcus arlettae		79.80	0.37	ORN 1	99	99	
Aerococcus viridans		20.20	0.36		1	1	
051673	**Low discrimination**						**051673**
					NOV	YEP	
Staphylococcus arlettae		69.18	0.40	ORN 1	99	99	
Aerococcus viridans		30.82	0.43		1	1	
051713	**Good identification**						**051713**
Aerococcus viridans		100.0	0.31	ORN 1			
051733	**Low discrimination**						**051733**
					NOV	YEP	
Aerococcus viridans		66.06	0.31	ORN 1	1	1	
Staphylococcus arlettae		33.94	0.14		99	99	
051743	**Good identification**						**051743**
Aerococcus viridans		100.0	0.34	ORN 1			
051752	**Low discrimination**						**051752**
					NOV	YEP	
Aerococcus viridans		81.56	0.36	ORN 1	1	1	
Staphylococcus arlettae		18.44	0.12		99	99	
051753	**Low discrimination**						**051753**
					NOV	YEP	
Aerococcus viridans		88.62	0.43	ORN 1	1	1	
Staphylococcus arlettae		11.38	0.14		99	99	
051763	**Low discrimination**						**051763**
					NOV	YEP	
Aerococcus viridans		72.18	0.34	ORN 1	1	1	
Staphylococcus arlettae		27.82	0.14		99	99	
051772	**Good identification**						**051772**
					NOV	YEP	
Staphylococcus arlettae		95.72	0.53	ORN 1	99	99	
Aerococcus viridans		4.28	0.36		1	1	
051773	**Good identification**						**051773**
					NOV	YEP	
Staphylococcus arlettae		92.71	0.55	ORN 1	99	99	
Aerococcus viridans		7.29	0.43		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
052000	**Good identification**					**052000**
Kocuria rosea		98.63	0.53	ORN 1		
Dermacoccus nishinomiyaensis		0.99	0.13			
Micrococcus sp.		0.26	-.09			
052010	**Good identification**					**052010**
Kocuria rosea		100.0	0.33	ORN 1		
052020	**Good identification**					**052020**
Kocuria rosea		100.0	0.45	ORN 1		
052200	**Good identification**					**052200**
Kocuria rosea		100.0	0.33	ORN 1		
053651	**Good identification**					**053651**
Staphylococcus sciuri		100.0	0.32	ORN 1		
053671	**Good identification**					**053671**
Staphylococcus sciuri		100.0	0.30	ORN 1		
053751	**Good identification**					**053751**
Staphylococcus sciuri		97.61	0.38	ORN 1		
Staphylococcus lentus		2.39	0.07	GLR 14		
053771	**Acceptable identification**					**053771**
					OXI	
Staphylococcus sciuri		92.34	0.35	ORN 1	99	
Staphylococcus equorum		4.66	0.06	GLR 14	1	
Staphylococcus lentus		3.00	0.07		99	
054412	**Good identification**					**054412**
Staphylococcus kloosii		100.0	0.27	ORN 1		
054450	**Low discrimination**					**054450**
					YEP	
Staphylococcus kloosii		87.03	0.52	ORN 1	99	
Staphylococcus cohnii subsp. cohnii		12.97	0.23		12	
054452	**Very good identification**					**054452**
Staphylococcus kloosii		100.0	0.53	ORN 1		
054470	**Good identification**					**054470**
Staphylococcus kloosii		100.0	0.40	ORN 1		
054472	**Good identification**					**054472**
Staphylococcus kloosii		100.0	0.41	ORN 1		
054510	**Good identification**					**054510**
Staphylococcus kloosii		100.0	0.32	ORN 1		
054512	**Good identification**					**054512**
Staphylococcus kloosii		100.0	0.33	ORN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
054550	**Very good identification**					**054550**
Staphylococcus kloosii		100.0	0.58	ORN 1		
054552	**Very good identification**					**054552**
Staphylococcus kloosii		100.0	0.59	ORN 1		
054570	**Good identification**					**054570**
Staphylococcus kloosii		100.0	0.46	ORN 1		
054572	**Acceptable identification**					**054572**
Staphylococcus kloosii		90.04	0.47	ORN 1		
Staphylococcus arlettae		9.96	0.15			
054650	**Good identification**					**054650**
Staphylococcus kloosii		100.0	0.32	ORN 1		
054652	**Good identification**					**054652**
Staphylococcus kloosii		100.0	0.32	ORN 1		
054672	**Good identification**					**054672**
Staphylococcus arlettae		97.43	0.41	ORN 1		
Staphylococcus kloosii		2.57	0.20			
054673	**Good identification**					**054673**
Staphylococcus arlettae		100.0	0.43	ORN 1		
054750	**Good identification**					**054750**
Staphylococcus kloosii		100.0	0.38	ORN 1		
054752	**Low discrimination**					**054752**
Staphylococcus kloosii		78.16	0.38	ORN 1		
Staphylococcus arlettae		21.85	0.15	SUC 9		
054772	**Very good identification**					**054772**
Staphylococcus arlettae		99.11	0.56	ORN 1		
Staphylococcus kloosii		0.90	0.26			
054773	**Very good identification**					**054773**
Staphylococcus arlettae		100.0	0.59	ORN 1		
055412	**Good identification**					**055412**
Staphylococcus kloosii		100.0	0.27	ORN 1		
055450	**Good identification**					**055450**
Staphylococcus kloosii		100.0	0.52	ORN 1		
055452	**Very good identification**					**055452**
Staphylococcus kloosii		100.0	0.53	ORN 1		
055470	**Good identification**					**055470**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.40	ORN 1		
055472	**Good identification**					**055472**
Staphylococcus kloosii		100.0	0.41	ORN 1		
055510	**Good identification**					**055510**
Staphylococcus kloosii		100.0	0.32	ORN 1		
055512	**Good identification**					**055512**
Staphylococcus kloosii		100.0	0.33	ORN 1		
055550	**Very good identification**					**055550**
Staphylococcus kloosii		100.0	0.58	ORN 1		
055552	**Very good identification**					**055552**
Staphylococcus kloosii		100.0	0.59	ORN 1		
055570	**Good identification**					**055570**
Staphylococcus kloosii		100.0	0.46	ORN 1		
055572	**Acceptable identification**					**055572**
Staphylococcus kloosii		90.04	0.47	ORN 1		
Staphylococcus arlettae		9.96	0.15			
055650	**Good identification**					**055650**
Staphylococcus kloosii		100.0	0.32	ORN 1		
055652	**Good identification**					**055652**
Staphylococcus kloosii		100.0	0.32	ORN 1		
055672	**Good identification**					**055672**
Staphylococcus arlettae		97.43	0.41	ORN 1		
Staphylococcus kloosii		2.57	0.20			
055673	**Good identification**					**055673**
Staphylococcus arlettae		100.0	0.43	ORN 1		
055750	**Good identification**					**055750**
Staphylococcus kloosii		100.0	0.38	ORN 1		
055752	**Low discrimination**					**055752**
Staphylococcus kloosii		78.16	0.38	ORN 1		
Staphylococcus arlettae		21.85	0.15	SUC 9		
055772	**Very good identification**					**055772**
Staphylococcus arlettae		99.11	0.56	ORN 1		
Staphylococcus kloosii		0.90	0.26			
055773	**Very good identification**					**055773**
Staphylococcus arlettae		99.71	0.59	ORN 1		
Staphylococcus gallinarum		0.29	-.04			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
056620	**Good identification**					**056620**
Staphylococcus muscae		100.0	0.39	ORN 1		
056703	**Good identification**					**056703**
Staphylococcus hyicus		100.0	0.29	ARG 96		
057651	**Good identification**					**057651**
Staphylococcus sciuri		100.0	0.37	ORN 1		
057671	**Good identification**					**057671**
Staphylococcus sciuri		100.0	0.34	ORN 1		
057711	**Good identification**					**057711**
Staphylococcus sciuri		100.0	0.31	ORN 1		
057731	**Good identification**					**057731**
Staphylococcus sciuri		100.0	0.28	ORN 1		
057751	**Good identification**					**057751**
Staphylococcus sciuri		100.0	0.43	ORN 1		
057753	**Low discrimination**					**057753**
					OXI	
Staphylococcus sciuri		87.62	0.27	ORN 1	99	
Staphylococcus lentus		6.86	0.05	GLR 14	99	
Staphylococcus gallinarum		5.52	-.04	LAC 14	1	
057771	**Low discrimination**					**057771**
					OXI	
Staphylococcus sciuri		62.74	0.40	ORN 1	99	
Staphylococcus gallinarum		34.47	0.29	GLR 14	1	
Staphylococcus equorum		2.79	0.09		1	
057773	**Low discrimination**					**057773**
					OXI	
Staphylococcus gallinarum		79.80	0.37	URE 92	1	
Staphylococcus sciuri		9.66	0.24	ORN 1	99	
Staphylococcus equorum		6.79	0.18		1	
060000	**Low discrimination**					**060000**
Kocuria rosea		59.24	0.33	BGA 1		
Micrococcus sp.		26.74	0.18	NIT 90		
Dermacoccus nishinomiyaensis		9.50	0.18			
060043	**Good identification**					**060043**
Aerococcus viridans		100.0	0.27	ESL 90		
060052	**Good identification**					**060052**
Aerococcus viridans		100.0	0.29	ESL 90		
060053	**Low discrimination**					**060053**
					NOV	
Aerococcus viridans		78.12	0.37	ESL 90	1	
Staphylococcus cohnii subsp. urealyticum		21.88	0.14	TRE 95	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
060063	**Good identification**						**060063**
Aerococcus viridans		100.0	0.27	ESL 90			
060072	**Good identification**						**060072**
Aerococcus viridans		100.0	0.29	ESL 90			
060073	**Good identification**						**060073**
Aerococcus viridans		100.0	0.37	ESL 90			
060143	**Good identification**						**060143**
Aerococcus viridans		100.0	0.27	ESL 90			
060152	**Low discrimination**						**060152**
					NOV	YEP	
Aerococcus viridans		79.26	0.29	ESL 90	1	1	
Staphylococcus kloosii		20.74	0.18	TRE 95	99	99	
060153	**Low discrimination**						**060153**
					NOV		
Aerococcus viridans		89.28	0.37	ESL 90	1		
Staphylococcus cohnii subsp. urealyticum		10.72	0.07	TRE 95	99		
060163	**Good identification**						**060163**
Aerococcus viridans		100.0	0.27	ESL 90			
060172	**Low discrimination**						**060172**
					NOV	YEP	
Aerococcus viridans		67.69	0.29	ESL 90	1	1	
Staphylococcus arlettae		32.31	0.12	TRE 95	99	99	
060173	**Low discrimination**						**060173**
					NOV	YEP	
Aerococcus viridans		78.67	0.37	ESL 90	1	1	
Staphylococcus arlettae		21.34	0.14	TRE 95	99	99	
060243	**Good identification**						**060243**
Aerococcus viridans		100.0	0.27	ESL 90			
060252	**Good identification**						**060252**
Aerococcus viridans		100.0	0.29	ESL 90			
060253	**Good identification**						**060253**
Aerococcus viridans		100.0	0.37	ESL 90			
060263	**Good identification**						**060263**
Aerococcus viridans		100.0	0.27	ESL 90			
060272	**Low discrimination**						**060272**
					NOV	YEP	
Staphylococcus arlettae		89.29	0.37	TRE 99	99	99	
Aerococcus viridans		10.71	0.29		1	1	
060273	**Low discrimination**						**060273**
					NOV	YEP	
Staphylococcus arlettae		82.57	0.40	TRE 99	99	99	
Aerococcus viridans		17.43	0.37		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
060343	**Good identification**						**060343**
Aerococcus viridans		100.0	0.27	ESL 90			
060352	**Low discrimination**						**060352**
					NOV	YEP	
Aerococcus viridans		67.69	0.29	ESL 90	1	1	
Staphylococcus arlettae		32.31	0.12	TRE 95	99	99	
060353	**Low discrimination**						**060353**
					NOV	YEP	
Aerococcus viridans		78.67	0.37	ESL 90	1	1	
Staphylococcus arlettae		21.34	0.14	TRE 95	99	99	
060363	**Low discrimination**						**060363**
					NOV	YEP	
Aerococcus viridans		55.14	0.27	ESL 90	1	1	
Staphylococcus arlettae		44.86	0.14	TRE 95	99	99	
060372	**Good identification**						**060372**
					NOV	YEP	
Staphylococcus arlettae		97.93	0.53	TRE 99	99	99	
Aerococcus viridans		2.07	0.29		1	1	
060373	**Good identification**						**060373**
					NOV	YEP	
Staphylococcus arlettae		94.39	0.55	TRE 99	99	99	
Aerococcus viridans		3.52	0.37		1	1	
Staphylococcus xylosus		2.10	0.12		99	40	
060400	**Low discrimination**						**060400**
					NOV		
Staphylococcus cohnii subsp. cohnii		69.94	0.31	BGA 1	99		
Staphylococcus auricularis		30.07	0.20		1		
060401	**Low discrimination**						**060401**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		88.78	0.32	BGA 1	99	1	
Aerococcus viridans		7.30	0.21		1	95	
Staphylococcus auricularis		3.92	0.00		1	40	
060402	**Low discrimination**						**060402**
Aerococcus viridans		87.80	0.33	ESL 90			
Staphylococcus auricularis		12.20	0.00				
060403	**Low discrimination**						**060403**
					NOV		
Aerococcus viridans		76.59	0.41	ESL 90	1		
Staphylococcus cohnii subsp. urealyticum		23.41	0.19		99		
060410	**Low discrimination**						**060410**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		73.51	0.33	BGA 1	1	12	
Macrococcus bovicus		11.75	0.10		99	99	
Staphylococcus kloosii		8.13	0.26		1	99	
060411	**Low discrimination**						**060411**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		72.90	0.34	BGA 1	99	1	
Aerococcus viridans		14.14	0.31		1	95	
Staphylococcus cohnii subsp. urealyticum		12.96	0.19		99	53	
060412	**Low discrimination**						**060412**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Aerococcus viridans		87.12	0.43	ESL 90	1	1	
Staphylococcus kloosii		12.88	0.27		99	99	
060413	**Low discrimination**						**060413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		50.75	0.40	URE 99	99		
Aerococcus viridans		49.25	0.51		1		
060422	**Good identification**						**060422**
Aerococcus viridans		100.0	0.33	ESL 90			
060423	**Good identification**						**060423**
Aerococcus viridans		100.0	0.41	ESL 90			
060431	**Good identification**						**060431**
Aerococcus viridans		100.0	0.31	ESL 90			
060432	**Good identification**						**060432**
Aerococcus viridans		100.0	0.43	ESL 90			
060433	**Good identification**						**060433**
Aerococcus viridans		100.0	0.51	ESL 90			
060440	**Low discrimination**						**060440**
					NOV		
Staphylococcus cohnii subsp. cohnii		70.24	0.41	BGA 1	99		
Staphylococcus auricularis		22.25	0.28		1		
Staphylococcus warneri		4.12	0.21		1		
060441	**Low discrimination**						**060441**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		85.00	0.42	BGA 1	99	1	
Aerococcus viridans		8.83	0.34		1	95	
Staphylococcus cohnii subsp. urealyticum		3.41	0.14		99	53	
060442	**Low discrimination**						**060442**
					PYR		
Aerococcus viridans		88.81	0.46	ESL 90	95		
Staphylococcus auricularis		7.20	0.08		40		
Staphylococcus warneri		4.00	0.11		8		
060443	**Low discrimination**						**060443**
					NOV		
Aerococcus viridans		69.78	0.53	ESL 90	1		
Staphylococcus cohnii subsp. urealyticum		30.22	0.35		99		
060450	**Low discrimination**						**060450**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		51.86	0.43	BGA 1	1	12	
Staphylococcus kloosii		34.41	0.52		1	99	
Macrococcus bovicus		6.11	0.18		99	99	
060451	**Low discrimination**						**060451**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		63.97	0.44	BGA 1	99	1	
Staphylococcus cohnii subsp. urealyticum		20.36	0.35		99	53	
Aerococcus viridans		15.67	0.43		1	95	
060452	**Low discrimination**						**060452**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		57.16	0.56	ESL 90	1	1		
Staphylococcus kloosii		40.15	0.53		99	99		
Staphylococcus cohnii subsp. urealyticum		1.97	0.14		99	64		
060453	**Low discrimination**							**060453**
					NOV			
Staphylococcus cohnii subsp. urealyticum		59.34	0.55	URE 99	99			
Aerococcus viridans		40.66	0.63		1			
060461	**Good identification**							**060461**
Aerococcus viridans		100.0	0.34	ESL 90				
060462	**Good identification**							**060462**
Aerococcus viridans		100.0	0.46	ESL 90				
060463	**Very good identification**							**060463**
Aerococcus viridans		100.0	0.53	ESL 90				
060470	**Low discrimination**							**060470**
					NOV	YEP		
Staphylococcus kloosii		59.33	0.40	PHS 99	99	99		
Aerococcus viridans		40.67	0.36		1	1		
060471	**Good identification**							**060471**
Aerococcus viridans		100.0	0.43	ESL 90				
060472	**Low discrimination**							**060472**
					NOV	YEP		
Aerococcus viridans		61.94	0.56	ESL 90	1	1		
Staphylococcus arlettae		27.19	0.37		99	99		
Staphylococcus kloosii		10.88	0.41		99	99		
060473	**Low discrimination**							**060473**
					NOV	YEP		
Aerococcus viridans		79.11	0.63	ESL 90	1	1		
Staphylococcus arlettae		19.73	0.40		99	99		
Staphylococcus cohnii subsp. urealyticum		1.17	0.14		99	64		
060502	**Good identification**							**060502**
Aerococcus viridans		100.0	0.33	ESL 90				
060503	**Low discrimination**							**060503**
					NOV			
Aerococcus viridans		88.42	0.41	ESL 90	1			
Staphylococcus cohnii subsp. urealyticum		11.58	0.12		99			
060510	**Low discrimination**							**060510**
					NOV	YEP	PYR	
Staphylococcus kloosii		44.33	0.32	PHS 99	99	99	52	
Staphylococcus cohnii subsp. cohnii		22.94	0.13	MLT 95	99	12	1	
Aerococcus viridans		18.59	0.23		1	1	95	
060511	**Low discrimination**							**060511**
					NOV	PYR		
Aerococcus viridans		50.86	0.31	ESL 90	1	95		
Staphylococcus cohnii subsp. cohnii		29.15	0.14	LAC 90	99	1		
Staphylococcus cohnii subsp. urealyticum		19.99	0.12		99	53		
060512	**Low discrimination**							**060512**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		77.70	0.43	ESL 90	1	1		
Staphylococcus kloosii		22.30	0.33		99	99		
060513	**Low discrimination**							**060513**
					NOV			
Aerococcus viridans		69.37	0.51	ESL 90	1			
Staphylococcus cohnii subsp. urealyticum		30.63	0.32		99			
060522	**Good identification**							**060522**
Aerococcus viridans		100.0	0.33	ESL 90				
060523	**Good identification**							**060523**
Aerococcus viridans		100.0	0.41	ESL 90				
060531	**Good identification**							**060531**
Aerococcus viridans		100.0	0.31	ESL 90				
060532	**Low discrimination**							**060532**
					NOV	YEP		
Aerococcus viridans		85.31	0.43	ESL 90	1	1		
Staphylococcus arlettae		8.57	0.12		99	99		
Staphylococcus kloosii		6.12	0.20		99	99		
060533	**Acceptable identification**							**060533**
					NOV	YEP		
Aerococcus viridans		94.60	0.51	ESL 90	1	1		
Staphylococcus arlettae		5.40	0.14		99	99		
060542	**Good identification**							**060542**
					NOV	YEP		
Aerococcus viridans		96.03	0.46	ESL 90	1	1		
Staphylococcus kloosii		3.97	0.18		99	99		
060543	**Low discrimination**							**060543**
					NOV			
Aerococcus viridans		84.35	0.53	ESL 90	1			
Staphylococcus cohnii subsp. urealyticum		15.66	0.27		99			
060550	**Low discrimination**							**060550**
					NOV	YEP	PYR	
Staphylococcus kloosii		82.39	0.58	PHS 99	99	99	52	
Aerococcus viridans		7.27	0.36		1	1	95	
Staphylococcus cohnii subsp. cohnii		7.11	0.24		99	12	1	
060551	**Low discrimination**							**060551**
					NOV	PYR		
Aerococcus viridans		48.56	0.43	ESL 90	1	95		
Staphylococcus cohnii subsp. urealyticum		27.04	0.27	LAC 90	99	53		
Staphylococcus cohnii subsp. cohnii		22.03	0.25		99	1		
060552	**Low discrimination**							**060552**
					NOV	YEP		
Staphylococcus kloosii		56.73	0.59	PHS 99	99	99		
Aerococcus viridans		41.61	0.56		1	1		
Staphylococcus arlettae		1.05	0.12		99	99		
060553	**Low discrimination**							**060553**
					NOV	YEP		
Aerococcus viridans		60.76	0.63	ESL 90	1	1		
Staphylococcus cohnii subsp. urealyticum		38.01	0.48		99	64		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		0.87	0.14		99	99	
060561	**Good identification**						**060561**
Aerococcus viridans		100.0	0.34	ESL 90			
060562	**Acceptable identification**						**060562**
					NOV	YEP	
Aerococcus viridans		92.99	0.46	ESL 90	1	1	
Staphylococcus arlettae		7.01	0.12		99	99	
060563	**Very good identification**						**060563**
					NOV	YEP	
Aerococcus viridans		95.89	0.53	ESL 90	1	1	
Staphylococcus arlettae		4.11	0.14		99	99	
060570	**Low discrimination**						**060570**
					NOV	YEP	
Staphylococcus kloosii		69.78	0.46	PHS 99	99	99	
Aerococcus viridans		24.64	0.36		1	1	
Staphylococcus arlettae		5.57	0.12		99	99	
060571	**Low discrimination**						**060571**
					NOV	YEP	
Aerococcus viridans		79.75	0.43	ESL 90	1	1	
Staphylococcus arlettae		10.25	0.14	LAC 90	99	99	
Staphylococcus xylosus		10.00	0.05		99	40	
060572	**Low discrimination**						**060572**
					NOV	YEP	
Staphylococcus arlettae		64.98	0.53	SUC 99	99	99	
Aerococcus viridans		26.12	0.56		1	1	
Staphylococcus kloosii		8.90	0.47		99	99	
060573	**Low discrimination**						**060573**
					NOV	YEP	
Staphylococcus arlettae		57.47	0.55	SUC 99	99	99	
Aerococcus viridans		40.66	0.63		1	1	
Staphylococcus xylosus		1.61	0.14		99	40	
060602	**Low discrimination**						**060602**
					PYR		
Aerococcus viridans		84.75	0.33	ESL 90	95		
Staphylococcus warneri		15.25	0.11		8		
060603	**Low discrimination**						**060603**
					OXI		
Aerococcus viridans		84.72	0.41	ESL 90	1		
Kocuria kristinae		7.68	-.03		99		
Staphylococcus simulans		7.60	0.03		1		
060611	**Good identification**						**060611**
Aerococcus viridans		100.0	0.31	ESL 90			
060612	**Acceptable identification**						**060612**
Aerococcus viridans		92.31	0.43	ESL 90			
Staphylococcus simulans		7.69	0.04				
060613	**Low discrimination**						**060613**
					NOV		
Aerococcus viridans		86.85	0.51	ESL 90	1		
Staphylococcus simulans		10.38	0.15		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		2.77	0.09		99			
060622	**Good identification**							**060622**
Aerococcus viridans		100.0	0.33	ESL 90				
060623	**Good identification**							**060623**
Aerococcus viridans		100.0	0.41	ESL 90				
060631	**Good identification**							**060631**
Aerococcus viridans		100.0	0.31	ESL 90				
060632	**Low discrimination**							**060632**
					NOV	YEP		
Staphylococcus arlettae		63.71	0.37	MLT 99	99	99		
Aerococcus viridans		36.29	0.43		1	1		
060633	**Low discrimination**							**060633**
					NOV	YEP		
Aerococcus viridans		49.51	0.51	ESL 90	1	1		
Staphylococcus arlettae		49.38	0.40		99	99		
Staphylococcus xylosus		1.11	-.04		99	40		
060640	**Low discrimination**							**060640**
					PYR			
Staphylococcus warneri		74.61	0.47	URE 98	8			
Staphylococcus auricularis		19.60	0.27		40			
Aerococcus viridans		3.23	0.26		95			
060641	**Low discrimination**							**060641**
					OXI			
Kocuria kristinae		83.13	0.30	VPT 90	99			
Aerococcus viridans		11.32	0.34	GLR 1	1			
Staphylococcus auricularis		3.27	0.07		1			
060642	**Low discrimination**							**060642**
					PYR			
Aerococcus viridans		51.83	0.46	ESL 90	95			
Staphylococcus warneri		44.29	0.37		8			
Staphylococcus auricularis		3.88	0.07		40			
060643	**Low discrimination**							**060643**
					OXI			
Aerococcus viridans		83.05	0.53	ESL 90	1			
Kocuria kristinae		16.95	0.17		99			
060650	**Low discrimination**							**060650**
					NOV	PYR	YEP	
Staphylococcus warneri		73.18	0.44	URE 98	1	8	43	
Aerococcus viridans		13.14	0.36		1	95	1	
Staphylococcus kloosii		7.58	0.32		99	52	99	
060651	**Low discrimination**							**060651**
					OXI	PYR		
Aerococcus viridans		86.90	0.43	ESL 90	1	95		
Kocuria kristinae		8.87	0.01	LAC 90	99	93		
Staphylococcus warneri		4.23	0.09		1	8		
060652	**Low discrimination**							**060652**
					NOV	PYR		
Aerococcus viridans		54.24	0.56	ESL 90	1	95		
Staphylococcus arlettae		23.81	0.37		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		11.19	0.34		1	8	
060653	**Low discrimination**						**060653**
					NOV	YEP	
Aerococcus viridans		76.08	0.63	ESL 90	1	1	
Staphylococcus arlettae		18.97	0.40		99	99	
Staphylococcus cohnii subsp. urealyticum		3.43	0.24		99	64	
060661	**Good identification**						**060661**
Aerococcus viridans		100.0	0.34	ESL 90			
060662	**Low discrimination**						**060662**
					NOV	YEP	
Staphylococcus arlettae		56.84	0.37	MAN 99	99	99	
Aerococcus viridans		43.16	0.46		1	1	
060663	**Low discrimination**						**060663**
					NOV	YEP	
Aerococcus viridans		57.20	0.53	ESL 90	1	1	
Staphylococcus arlettae		42.80	0.40		99	99	
060670	**Low discrimination**						**060670**
					NOV	YEP	
Staphylococcus arlettae		77.54	0.37	LAC 99	99	99	
Aerococcus viridans		19.63	0.36		1	1	
Staphylococcus kloosii		2.83	0.19		99	99	
060671	**Low discrimination**						**060671**
					NOV	YEP	
Staphylococcus arlettae		64.49	0.40	LAC 99	99	99	
Aerococcus viridans		28.73	0.43		1	1	
Staphylococcus xylosus		6.77	0.10		99	40	
060672	**Very good identification**						**060672**
					NOV	YEP	
Staphylococcus arlettae		97.71	0.78		99	99	
Aerococcus viridans		2.25	0.56		1	1	
Staphylococcus kloosii		0.04	0.20		99	99	
060673	**Very good identification**						**060673**
					NOV	YEP	
Staphylococcus arlettae		95.83	0.81		99	99	
Aerococcus viridans		3.88	0.63		1	1	
Staphylococcus xylosus		0.29	0.20		99	40	
060702	**Good identification**						**060702**
Aerococcus viridans		100.0	0.33	ESL 90			
060703	**Good identification**						**060703**
Aerococcus viridans		100.0	0.41	ESL 90			
060711	**Good identification**						**060711**
Aerococcus viridans		100.0	0.31	ESL 90			
060712	**Acceptable identification**						**060712**
					NOV	YEP	
Aerococcus viridans		90.87	0.43	ESL 90	1	1	
Staphylococcus arlettae		9.13	0.12		99	99	
060713	**Acceptable identification**						**060713**
					NOV	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		92.87	0.51	ESL 90	1	1		
Staphylococcus arlettae		5.30	0.14		99	99		
Staphylococcus xylosus		1.83	-.05		99	40		
060722	**Low discrimination**							**060722**
					NOV	YEP		
Aerococcus viridans		76.83	0.33	ESL 90	1	1		
Staphylococcus arlettae		23.17	0.12		99	99		
060723	**Low discrimination**							**060723**
					NOV	YEP		
Aerococcus viridans		85.38	0.41	ESL 90	1	1		
Staphylococcus arlettae		14.62	0.14		99	99		
060731	**Low discrimination**							**060731**
					NOV	YEP		
Aerococcus viridans		44.81	0.31	ESL 90	1	1		
Staphylococcus xylosus		32.16	0.08	LAC 90	99	40		
Staphylococcus arlettae		23.03	0.14		99	99		
060732	**Acceptable identification**							**060732**
					NOV	YEP		
Staphylococcus arlettae		90.87	0.53	MLT 99	99	99		
Aerococcus viridans		9.13	0.43		1	1		
060733	**Low discrimination**							**060733**
					NOV	YEP		
Staphylococcus arlettae		82.17	0.55	MLT 99	99	99		
Aerococcus viridans		14.54	0.51		1	1		
Staphylococcus xylosus		3.30	0.17		99	40		
060740	**Low discrimination**							**060740**
					PYR			
Staphylococcus warneri		81.32	0.37	URE 98	8			
Aerococcus viridans		11.16	0.26		95			
Staphylococcus auricularis		7.52	0.07		40			
060742	**Low discrimination**							**060742**
					NOV	PYR		
Aerococcus viridans		74.34	0.46	ESL 90	1	95		
Staphylococcus warneri		20.06	0.27		1	8		
Staphylococcus arlettae		5.60	0.12		99	1		
060743	**Good identification**							**060743**
					OXI	NOV	YEP	
Aerococcus viridans		90.02	0.53	ESL 90	1	1	1	
Kocuria kristinae		6.12	0.07		99	1	15	
Staphylococcus arlettae		3.86	0.14		1	99	99	
060750	**Low discrimination**							**060750**
					NOV	PYR	YEP	
Staphylococcus warneri		39.69	0.34	URE 98	1	8	43	
Staphylococcus kloosii		25.28	0.38		99	52	99	
Aerococcus viridans		22.56	0.36		1	95	1	
060751	**Low discrimination**							**060751**
					NOV	YEP		
Aerococcus viridans		74.87	0.43	ESL 90	1	1		
Staphylococcus xylosus		15.52	0.09	LAC 90	99	40		
Staphylococcus arlettae		9.62	0.14		99	99		
060752	**Low discrimination**							**060752**
					NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		67.46	0.53	XYL 99	99	99	
Aerococcus viridans		27.12	0.56		1	1	
Staphylococcus kloosii		3.66	0.38		99	99	
060753	**Low discrimination**						**060753**
					NOV	YEP	
Staphylococcus arlettae		56.57	0.55	XYL 99	99	99	
Aerococcus viridans		40.03	0.63		1	1	
Staphylococcus xylosus		2.62	0.18		99	40	
060761	**Low discrimination**						**060761**
					NOV	YEP	
Aerococcus viridans		54.29	0.34	ESL 90	1	1	
Staphylococcus xylosus		24.78	0.06	LAC 90	99	40	
Staphylococcus arlettae		20.93	0.14		99	99	
060762	**Low discrimination**						**060762**
					NOV	YEP	
Staphylococcus arlettae		88.18	0.53	MAN 99	99	99	
Aerococcus viridans		11.82	0.46		1	1	
060763	**Low discrimination**						**060763**
					NOV	YEP	
Staphylococcus arlettae		78.75	0.55	MAN 99	99	99	
Aerococcus viridans		18.57	0.53		1	1	
Staphylococcus xylosus		2.68	0.16		99	40	
060770	**Acceptable identification**						**060770**
					NOV	YEP	
Staphylococcus arlettae		94.59	0.53	LAC 99	99	99	
Aerococcus viridans		4.23	0.36		1	1	
Staphylococcus kloosii		1.18	0.25		99	99	
060771	**Low discrimination**						**060771**
					NOV	YEP	
Staphylococcus arlettae		78.99	0.55	LAC 99	99	99	
Staphylococcus xylosus		14.80	0.31		99	40	
Aerococcus viridans		6.21	0.43		1	1	
060772	**Excellent identification**						**060772**
					NOV	YEP	
Staphylococcus arlettae		99.58	0.94		99	99	
Aerococcus viridans		0.40	0.56		1	1	
Staphylococcus kloosii		0.01	0.26		99	99	
060773	**Very good identification**						**060773**
					NOV	YEP	
Staphylococcus arlettae		98.76	0.96		99	99	
Aerococcus viridans		0.71	0.63		1	1	
Staphylococcus xylosus		0.53	0.40		99	40	
061002	**Good identification**						**061002**
Aerococcus viridans		100.0	0.27	TRE 95			
061003	**Good identification**						**061003**
Aerococcus viridans		100.0	0.34	TRE 95			
061012	**Good identification**						**061012**
Aerococcus viridans		100.0	0.36	TRE 95			
061013	**Good identification**						**061013**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.44	TRE 95			
061022	**Good identification**						**061022**
Aerococcus viridans		100.0	0.27	TRE 95			
061023	**Good identification**						**061023**
Aerococcus viridans		100.0	0.34	TRE 95			
061032	**Good identification**						**061032**
Aerococcus viridans		100.0	0.36	TRE 95			
061033	**Good identification**						**061033**
Aerococcus viridans		100.0	0.44	TRE 95			
061041	**Good identification**						**061041**
Aerococcus viridans		100.0	0.27	TRE 95			
061042	**Good identification**						**061042**
Aerococcus viridans		100.0	0.39	TRE 95			
061043	**Good identification**						**061043**
Aerococcus viridans		100.0	0.47	TRE 95			
061050	**Low discrimination**						**061050**
					OXI	NOV	
Aerococcus viridans		76.83	0.29	TRE 95	1	1	
Macrococcus equipersicus		23.17	0.02	LAC 90	99	99	
061051	**Good identification**						**061051**
Aerococcus viridans		100.0	0.37	TRE 95			
061052	**Good identification**						**061052**
Aerococcus viridans		100.0	0.49	TRE 95			
061053	**Very good identification**						**061053**
					NOV		
Aerococcus viridans		98.80	0.56	TRE 95	1		
Staphylococcus cohnii subsp. urealyticum		1.20	0.06		99		
061061	**Good identification**						**061061**
Aerococcus viridans		100.0	0.27	TRE 95			
061062	**Good identification**						**061062**
Aerococcus viridans		100.0	0.39	TRE 95			
061063	**Good identification**						**061063**
Aerococcus viridans		100.0	0.47	TRE 95			
061070	**Good identification**						**061070**
Aerococcus viridans		100.0	0.29	TRE 95			
061071	**Good identification**						**061071**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.37	TRE 95			
061072	**Good identification**						**061072**
Aerococcus viridans		100.0	0.49	TRE 95			
061073	**Very good identification**						**061073**
Aerococcus viridans		100.0	0.56	TRE 95			
061102	**Good identification**						**061102**
Aerococcus viridans		100.0	0.27	TRE 95			
061103	**Good identification**						**061103**
Aerococcus viridans		100.0	0.34	TRE 95			
061112	**Good identification**						**061112**
Aerococcus viridans		100.0	0.36	TRE 95			
061113	**Good identification**						**061113**
Aerococcus viridans		100.0	0.44	TRE 95			
061122	**Good identification**						**061122**
Aerococcus viridans		100.0	0.27	TRE 95			
061123	**Good identification**						**061123**
Aerococcus viridans		100.0	0.34	TRE 95			
061132	**Good identification**						**061132**
Aerococcus viridans		100.0	0.36	TRE 95			
061133	**Good identification**						**061133**
Aerococcus viridans		100.0	0.44	TRE 95			
061141	**Good identification**						**061141**
Aerococcus viridans		100.0	0.27	TRE 95			
061142	**Good identification**						**061142**
Aerococcus viridans		100.0	0.39	TRE 95			
061143	**Good identification**						**061143**
Aerococcus viridans		100.0	0.47	TRE 95			
061150	**Low discrimination**						**061150**
					NOV	YEP	
Aerococcus viridans		80.55	0.29	TRE 95	1	1	
Staphylococcus kloosii		19.45	0.17	LAC 90	99	99	
061151	**Good identification**						**061151**
Aerococcus viridans		100.0	0.37	TRE 95			
061152	**Good identification**						**061152**
					NOV	YEP	
Aerococcus viridans		97.18	0.49	TRE 95	1	1	
Staphylococcus kloosii		2.83	0.18		99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
061153	**Very good identification**						**061153**
Aerococcus viridans		100.0	0.56	TRE 95			
061161	**Good identification**						**061161**
Aerococcus viridans		100.0	0.27	TRE 95			
061162	**Good identification**						**061162**
Aerococcus viridans		100.0	0.39	TRE 95			
061163	**Good identification**						**061163**
Aerococcus viridans		100.0	0.47	TRE 95			
061170	**Good identification**						**061170**
Aerococcus viridans		100.0	0.29	TRE 95			
061171	**Good identification**						**061171**
Aerococcus viridans		100.0	0.37	TRE 95			
061172	**Acceptable identification**						**061172**
					NOV	YEP	
Aerococcus viridans		94.96	0.49	TRE 95	1	1	
Staphylococcus arlettae		5.04	0.12		99	99	
061173	**Very good identification**						**061173**
					NOV	YEP	
Aerococcus viridans		97.08	0.56	TRE 95	1	1	
Staphylococcus arlettae		2.93	0.14		99	99	
061202	**Good identification**						**061202**
Aerococcus viridans		100.0	0.27	TRE 95			
061203	**Good identification**						**061203**
Aerococcus viridans		100.0	0.34	TRE 95			
061212	**Good identification**						**061212**
Aerococcus viridans		100.0	0.36	TRE 95			
061213	**Good identification**						**061213**
Aerococcus viridans		100.0	0.44	TRE 95			
061222	**Good identification**						**061222**
Aerococcus viridans		100.0	0.27	TRE 95			
061223	**Good identification**						**061223**
Aerococcus viridans		100.0	0.34	TRE 95			
061232	**Good identification**						**061232**
Aerococcus viridans		100.0	0.36	TRE 95			
061233	**Good identification**						**061233**
Aerococcus viridans		100.0	0.44	TRE 95			
061241	**Good identification**						**061241**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.27	TRE 95			
061242	**Good identification**						**061242**
Aerococcus viridans		100.0	0.39	TRE 95			
061243	**Good identification**						**061243**
Aerococcus viridans		100.0	0.47	TRE 95			
061250	**Good identification**						**061250**
Aerococcus viridans		100.0	0.29	TRE 95			
061251	**Good identification**						**061251**
Aerococcus viridans		100.0	0.37	TRE 95			
061252	**Good identification**						**061252**
Aerococcus viridans		100.0	0.49	TRE 95			
061253	**Very good identification**						**061253**
Aerococcus viridans		100.0	0.56	TRE 95			
061261	**Good identification**						**061261**
Aerococcus viridans		100.0	0.27	TRE 95			
061262	**Good identification**						**061262**
Aerococcus viridans		100.0	0.39	TRE 95			
061263	**Good identification**						**061263**
Aerococcus viridans		100.0	0.47	TRE 95			
061270	**Good identification**						**061270**
Aerococcus viridans		100.0	0.29	TRE 95			
061271	**Good identification**						**061271**
Aerococcus viridans		100.0	0.37	TRE 95			
061272	**Low discrimination**						**061272**
					NOV	YEP	
Aerococcus viridans		51.90	0.49	TRE 95	1	1	
Staphylococcus arlettae		48.10	0.37		99	99	
061273	**Low discrimination**						**061273**
					NOV	YEP	
Aerococcus viridans		65.51	0.56	TRE 95	1	1	
Staphylococcus arlettae		34.49	0.40		99	99	
061302	**Good identification**						**061302**
Aerococcus viridans		100.0	0.27	TRE 95			
061303	**Good identification**						**061303**
Aerococcus viridans		100.0	0.34	TRE 95			
061312	**Good identification**						**061312**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.36	TRE 95			
061313	**Good identification**						**061313**
Aerococcus viridans		100.0	0.44	TRE 95			
061322	**Good identification**						**061322**
Aerococcus viridans		100.0	0.27	TRE 95			
061323	**Good identification**						**061323**
Aerococcus viridans		100.0	0.34	TRE 95			
061332	**Low discrimination**						**061332**
					NOV	YEP	
Aerococcus viridans		82.50	0.36	TRE 95	1	1	
Staphylococcus arlettae		17.50	0.12		99	99	
061333	**Low discrimination**						**061333**
					NOV	YEP	
Aerococcus viridans		89.24	0.44	TRE 95	1	1	
Staphylococcus arlettae		10.76	0.14		99	99	
061341	**Good identification**						**061341**
Aerococcus viridans		100.0	0.27	TRE 95			
061342	**Good identification**						**061342**
Aerococcus viridans		100.0	0.39	TRE 95			
061343	**Good identification**						**061343**
Aerococcus viridans		100.0	0.47	TRE 95			
061350	**Good identification**						**061350**
Aerococcus viridans		100.0	0.29	TRE 95			
061351	**Good identification**						**061351**
Aerococcus viridans		100.0	0.37	TRE 95			
061352	**Acceptable identification**						**061352**
					NOV	YEP	
Aerococcus viridans		94.96	0.49	TRE 95	1	1	
Staphylococcus arlettae		5.04	0.12		99	99	
061353	**Very good identification**						**061353**
					NOV	YEP	
Aerococcus viridans		97.08	0.56	TRE 95	1	1	
Staphylococcus arlettae		2.93	0.14		99	99	
061361	**Good identification**						**061361**
Aerococcus viridans		100.0	0.27	TRE 95			
061362	**Low discrimination**						**061362**
					NOV	YEP	
Aerococcus viridans		86.27	0.39	TRE 95	1	1	
Staphylococcus arlettae		13.73	0.12		99	99	
061363	**Acceptable identification**						**061363**
					NOV	YEP	
Aerococcus viridans		91.71	0.47	TRE 95	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		8.29	0.14		99	99		
061370	**Low discrimination**							**061370**
					NOV	YEP		
Aerococcus viridans		67.69	0.29	TRE 95	1	1		
Staphylococcus arlettae		32.31	0.12	LAC 90	99	99		
061371	**Low discrimination**							**061371**
					NOV	YEP		
Aerococcus viridans		78.67	0.37	TRE 95	1	1		
Staphylococcus arlettae		21.34	0.14	LAC 90	99	99		
061372	**Low discrimination**							**061372**
					NOV	YEP		
Staphylococcus arlettae		84.00	0.53	TRE 99	99	99		
Aerococcus viridans		16.00	0.49		1	1		
061373	**Low discrimination**							**061373**
					NOV	YEP		
Staphylococcus arlettae		74.59	0.55	TRE 99	99	99		
Aerococcus viridans		25.00	0.56		1	1		
Staphylococcus xylosus		0.41	-.01		99	40		
061401	**Good identification**							**061401**
Aerococcus viridans		100.0	0.41	LAC 90				
061402	**Very good identification**							**061402**
Aerococcus viridans		100.0	0.53					
061403	**Very good identification**							**061403**
					NOV			
Aerococcus viridans		98.70	0.61		1			
Staphylococcus cohnii subsp. urealyticum		1.30	0.11		99			
061410	**Low discrimination**							**061410**
					OXI	NOV	YEP	
Aerococcus viridans		73.52	0.43	LAC 90	1	1	1	
Macrococcus carouselicus		16.44	0.08		99	99	67	
Staphylococcus kloosii		10.04	0.26		1	99	99	
061411	**Good identification**							**061411**
					NOV			
Aerococcus viridans		96.19	0.51	LAC 90	1			
Staphylococcus cohnii subsp. urealyticum		3.81	0.11		99			
061412	**Very good identification**							**061412**
					NOV	YEP		
Aerococcus viridans		98.38	0.63		1	1		
Staphylococcus kloosii		1.62	0.27		99	99		
061413	**Very good identification**							**061413**
					NOV			
Aerococcus viridans		95.74	0.70		1			
Staphylococcus cohnii subsp. urealyticum		4.26	0.31		99			
061420	**Good identification**							**061420**
Aerococcus viridans		100.0	0.33	LAC 90				
061421	**Good identification**							**061421**
Aerococcus viridans		100.0	0.41	LAC 90				
061422	**Very good identification**							**061422**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.53					
061423	**Very good identification**							**061423**
Aerococcus viridans		100.0	0.61					
061430	**Good identification**							**061430**
Aerococcus viridans		100.0	0.43	LAC 90				
061431	**Good identification**							**061431**
Aerococcus viridans		100.0	0.51	LAC 90				
061432	**Very good identification**							**061432**
Aerococcus viridans		100.0	0.63					
061433	**Very good identification**							**061433**
Aerococcus viridans		100.0	0.70					
061440	**Good identification**							**061440**
Aerococcus viridans		100.0	0.46	LAC 90				
061441	**Very good identification**							**061441**
					NOV			
Aerococcus viridans		98.36	0.53	LAC 90	1			
Staphylococcus cohnii subsp. urealyticum		1.64	0.06		99			
061442	**Very good identification**							**061442**
Aerococcus viridans		100.0	0.65					
061443	**Very good identification**							**061443**
					NOV			
Aerococcus viridans		98.16	0.73		1			
Staphylococcus cohnii subsp. urealyticum		1.84	0.26		99			
061450	**Low discrimination**							**061450**
					OXI	NOV	YEP	
Aerococcus viridans		59.63	0.56	LAC 90	1	1	1	
Staphylococcus kloosii		38.66	0.52		1	99	99	
Macrococcus equipersicus		0.95	0.02		99	99	99	
061451	**Good identification**							**061451**
					NOV			
Aerococcus viridans		94.68	0.63	LAC 90	1			
Staphylococcus cohnii subsp. urealyticum		5.32	0.26		99			
061452	**Good identification**							**061452**
					NOV	YEP		
Aerococcus viridans		92.63	0.75		1	1		
Staphylococcus kloosii		7.23	0.53		99	99		
Staphylococcus cohnii subsp. urealyticum		0.14	0.06		99	64		
061453	**Good identification**							**061453**
					NOV			
Aerococcus viridans		94.07	0.83		1			
Staphylococcus cohnii subsp. urealyticum		5.93	0.47		99			
061460	**Good identification**							**061460**
Aerococcus viridans		100.0	0.46	LAC 90				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
061461	**Very good identification**							**061461**
Aerococcus viridans		100.0	0.53	LAC 90				
061462	**Very good identification**							**061462**
Aerococcus viridans		100.0	0.65					
061463	**Very good identification**							**061463**
Aerococcus viridans		100.0	0.73					
061470	**Low discrimination**							**061470**
					NOV	YEP		
Aerococcus viridans		86.05	0.56	LAC 90	1	1		
Staphylococcus kloosii		13.95	0.40		99	99		
061471	**Very good identification**							**061471**
Aerococcus viridans		100.0	0.63	LAC 90				
061472	**Good identification**							**061472**
					NOV	YEP		
Aerococcus viridans		93.61	0.75		1	1		
Staphylococcus arlettae		4.57	0.37		99	99		
Staphylococcus kloosii		1.83	0.41		99	99		
061473	**Very good identification**							**061473**
					NOV	YEP		
Aerococcus viridans		97.24	0.83		1	1		
Staphylococcus arlettae		2.70	0.40		99	99		
Staphylococcus cohnii subsp. urealyticum		0.06	0.06		99	64		
061500	**Low discrimination**							**061500**
					OXI	NOV		
Aerococcus viridans		68.98	0.33	LAC 90	1	1		
Macrococcus carouselicus		31.02	0.04		99	99		
061501	**Good identification**							**061501**
Aerococcus viridans		100.0	0.41	LAC 90				
061502	**Very good identification**							**061502**
Aerococcus viridans		100.0	0.53					
061503	**Very good identification**							**061503**
Aerococcus viridans		100.0	0.61					
061510	**Low discrimination**							**061510**
					OXI	NOV	YEP	
Aerococcus viridans		73.49	0.43	LAC 90	1	1	1	
Staphylococcus kloosii		19.47	0.32		1	99	99	
Macrococcus carouselicus		7.04	0.00		99	99	67	
061511	**Good identification**							**061511**
Aerococcus viridans		100.0	0.51	LAC 90				
061512	**Very good identification**							**061512**
					NOV	YEP		
Aerococcus viridans		96.91	0.63		1	1		
Staphylococcus kloosii		3.09	0.33		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
061513	**Very good identification**						**061513**
					NOV		
Aerococcus viridans		98.13	0.70		1		
Staphylococcus cohnii subsp. urealyticum		1.87	0.24		99		
061520	**Good identification**						**061520**
Aerococcus viridans		100.0	0.33	LAC 90			
061521	**Good identification**						**061521**
Aerococcus viridans		100.0	0.41	LAC 90			
061522	**Very good identification**						**061522**
Aerococcus viridans		100.0	0.53				
061523	**Very good identification**						**061523**
Aerococcus viridans		100.0	0.61				
061530	**Acceptable identification**						**061530**
					NOV	YEP	
Aerococcus viridans		93.79	0.43	LAC 90	1	1	
Staphylococcus kloosii		6.21	0.20		99	99	
061531	**Good identification**						**061531**
Aerococcus viridans		100.0	0.51	LAC 90			
061532	**Very good identification**						**061532**
					NOV	YEP	
Aerococcus viridans		98.12	0.63		1	1	
Staphylococcus arlettae		1.10	0.12		99	99	
Staphylococcus kloosii		0.78	0.20		99	99	
061533	**Very good identification**						**061533**
					NOV	YEP	
Aerococcus viridans		99.37	0.70		1	1	
Staphylococcus arlettae		0.63	0.14		99	99	
061540	**Good identification**						**061540**
					NOV	YEP	
Aerococcus viridans		96.33	0.46	LAC 90	1	1	
Staphylococcus kloosii		3.67	0.17		99	99	
061541	**Very good identification**						**061541**
Aerococcus viridans		100.0	0.53	LAC 90			
061542	**Very good identification**						**061542**
					NOV	YEP	
Aerococcus viridans		99.54	0.65		1	1	
Staphylococcus kloosii		0.46	0.18		99	99	
061543	**Very good identification**						**061543**
					NOV		
Aerococcus viridans		99.20	0.73		1		
Staphylococcus cohnii subsp. urealyticum		0.80	0.19		99		
061550	**Low discrimination**						**061550**
					NOV	YEP	
Staphylococcus kloosii		55.72	0.58	PHS 99	99	99	
Aerococcus viridans		44.28	0.56		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
061551	**Very good identification**						**061551**
					NOV	YEP	
Aerococcus viridans		97.13	0.63	LAC 90	1	1	
Staphylococcus cohnii subsp. urealyticum		2.34	0.19		99	64	
Staphylococcus kloosii		0.53	0.17		99	99	
061552	**Low discrimination**						**061552**
					NOV	YEP	
Aerococcus viridans		86.63	0.75		1	1	
Staphylococcus kloosii		13.12	0.59		99	99	
Staphylococcus arlettae		0.24	0.12		99	99	
061553	**Very good identification**						**061553**
					NOV	YEP	
Aerococcus viridans		97.16	0.83		1	1	
Staphylococcus cohnii subsp. urealyticum		2.63	0.39		99	64	
Staphylococcus arlettae		0.15	0.14		99	99	
061560	**Good identification**						**061560**
Aerococcus viridans		100.0	0.46	LAC 90			
061561	**Very good identification**						**061561**
Aerococcus viridans		100.0	0.53	LAC 90			
061562	**Very good identification**						**061562**
					NOV	YEP	
Aerococcus viridans		99.17	0.65		1	1	
Staphylococcus arlettae		0.83	0.12		99	99	
061563	**Very good identification**						**061563**
					NOV	YEP	
Aerococcus viridans		99.53	0.73		1	1	
Staphylococcus arlettae		0.47	0.14		99	99	
061570	**Low discrimination**						**061570**
					NOV	YEP	
Aerococcus viridans		74.64	0.56	LAC 90	1	1	
Staphylococcus kloosii		23.48	0.46		99	99	
Staphylococcus arlettae		1.88	0.12		99	99	
061571	**Very good identification**						**061571**
					NOV	YEP	
Aerococcus viridans		98.59	0.63	LAC 90	1	1	
Staphylococcus arlettae		1.41	0.14		99	99	
061572	**Low discrimination**						**061572**
					NOV	YEP	
Aerococcus viridans		76.09	0.75		1	1	
Staphylococcus arlettae		21.03	0.53		99	99	
Staphylococcus kloosii		2.88	0.47		99	99	
061573	**Low discrimination**						**061573**
					NOV	YEP	
Aerococcus viridans		86.35	0.83		1	1	
Staphylococcus arlettae		13.56	0.55		99	99	
Staphylococcus xylosus		0.10	0.02		99	40	
061601	**Acceptable identification**						**061601**
					OXI		
Aerococcus viridans		91.69	0.41	LAC 90	1		
Kocuria kristinae		8.32	-.03		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
061602	**Very good identification**						**061602**
Aerococcus viridans		100.0	0.53				
061603	**Very good identification**						**061603**
Aerococcus viridans		100.0	0.61				
061610	**Low discrimination**						**061610**
					OXI	NOV	
Aerococcus viridans		68.77	0.43	LAC 90	1	1	
Macrococcus carouselicus		31.23	0.14		99	99	
061611	**Good identification**						**061611**
Aerococcus viridans		100.0	0.51	LAC 90			
061612	**Very good identification**						**061612**
Aerococcus viridans		100.0	0.63				
061613	**Very good identification**						**061613**
Aerococcus viridans		100.0	0.70				
061620	**Good identification**						**061620**
Aerococcus viridans		100.0	0.33	LAC 90			
061621	**Good identification**						**061621**
Aerococcus viridans		100.0	0.41	LAC 90			
061622	**Very good identification**						**061622**
Aerococcus viridans		100.0	0.53				
061623	**Very good identification**						**061623**
Aerococcus viridans		100.0	0.61				
061630	**Good identification**						**061630**
					OXI	NOV	
Aerococcus viridans		95.19	0.43	LAC 90	1	1	
Staphylococcus vitulinus		4.81	-.08		99	99	
061631	**Good identification**						**061631**
Aerococcus viridans		100.0	0.51	LAC 90			
061632	**Low discrimination**						**061632**
					NOV	YEP	
Aerococcus viridans		83.68	0.63		1	1	
Staphylococcus arlettae		16.33	0.37		99	99	
061633	**Good identification**						**061633**
					NOV	YEP	
Aerococcus viridans		90.02	0.70		1	1	
Staphylococcus arlettae		9.98	0.40		99	99	
061640	**Good identification**						**061640**
Aerococcus viridans		100.0	0.46	LAC 90			
061641	**Low discrimination**						**061641**
					OXI		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		83.05	0.53	LAC 90	1			
Kocuria kristinae		16.95	0.17		99			
061642	**Very good identification**							**061642**
Aerococcus viridans		100.0	0.65					
061643	**Very good identification**							**061643**
					OXI			
Aerococcus viridans		99.44	0.73		1			
Kocuria kristinae		0.56	0.05		99			
061650	**Good identification**							**061650**
					NOV	YEP		
Aerococcus viridans		93.97	0.56	LAC 90	1	1		
Staphylococcus kloosii		6.03	0.32		99	99		
061651	**Very good identification**							**061651**
Aerococcus viridans		100.0	0.63	LAC 90				
061652	**Good identification**							**061652**
					NOV	YEP		
Aerococcus viridans		94.65	0.75		1	1		
Staphylococcus arlettae		4.62	0.37		99	99		
Staphylococcus kloosii		0.73	0.32		99	99		
061653	**Very good identification**							**061653**
					NOV	YEP		
Aerococcus viridans		97.12	0.83		1	1		
Staphylococcus arlettae		2.69	0.40		99	99		
Staphylococcus cohnii subsp. urealyticum		0.19	0.16		99	64		
061660	**Good identification**							**061660**
Aerococcus viridans		100.0	0.46	LAC 90				
061661	**Very good identification**							**061661**
Aerococcus viridans		100.0	0.53	LAC 90				
061662	**Low discrimination**							**061662**
					NOV	YEP		
Aerococcus viridans		87.24	0.65		1	1		
Staphylococcus arlettae		12.77	0.37		99	99		
061663	**Good identification**							**061663**
					NOV	YEP		
Aerococcus viridans		92.33	0.73		1	1		
Staphylococcus arlettae		7.68	0.40		99	99		
061670	**Low discrimination**							**061670**
					NOV	YEP		
Aerococcus viridans		68.73	0.56	LAC 90	1	1		
Staphylococcus arlettae		30.17	0.37		99	99		
Staphylococcus kloosii		1.10	0.19		99	99		
061671	**Low discrimination**							**061671**
					NOV	YEP	PYR	
Aerococcus viridans		79.02	0.63	LAC 90	1	1	95	
Staphylococcus arlettae		19.71	0.40		99	99	1	
Staphylococcus equorum		0.75	0.03		99	1	5	
061672	**Low discrimination**							**061672**
					NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		82.82	0.78		99	99		
Aerococcus viridans		17.15	0.75		1	1		
Staphylococcus kloosii		0.03	0.20		99	99		
061673	**Low discrimination**							**061673**
					NOV	YEP	PYR	
Staphylococcus arlettae		73.19	0.81		99	99	1	
Aerococcus viridans		26.68	0.83		1	1	95	
Staphylococcus equorum		0.07	0.11		99	1	5	
061700	**Low discrimination**							**061700**
					OXI	NOV		
Aerococcus viridans		52.27	0.33	LAC 90	1	1		
Macrococcus carouselicus		47.73	0.10		99	99		
061701	**Good identification**							**061701**
Aerococcus viridans		100.0	0.41	LAC 90				
061702	**Very good identification**							**061702**
Aerococcus viridans		100.0	0.53					
061703	**Very good identification**							**061703**
Aerococcus viridans		100.0	0.61					
061710	**Low discrimination**							**061710**
					OXI	NOV		
Aerococcus viridans		83.71	0.43	LAC 90	1	1		
Macrococcus carouselicus		16.29	0.06		99	99		
061711	**Good identification**							**061711**
Aerococcus viridans		100.0	0.51	LAC 90				
061712	**Very good identification**							**061712**
					NOV	YEP		
Aerococcus viridans		98.90	0.63		1	1		
Staphylococcus arlettae		1.10	0.12		99	99		
061713	**Very good identification**							**061713**
					OXI	NOV	YEP	
Aerococcus viridans		98.70	0.70		1	1	1	
Staphylococcus lentus		0.67	0.12		99	99	1	
Staphylococcus arlettae		0.63	0.14		1	99	99	
061720	**Good identification**							**061720**
Aerococcus viridans		100.0	0.33	LAC 90				
061721	**Good identification**							**061721**
Aerococcus viridans		100.0	0.41	LAC 90				
061722	**Very good identification**							**061722**
					NOV	YEP		
Aerococcus viridans		96.76	0.53		1	1		
Staphylococcus arlettae		3.24	0.12		99	99		
061723	**Very good identification**							**061723**
					NOV	YEP		
Aerococcus viridans		98.13	0.61		1	1		
Staphylococcus arlettae		1.87	0.14		99	99		
061730	**Acceptable identification**							**061730**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		90.87	0.43	LAC 90	1	1		
Staphylococcus arlettae		9.13	0.12		99	99		
061731	**Low discrimination**							**061731**
					NOV	YEP	PYR	
Aerococcus viridans		89.68	0.51	LAC 90	1	1	95	
Staphylococcus arlettae		5.12	0.14		99	99	1	
Staphylococcus equorum		3.41	0.03		99	1	5	
061732	**Low discrimination**							**061732**
					NOV	YEP		
Staphylococcus arlettae		52.51	0.53	MLT 99	99	99		
Aerococcus viridans		47.49	0.63		1	1		
061733	**Low discrimination**							**061733**
					NOV	YEP	PYR	
Aerococcus viridans		60.53	0.70		1	1	95	
Staphylococcus arlettae		38.02	0.55		99	99	1	
Staphylococcus equorum		0.66	0.11		99	1	5	
061740	**Good identification**							**061740**
Aerococcus viridans		100.0	0.46	LAC 90				
061741	**Good identification**							**061741**
					OXI			
Aerococcus viridans		93.63	0.53	LAC 90	1			
Kocuria kristinae		6.37	0.07		99			
061742	**Very good identification**							**061742**
					NOV	YEP		
Aerococcus viridans		99.17	0.65		1	1		
Staphylococcus arlettae		0.83	0.12		99	99		
061743	**Very good identification**							**061743**
					OXI	NOV	YEP	
Aerococcus viridans		99.34	0.73		1	1	1	
Staphylococcus arlettae		0.47	0.14		1	99	99	
Kocuria kristinae		0.19	-.05		99	1	15	
061750	**Low discrimination**							**061750**
					NOV	YEP		
Aerococcus viridans		86.99	0.56	LAC 90	1	1		
Staphylococcus kloosii		10.83	0.38		99	99		
Staphylococcus arlettae		2.19	0.12		99	99		
061751	**Very good identification**							**061751**
					OXI	NOV	YEP	
Aerococcus viridans		96.51	0.63	LAC 90	1	1	1	
Staphylococcus arlettae		1.38	0.14		1	99	99	
Staphylococcus lentus		0.85	0.07		99	99	1	
061752	**Low discrimination**							**061752**
					NOV	YEP		
Aerococcus viridans		77.44	0.75		1	1		
Staphylococcus arlettae		21.40	0.53		99	99		
Staphylococcus kloosii		1.16	0.38		99	99		
061753	**Low discrimination**							**061753**
					OXI	NOV	YEP	
Aerococcus viridans		85.98	0.83		1	1	1	
Staphylococcus arlettae		13.50	0.55		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		0.30	0.18		99	99	1	
061760	**Acceptable identification**							**061760**
					NOV	YEP		
Aerococcus viridans		92.99	0.46	LAC 90	1	1		
Staphylococcus arlettae		7.01	0.12		99	99		
061761	**Very good identification**							**061761**
					NOV	YEP		
Aerococcus viridans		95.89	0.53	LAC 90	1	1		
Staphylococcus arlettae		4.11	0.14		99	99		
061762	**Low discrimination**							**061762**
					NOV	YEP		
Aerococcus viridans		54.67	0.65		1	1		
Staphylococcus arlettae		45.33	0.53		99	99		
061763	**Low discrimination**							**061763**
					NOV	YEP		
Aerococcus viridans		67.79	0.73		1	1		
Staphylococcus arlettae		31.94	0.55		99	99		
Staphylococcus xylosus		0.27	0.03		99	40		
061770	**Low discrimination**							**061770**
					NOV	YEP		
Staphylococcus arlettae		70.70	0.53	LAC 99	99	99		
Aerococcus viridans		28.42	0.56		1	1		
Staphylococcus kloosii		0.88	0.25		99	99		
061771	**Low discrimination**							**061771**
					NOV	YEP	PYR	
Staphylococcus arlettae		54.56	0.55	LAC 99	99	99	1	
Aerococcus viridans		38.61	0.63		1	1	95	
Staphylococcus equorum		3.72	0.23		99	1	5	
061772	**Very good identification**							**061772**
					NOV	YEP		
Staphylococcus arlettae		96.46	0.94		99	99		
Aerococcus viridans		3.53	0.75		1	1		
Staphylococcus kloosii		0.01	0.26		99	99		
061773	**Good identification**							**061773**
					NOV	YEP	PYR	
Staphylococcus arlettae		93.66	0.96		99	99	1	
Aerococcus viridans		6.03	0.83		1	1	95	
Staphylococcus equorum		0.17	0.32		99	1	5	
062000	**Acceptable identification**							**062000**
					OXI			
Kocuria rosea		94.11	0.53	BGA 1	35			
Kocuria varians		3.89	0.33		13			
Dermacoccus nishinomiyaensis		0.94	0.13		93			
062010	**Good identification**							**062010**
Kocuria rosea		100.0	0.33	BGA 1				
062020	**Good identification**							**062020**
Kocuria rosea		96.72	0.45	BGA 1				
Kocuria varians		3.28	0.24					
062100	**Low discrimination**							**062100**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Kocuria varians		83.79	0.37	GLR 1	13	
Kocuria rosea		13.64	0.12		35	
Dermacoccus nishinomiyaensis		2.58	-.02		93	
062120	**Low discrimination**					**062120**
Kocuria varians		83.44	0.27	GLR 1		
Kocuria rosea		16.56	0.04			
062200	**Low discrimination**					**062200**
Kocuria rosea		89.16	0.33	BGA 1		
Staphylococcus auricularis		6.32	0.17	SUC 10		
Kocuria varians		4.53	0.15			
062373	**Acceptable identification**					**062373**
Staphylococcus xylosus		94.68	0.31	URE 99		
Staphylococcus arlettae		5.32	0.14	TRE 96		
062400	**Low discrimination**					**062400**
Staphylococcus auricularis		86.64	0.40	BGA 14		
Kocuria rosea		8.92	0.12	GLR 3		
Kocuria varians		2.75	0.10			
062440	**Good identification**					**062440**
					NOV	
Staphylococcus auricularis		96.60	0.47	BGA 14	1	
Staphylococcus cohnii subsp. cohnii		2.55	0.18	GLR 3	99	
Staphylococcus warneri		0.85	0.13		1	
062441	**Low discrimination**					**062441**
					NOV	
Staphylococcus auricularis		79.53	0.28	BGA 14	1	
Staphylococcus cohnii subsp. cohnii		20.47	0.19	GLR 3	99	
062442	**Good identification**					**062442**
Staphylococcus auricularis		100.0	0.28	BGA 14		
062453	**Acceptable identification**					**062453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		91.63	0.40	URE 99	99	
Staphylococcus simulans		4.78	0.08		1	
Aerococcus viridans		3.59	0.22		1	
062540	**Good identification**					**062540**
Staphylococcus auricularis		100.0	0.28	BGA 14		
062550	**Good identification**					**062550**
Staphylococcus saprophyticus subsp. bovis		97.90	0.34	URE 99		
Staphylococcus kloosii		2.10	0.17			
062553	**Low discrimination**					**062553**
Staphylococcus cohnii subsp. urealyticum		67.19	0.32	URE 99		
Staphylococcus xylosus		16.99	0.11			
Staphylococcus saprophyticus subsp. bovis		9.68	0.09			
062573	**Acceptable identification**					**062573**
					NOV YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		92.94	0.33	URE 99	99	40	
Staphylococcus arlettae		4.13	0.14	SUC 95	99	99	
Aerococcus viridans		2.93	0.22		1	1	
062600	**Good identification**						**062600**
Staphylococcus auricularis		97.82	0.39	BGA 14			
Staphylococcus warneri		2.18	0.13	GLR 3			
062603	**Low discrimination**						**062603**
Staphylococcus simulans		87.50	0.31	URE 98			
Staphylococcus hyicus		12.50	0.07	ARG 92			
062612	**Good identification**						**062612**
Staphylococcus simulans		100.0	0.33	URE 98			
062613	**Good identification**						**062613**
Staphylococcus simulans		100.0	0.43	URE 98			
062640	**Low discrimination**						**062640**
					PYR		
Staphylococcus auricularis		66.85	0.46	BGA 14	40		
Staphylococcus haemolyticus		21.04	0.37	GLR 3	98		
Staphylococcus warneri		12.12	0.39		8		
062641	**Low discrimination**						**062641**
					OXI		
Staphylococcus auricularis		76.90	0.27	BGA 14	1		
Kocuria kristinae		16.37	0.06	GLR 3	99		
Staphylococcus haemolyticus		6.74	0.06	MNS 10	1		
062642	**Low discrimination**						**062642**
					PYR		
Staphylococcus haemolyticus		47.04	0.32	VPT 86	98		
Staphylococcus auricularis		29.52	0.27	ARG 92	40		
Staphylococcus warneri		16.06	0.30		8		
062650	**Low discrimination**						**062650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		47.39	0.34	URE 99	99	99	
Staphylococcus haemolyticus		29.81	0.32		2	98	
Staphylococcus warneri		21.17	0.37		1	8	
062653	**Low discrimination**						**062653**
					NOV		
Staphylococcus simulans		78.85	0.36	URE 98	1		
Staphylococcus xylosus		12.84	0.16	ARG 92	99		
Staphylococcus saprophyticus subsp. bovis		3.89	0.09		99		
062671	**Good identification**						**062671**
Staphylococcus xylosus		100.0	0.29	URE 99			
062672	**Good identification**						**062672**
Staphylococcus arlettae		96.86	0.37	NIT 1			
Staphylococcus xylosus		3.14	-.03				
062673	**Low discrimination**						**062673**
					NOV	YEP	
Staphylococcus xylosus		69.92	0.38	URE 99	99	40	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		28.91	0.40	GAL 91	99	99	
Aerococcus viridans		1.17	0.22		1	1	
062703	**Acceptable identification**						**062703**
					COA	PYR	
Staphylococcus hyicus		91.19	0.28	ARG 96	24	1	
Staphylococcus simulans		4.83	0.09	BGA 10	1	70	
Staphylococcus intermedius		3.98	0.06	PHS 92	99	90	
062731	**Good identification**						**062731**
Staphylococcus xylosus		100.0	0.26	URE 99			
062733	**Acceptable identification**						**062733**
					OXI	YEP	
Staphylococcus xylosus		93.94	0.36	URE 99	1	40	
Staphylococcus lentus		3.14	0.12	MLT 93	99	1	
Staphylococcus arlettae		2.92	0.14		1	99	
062740	**Low discrimination**						**062740**
					PYR		
Staphylococcus haemolyticus		65.54	0.38	VPT 86	98		
Staphylococcus auricularis		20.52	0.27	ARG 92	40		
Staphylococcus warneri		10.57	0.29		8		
062742	**Low discrimination**						**062742**
					PYR		
Staphylococcus haemolyticus		86.40	0.33	VPT 86	98		
Staphylococcus warneri		8.26	0.19	ARG 92	8		
Staphylococcus auricularis		5.34	0.07		40		
062750	**Low discrimination**						**062750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		82.47	0.47	URE 99	99	99	
Staphylococcus haemolyticus		14.63	0.34		2	98	
Staphylococcus warneri		2.91	0.26		1	8	
062751	**Low discrimination**						**062751**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		54.88	0.34	URE 99	1	99	
Staphylococcus xylosus		40.19	0.28		1	65	
Staphylococcus lentus		1.89	0.07		99	6	
062752	**Low discrimination**						**062752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		64.84	0.34	URE 99	99	99	
Staphylococcus haemolyticus		25.87	0.28		2	98	
Staphylococcus warneri		3.05	0.16		1	8	
062753	**Low discrimination**						**062753**
					COA	NOV	
Staphylococcus xylosus		72.69	0.37	URE 99	1	99	
Staphylococcus saprophyticus subsp. bovis		8.72	0.22	XYL 92	1	99	
Staphylococcus intermedius		7.15	0.18		99	1	
062763	**Good identification**						**062763**
Staphylococcus xylosus		96.46	0.34	URE 99			
Staphylococcus arlettae		3.54	0.14	MAN 94			
062771	**Good identification**						**062771**
					YEP		
Staphylococcus xylosus		98.43	0.50	URE 99	40		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		0.66	0.14		99		
Staphylococcus equorum		0.51	0.04		1		
062772	**Acceptable identification**						**062772**
Staphylococcus arlettae		94.54	0.53	NIT 1			
Staphylococcus xylosus		5.46	0.18				
062773	**Low discrimination**						**062773**
					OXI	YEP	
Staphylococcus xylosus		80.44	0.59	URE 99	1	40	
Staphylococcus arlettae		18.64	0.55		1	99	
Staphylococcus lentus		0.41	0.18		99	1	
063413	**Low discrimination**						**063413**
					NOV		
Aerococcus viridans		56.25	0.29	NIT 1	1		
Staphylococcus cohnii subsp. urealyticum		43.75	0.16		99		
063433	**Good identification**						**063433**
Aerococcus viridans		100.0	0.29	NIT 1			
063443	**Low discrimination**						**063443**
					NOV		
Aerococcus viridans		75.36	0.32	NIT 1	1		
Staphylococcus cohnii subsp. urealyticum		24.64	0.11		99		
063452	**Good identification**						**063452**
Aerococcus viridans		100.0	0.34	NIT 1			
063453	**Low discrimination**						**063453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		52.42	0.31	URE 99	99		
Aerococcus viridans		47.58	0.42		1		
063463	**Good identification**						**063463**
Aerococcus viridans		100.0	0.32	NIT 1			
063472	**Good identification**						**063472**
Aerococcus viridans		100.0	0.34	NIT 1			
063473	**Acceptable identification**						**063473**
					NOV		
Aerococcus viridans		91.97	0.42	NIT 1	1		
Staphylococcus xylosus		8.03	-.01		99		
063513	**Low discrimination**						**063513**
					OXI	NOV	
Aerococcus viridans		49.80	0.29	NIT 1	1	1	
Staphylococcus lentus		33.60	0.12		99	99	
Staphylococcus cohnii subsp. urealyticum		16.60	0.08		1	99	
063533	**Low discrimination**						**063533**
					OXI	NOV	
Aerococcus viridans		51.54	0.29	NIT 1	1	1	
Staphylococcus lentus		34.77	0.12		99	99	
Staphylococcus xylosus		13.70	-.03		1	99	
063543	**Good identification**						**063543**
Aerococcus viridans		100.0	0.32	NIT 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
063552	**Low discrimination**						**063552**
					NOV	YEP	
Aerococcus viridans		86.84	0.34	NIT 1	1	1	
Staphylococcus kloosii		13.16	0.18		99	99	
063553	**Low discrimination**						**063553**
					OXI	NOV	
Aerococcus viridans		52.87	0.42	NIT 1	1	1	
Staphylococcus cohnii subsp. urealyticum		24.97	0.24		1	99	
Staphylococcus lentus		18.10	0.18		99	99	
063563	**Low discrimination**						**063563**
					NOV		
Aerococcus viridans		85.54	0.32	NIT 1	1		
Staphylococcus xylosus		14.46	-.04		99		
063572	**Low discrimination**						**063572**
					NOV	YEP	
Aerococcus viridans		78.35	0.34	NIT 1	1	1	
Staphylococcus arlettae		21.65	0.12		99	99	
063573	**Low discrimination**						**063573**
					OXI	NOV	
Aerococcus viridans		37.06	0.42	NIT 1	1	1	
Staphylococcus xylosus		32.71	0.20		1	99	
Staphylococcus lentus		12.69	0.18		99	99	
063613	**Low discrimination**						**063613**
					OXI	NOV	
Aerococcus viridans		42.64	0.29	NIT 1	1	1	
Staphylococcus lentus		28.77	0.12		99	99	
Staphylococcus sciuri		15.00	0.04		99	99	
063643	**Good identification**						**063643**
Aerococcus viridans		100.0	0.32	NIT 1			
063651	**Acceptable identification**						**063651**
					OXI	NOV	
Staphylococcus sciuri		90.73	0.32	BGA 1	99	99	
Aerococcus viridans		4.96	0.22	GLR 14	1	1	
Staphylococcus lentus		4.31	0.07		99	99	
063652	**Good identification**						**063652**
Aerococcus viridans		100.0	0.34	NIT 1			
063653	**Low discrimination**						**063653**
					OXI	NOV	
Aerococcus viridans		55.02	0.42	NIT 1	1	1	
Staphylococcus lentus		18.84	0.18		99	99	
Staphylococcus sciuri		18.20	0.16		99	99	
063663	**Low discrimination**						**063663**
					NOV		
Aerococcus viridans		75.90	0.32	NIT 1	1		
Staphylococcus xylosus		24.11	0.01		99		
063671	**Low discrimination**						**063671**
					OXI		
Staphylococcus sciuri		43.46	0.30	BGA 1	99		
Staphylococcus equorum		34.13	0.24	GLR 14	1		
Staphylococcus xylosus		16.52	0.17		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
063672	**Low discrimination**							**063672**
					NOV	YEP		
Staphylococcus arlettae		82.84	0.37	NIT 1	99	99		
Aerococcus viridans		17.16	0.34		1	1		
063673	**Low discrimination**							**063673**
					YEP			
Staphylococcus equorum		34.06	0.33	URE 99	1			
Staphylococcus arlettae		30.18	0.40	GAL 91	99			
Staphylococcus xylosus		18.25	0.26		40			
063711	**Low discrimination**							**063711**
Staphylococcus lentus		81.79	0.41	GLR 1				
Staphylococcus sciuri		18.21	0.26					
063713	**Good identification**							**063713**
					OXI	NOV		
Staphylococcus lentus		96.92	0.53	GLR 1	99	99		
Aerococcus viridans		1.45	0.29		1	1		
Staphylococcus sciuri		0.99	0.10		99	99		
063731	**Low discrimination**							**063731**
					OXI			
Staphylococcus lentus		65.89	0.41	GLR 1	99			
Staphylococcus equorum		16.84	0.24		1			
Staphylococcus sciuri		11.07	0.24		99			
063733	**Low discrimination**							**063733**
					OXI			
Staphylococcus lentus		77.33	0.53	GLR 1	99			
Staphylococcus equorum		14.34	0.33		1			
Staphylococcus xylosus		5.85	0.23		1			
063751	**Low discrimination**							**063751**
					OXI			
Staphylococcus lentus		67.09	0.48	GLR 1	99			
Staphylococcus sciuri		27.68	0.38		99			
Staphylococcus xylosus		3.60	0.15		1			
063752	**Low discrimination**							**063752**
					OXI	NOV	YEP	
Aerococcus viridans		44.24	0.34	NIT 1	1	1	1	
Staphylococcus lentus		35.34	0.18		99	99	1	
Staphylococcus arlettae		12.23	0.12		1	99	99	
063753	**Good identification**							**063753**
					OXI	NOV		
Staphylococcus lentus		90.46	0.59	GLR 1	99	99		
Staphylococcus xylosus		3.89	0.25		1	99		
Aerococcus viridans		2.67	0.42		1	1		
063771	**Low discrimination**							**063771**
					OXI			
Staphylococcus equorum		36.92	0.45	URE 99	1			
Staphylococcus lentus		29.00	0.48		99			
Staphylococcus xylosus		17.87	0.37		1			
063772	**Low discrimination**							**063772**
					OXI	NOV	YEP	
Staphylococcus arlettae		89.91	0.53	NIT 1	1	99	99	
Aerococcus viridans		3.29	0.34		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		2.63	0.18		99	99	1	
063773	**Low discrimination**							**063773**
					OXI			
Staphylococcus lentus		32.55	0.59	GLR 1	99			
Staphylococcus equorum		30.06	0.54		1			
Staphylococcus xylosus		16.10	0.47		1			
064012	**Good identification**							**064012**
Staphylococcus kloosii		100.0	0.27	TRE 99				
064050	**Good identification**							**064050**
Staphylococcus kloosii		100.0	0.52	TRE 99				
064052	**Very good identification**							**064052**
Staphylococcus kloosii		100.0	0.53	TRE 99				
064070	**Good identification**							**064070**
Staphylococcus kloosii		100.0	0.40	TRE 99				
064072	**Good identification**							**064072**
Staphylococcus kloosii		100.0	0.41	TRE 99				
064110	**Good identification**							**064110**
Staphylococcus kloosii		100.0	0.32	TRE 99				
064112	**Good identification**							**064112**
Staphylococcus kloosii		100.0	0.33	TRE 99				
064150	**Very good identification**							**064150**
Staphylococcus kloosii		100.0	0.58	TRE 99				
064152	**Very good identification**							**064152**
Staphylococcus kloosii		100.0	0.59	TRE 99				
064170	**Good identification**							**064170**
Staphylococcus kloosii		100.0	0.46	TRE 99				
064172	**Acceptable identification**							**064172**
Staphylococcus kloosii		90.04	0.47	TRE 99				
Staphylococcus arlettae		9.96	0.15					
064250	**Good identification**							**064250**
Staphylococcus kloosii		100.0	0.32	SUC 9				
064252	**Good identification**							**064252**
Staphylococcus kloosii		100.0	0.32	SUC 9				
064272	**Good identification**							**064272**
Staphylococcus arlettae		97.43	0.41	TRE 99				
Staphylococcus kloosii		2.57	0.20					
064273	**Good identification**							**064273**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus arlettae		100.0	0.43	TRE 99		
064350	**Good identification**					**064350**
Staphylococcus kloosii		100.0	0.38	SUC 9		
064352	**Low discrimination**					**064352**
Staphylococcus kloosii		78.16	0.38	SUC 9		
Staphylococcus arlettae		21.85	0.15	TRE 99		
064372	**Very good identification**					**064372**
Staphylococcus arlettae		99.11	0.56	TRE 99		
Staphylococcus kloosii		0.90	0.26			
064373	**Very good identification**					**064373**
Staphylococcus arlettae		99.58	0.59	TRE 99		
Staphylococcus xylosus		0.42	0.01			
064402	**Good identification**					**064402**
Staphylococcus kloosii		100.0	0.27	MAN 99		
064410	**Very good identification**					**064410**
					YEP	
Staphylococcus kloosii		99.11	0.67	MLT 95	99	
Staphylococcus cohnii subsp. cohnii		0.90	0.12		12	
064412	**Very good identification**					**064412**
Staphylococcus kloosii		100.0	0.68	MLT 95		
064413	**Good identification**					**064413**
Staphylococcus cohnii subsp. urealyticum		96.06	0.43	URE 99		
Staphylococcus kloosii		3.94	0.27			
064430	**Very good identification**					**064430**
Staphylococcus kloosii		100.0	0.55	MLT 95		
064432	**Very good identification**					**064432**
Staphylococcus kloosii		100.0	0.55	MLT 95		
064440	**Low discrimination**					**064440**
					YEP	
Staphylococcus kloosii		89.52	0.52	MAN 99	99	
Staphylococcus cohnii subsp. cohnii		10.48	0.21		12	
064442	**Very good identification**					**064442**
Staphylococcus kloosii		100.0	0.53	MAN 99		
064443	**Good identification**					**064443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	URE 99		
064450	**Very good identification**					**064450**
					YEP	
Staphylococcus kloosii		99.85	0.93		99	
Staphylococcus cohnii subsp. cohnii		0.15	0.23		12	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
064451	**Low discrimination**						**064451**
					YEP		
Staphylococcus kloosii		51.53	0.52	MNS 1	99		
Staphylococcus cohnii subsp. urealyticum		40.16	0.38		64		
Staphylococcus cohnii subsp. cohnii		8.32	0.23		12		
064452	**Excellent identification**						**064452**
Staphylococcus kloosii		99.93	0.94				
Staphylococcus cohnii subsp. urealyticum		0.07	0.18				
064453	**Low discrimination**						**064453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		87.56	0.59	URE 99	99	64	
Staphylococcus kloosii		12.04	0.53		99	99	
Aerococcus viridans		0.40	0.22		1	1	
064460	**Good identification**						**064460**
Staphylococcus kloosii		100.0	0.40	MAN 99			
064462	**Good identification**						**064462**
Staphylococcus kloosii		100.0	0.41	MAN 99			
064470	**Very good identification**						**064470**
Staphylococcus kloosii		100.0	0.81				
064471	**Good identification**						**064471**
Staphylococcus kloosii		100.0	0.40	MNS 1			
064472	**Very good identification**						**064472**
Staphylococcus kloosii		96.35	0.82				
Staphylococcus arlettae		3.65	0.41				
064473	**Low discrimination**						**064473**
Staphylococcus arlettae		77.68	0.43	SUC 99			
Staphylococcus kloosii		15.63	0.41				
Staphylococcus cohnii subsp. urealyticum		4.59	0.18				
064500	**Good identification**						**064500**
Staphylococcus kloosii		100.0	0.32	MAN 99			
064502	**Good identification**						**064502**
Staphylococcus kloosii		100.0	0.33	MAN 99			
064510	**Very good identification**						**064510**
Staphylococcus kloosii		100.0	0.73	MLT 95			
064511	**Low discrimination**						**064511**
Staphylococcus kloosii		63.42	0.32	MLT 95			
Staphylococcus cohnii subsp. urealyticum		36.59	0.15	MNS 1			
064512	**Very good identification**						**064512**
Staphylococcus kloosii		100.0	0.74	MLT 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
064513	**Low discrimination**					**064513**
Staphylococcus cohnii subsp. urealyticum		84.34	0.36	URE 99		
Staphylococcus kloosii		15.66	0.33			
064530	**Very good identification**					**064530**
Staphylococcus kloosii		100.0	0.61	MLT 95		
064532	**Very good identification**					**064532**
Staphylococcus kloosii		97.92	0.61	MLT 95		
Staphylococcus arlettae		2.08	0.15			
064540	**Very good identification**					**064540**
Staphylococcus kloosii		100.0	0.58	MAN 99		
064542	**Very good identification**					**064542**
Staphylococcus kloosii		100.0	0.59	MAN 99		
064543	**Acceptable identification**					**064543**
Staphylococcus cohnii subsp. urealyticum		94.02	0.31	URE 99		
Staphylococcus kloosii		5.98	0.18	MAN 91		
064550	**Excellent identification**					**064550**
Staphylococcus kloosii		100.0	0.99			
064551	**Low discrimination**					**064551**
					YEP	
Staphylococcus kloosii		84.65	0.58	MNS 1	99	
Staphylococcus cohnii subsp. urealyticum		14.57	0.31		64	
Staphylococcus cohnii subsp. cohnii		0.78	0.04		12	
064552	**Excellent identification**					**064552**
Staphylococcus kloosii		99.96	1.00			
Staphylococcus arlettae		0.03	0.15			
Staphylococcus cohnii subsp. urealyticum		0.02	0.10			
064553	**Low discrimination**					**064553**
Staphylococcus cohnii subsp. urealyticum		60.38	0.51	URE 99		
Staphylococcus kloosii		37.60	0.59			
Staphylococcus arlettae		1.38	0.18			
064560	**Good identification**					**064560**
Staphylococcus kloosii		100.0	0.46	MAN 99		
064562	**Acceptable identification**					**064562**
Staphylococcus kloosii		90.04	0.47	MAN 99		
Staphylococcus arlettae		9.96	0.15			
064570	**Very good identification**					**064570**
Staphylococcus kloosii		99.88	0.87			
Staphylococcus arlettae		0.12	0.15			
064571	**Low discrimination**					**064571**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		86.30	0.46	MNS 1		
Staphylococcus arlettae		13.70	0.18			
064572	**Good identification**					**064572**
Staphylococcus kloosii		90.04	0.88			
Staphylococcus arlettae		9.96	0.56			
064573	**Good identification**					**064573**
Staphylococcus arlettae		92.30	0.59	SUC 99		
Staphylococcus kloosii		6.36	0.47			
Staphylococcus xylosus		0.49	0.03			
064610	**Good identification**					**064610**
Staphylococcus kloosii		100.0	0.46	SUC 9		
064612	**Good identification**					**064612**
Staphylococcus kloosii		100.0	0.47	SUC 9		
064630	**Good identification**					**064630**
Staphylococcus kloosii		100.0	0.34	SUC 9		
064632	**Low discrimination**					**064632**
Staphylococcus arlettae		87.92	0.41	MLT 99		
Staphylococcus kloosii		12.08	0.35			
064633	**Good identification**					**064633**
Staphylococcus arlettae		100.0	0.43	MLT 99		
064640	**Good identification**					**064640**
Staphylococcus kloosii		100.0	0.32	SUC 9		
064642	**Good identification**					**064642**
Staphylococcus kloosii		100.0	0.32	SUC 9		
064650	**Very good identification**					**064650**
Staphylococcus kloosii		100.0	0.73	SUC 9		
064651	**Low discrimination**					**064651**
Staphylococcus kloosii		80.41	0.32	SUC 9		
Staphylococcus cohnii subsp. urealyticum		19.60	0.07	MNS 1		
064652	**Good identification**					**064652**
Staphylococcus kloosii		91.26	0.73	SUC 9		
Staphylococcus arlettae		8.74	0.41			
064653	**Low discrimination**					**064653**
Staphylococcus arlettae		77.66	0.43	XYL 99		
Staphylococcus cohnii subsp. urealyticum		14.06	0.28			
Staphylococcus kloosii		6.18	0.32			
064662	**Good identification**					**064662**
Staphylococcus arlettae		97.43	0.41	MAN 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		2.57	0.20				
064663	**Good identification**						**064663**
Staphylococcus arlettae		100.0	0.43	MAN 99			
064670	**Low discrimination**						**064670**
Staphylococcus kloosii		70.68	0.60	SUC 9			
Staphylococcus arlettae		29.33	0.41				
064671	**Good identification**						**064671**
Staphylococcus arlettae		96.33	0.43	LAC 99			
Staphylococcus xylosus		1.90	-.01				
Staphylococcus kloosii		1.77	0.19				
064672	**Very good identification**						**064672**
Staphylococcus arlettae		97.43	0.82				
Staphylococcus kloosii		2.57	0.61				
064673	**Very good identification**						**064673**
					NOV	YEP	
Staphylococcus arlettae		99.90	0.85		99	99	
Staphylococcus xylosus		0.06	0.08		99	40	
Aerococcus viridans		0.03	0.22		1	1	
064710	**Good identification**						**064710**
Staphylococcus kloosii		100.0	0.52	SUC 9			
064712	**Good identification**						**064712**
Staphylococcus kloosii		94.91	0.53	SUC 9			
Staphylococcus arlettae		5.09	0.15	MLT 95			
064730	**Low discrimination**						**064730**
Staphylococcus kloosii		81.14	0.40	SUC 9			
Staphylococcus arlettae		18.86	0.15	MLT 95			
064732	**Very good identification**						**064732**
Staphylococcus arlettae		95.50	0.56	MLT 99			
Staphylococcus kloosii		4.50	0.41				
064733	**Very good identification**						**064733**
Staphylococcus arlettae		99.25	0.59	MLT 99			
Staphylococcus xylosus		0.75	0.06				
064740	**Good identification**						**064740**
Staphylococcus kloosii		100.0	0.38	SUC 9			
064742	**Low discrimination**						**064742**
Staphylococcus kloosii		78.16	0.38	SUC 9			
Staphylococcus arlettae		21.85	0.15	MAN 99			
064750	**Very good identification**						**064750**
Staphylococcus kloosii		99.70	0.79	SUC 9			
Staphylococcus arlettae		0.31	0.15				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
064751	**Low discrimination**						**064751**
					COA	NOV	
Staphylococcus kloosii		52.34	0.38	SUC 9	1	99	
Staphylococcus arlettae		21.01	0.18	MNS 1	1	99	
Staphylococcus intermedius		20.28	0.10		99	1	
064752	**Low discrimination**						**064752**
Staphylococcus kloosii		78.16	0.79	SUC 9			
Staphylococcus arlettae		21.85	0.56				
064753	**Low discrimination**						**064753**
					COA	NOV	
Staphylococcus arlettae		89.74	0.59	XYL 99	1	99	
Staphylococcus intermedius		5.38	0.26		99	1	
Staphylococcus kloosii		2.45	0.38		1	99	
064762	**Very good identification**						**064762**
Staphylococcus arlettae		99.11	0.56	MAN 99			
Staphylococcus kloosii		0.90	0.26				
064763	**Very good identification**						**064763**
Staphylococcus arlettae		99.36	0.59	MAN 99			
Staphylococcus xylosus		0.64	0.04				
064770	**Low discrimination**						**064770**
Staphylococcus arlettae		54.78	0.56	LAC 99			
Staphylococcus kloosii		45.22	0.66				
064771	**Very good identification**						**064771**
Staphylococcus arlettae		96.01	0.59	LAC 99			
Staphylococcus xylosus		3.38	0.20				
Staphylococcus kloosii		0.60	0.25				
064772	**Excellent identification**						**064772**
Staphylococcus arlettae		99.11	0.98				
Staphylococcus kloosii		0.90	0.67				
064773	**Excellent identification**						**064773**
Staphylococcus arlettae		99.89	1.00				
Staphylococcus xylosus		0.10	0.29				
Staphylococcus kloosii		0.01	0.26				
065012	**Good identification**						**065012**
Staphylococcus kloosii		100.0	0.27	TRE 99			
065050	**Good identification**						**065050**
Staphylococcus kloosii		100.0	0.52	TRE 99			
065052	**Very good identification**						**065052**
Staphylococcus kloosii		100.0	0.53	TRE 99			
065070	**Good identification**						**065070**
Staphylococcus kloosii		100.0	0.40	TRE 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
065072	**Good identification**					**065072**
Staphylococcus kloosii		100.0	0.41	TRE 99		
065110	**Good identification**					**065110**
Staphylococcus kloosii		100.0	0.32	TRE 99		
065112	**Good identification**					**065112**
Staphylococcus kloosii		100.0	0.33	TRE 99		
065150	**Very good identification**					**065150**
Staphylococcus kloosii		100.0	0.58	TRE 99		
065152	**Very good identification**					**065152**
Staphylococcus kloosii		100.0	0.59	TRE 99		
065170	**Good identification**					**065170**
Staphylococcus kloosii		100.0	0.46	TRE 99		
065172	**Acceptable identification**					**065172**
Staphylococcus kloosii		90.04	0.47	TRE 99		
Staphylococcus arlettae		9.96	0.15			
065250	**Good identification**					**065250**
Staphylococcus kloosii		100.0	0.32	SUC 9		
065252	**Good identification**					**065252**
Staphylococcus kloosii		100.0	0.32	SUC 9		
065272	**Good identification**					**065272**
Staphylococcus arlettae		97.43	0.41	TRE 99		
Staphylococcus kloosii		2.57	0.20			
065273	**Good identification**					**065273**
Staphylococcus arlettae		100.0	0.43	TRE 99		
065350	**Good identification**					**065350**
Staphylococcus kloosii		100.0	0.38	SUC 9		
065352	**Low discrimination**					**065352**
Staphylococcus kloosii		78.16	0.38	SUC 9		
Staphylococcus arlettae		21.85	0.15	TRE 99		
065372	**Very good identification**					**065372**
Staphylococcus arlettae		99.11	0.56	TRE 99		
Staphylococcus kloosii		0.90	0.26			
065373	**Very good identification**					**065373**
Staphylococcus arlettae		100.0	0.59	TRE 99		
065402	**Good identification**					**065402**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.27	MAN 99			
065410	**Very good identification**						**065410**
Staphylococcus kloosii		100.0	0.67	MLT 95			
065412	**Very good identification**						**065412**
					NOV	YEP	
Staphylococcus kloosii		99.38	0.68	MLT 95	99	99	
Aerococcus viridans		0.62	0.22		1	1	
065413	**Low discrimination**						**065413**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		79.57	0.35	URE 99	99	64	
Aerococcus viridans		12.04	0.29		1	1	
Staphylococcus kloosii		8.39	0.27		99	99	
065430	**Very good identification**						**065430**
Staphylococcus kloosii		100.0	0.55	MLT 95			
065432	**Very good identification**						**065432**
					NOV	YEP	
Staphylococcus kloosii		97.58	0.55	MLT 95	99	99	
Aerococcus viridans		2.42	0.22		1	1	
065433	**Good identification**						**065433**
Aerococcus viridans		100.0	0.29	PHS 1			
065440	**Good identification**						**065440**
Staphylococcus kloosii		100.0	0.52	MAN 99			
065442	**Very good identification**						**065442**
					NOV	YEP	
Staphylococcus kloosii		95.86	0.53	MAN 99	99	99	
Aerococcus viridans		4.14	0.24		1	1	
065443	**Low discrimination**						**065443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		73.54	0.30	URE 99	99		
Aerococcus viridans		26.46	0.32	MAN 91	1		
065450	**Very good identification**						**065450**
Staphylococcus kloosii		100.0	0.93				
065451	**Low discrimination**						**065451**
					NOV	YEP	
Staphylococcus kloosii		74.66	0.52	MNS 1	99	99	
Staphylococcus cohnii subsp. urealyticum		22.63	0.30		99	64	
Aerococcus viridans		2.71	0.22		1	1	
065452	**Excellent identification**						**065452**
					NOV	YEP	
Staphylococcus kloosii		99.84	0.94		99	99	
Aerococcus viridans		0.13	0.34		1	1	
Staphylococcus cohnii subsp. urealyticum		0.03	0.09		99	64	
065453	**Low discrimination**						**065453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		68.48	0.51	URE 99	99	64	
Staphylococcus kloosii		24.21	0.53		99	99	
Aerococcus viridans		7.31	0.42		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
065460	**Good identification**						**065460**
Staphylococcus kloosii		100.0	0.40	MAN 99			
065462	**Low discrimination**						**065462**
					NOV	YEP	
Staphylococcus kloosii		85.28	0.41	MAN 99	99	99	
Aerococcus viridans		14.72	0.24		1	1	
065463	**Good identification**						**065463**
Aerococcus viridans		100.0	0.32	PHS 1			
065470	**Very good identification**						**065470**
Staphylococcus kloosii		100.0	0.81				
065471	**Low discrimination**						**065471**
					NOV	YEP	
Staphylococcus kloosii		87.31	0.40	MNS 1	99	99	
Aerococcus viridans		12.70	0.22		1	1	
065472	**Very good identification**						**065472**
					NOV	YEP	
Staphylococcus kloosii		95.87	0.82		99	99	
Staphylococcus arlettae		3.63	0.41		99	99	
Aerococcus viridans		0.50	0.34		1	1	
065473	**Low discrimination**						**065473**
					NOV	YEP	
Staphylococcus arlettae		68.16	0.43	SUC 99	99	99	
Aerococcus viridans		16.56	0.42		1	1	
Staphylococcus kloosii		13.71	0.41		99	99	
065500	**Good identification**						**065500**
Staphylococcus kloosii		100.0	0.32	MAN 99			
065502	**Good identification**						**065502**
Staphylococcus kloosii		100.0	0.33	MAN 99			
065510	**Very good identification**						**065510**
Staphylococcus kloosii		100.0	0.73	MLT 95			
065511	**Low discrimination**						**065511**
Staphylococcus kloosii		81.68	0.32	MLT 95			
Staphylococcus cohnii subsp. urealyticum		18.33	0.07	MNS 1			
065512	**Very good identification**						**065512**
					NOV	YEP	
Staphylococcus kloosii		99.68	0.74	MLT 95	99	99	
Aerococcus viridans		0.32	0.22		1	1	
065513	**Low discrimination**						**065513**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		54.63	0.27	URE 99	99	64	
Staphylococcus kloosii		26.09	0.33		99	99	
Aerococcus viridans		19.28	0.29		1	1	
065530	**Very good identification**						**065530**
Staphylococcus kloosii		100.0	0.61	MLT 95			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
065532	**Very good identification**						**065532**
					NOV	YEP	
Staphylococcus kloosii		96.71	0.61	MLT 95	99	99	
Staphylococcus arlettae		2.05	0.15		99	99	
Aerococcus viridans		1.24	0.22		1	1	
065533	**Low discrimination**						**065533**
					NOV	YEP	
Aerococcus viridans		43.85	0.29	PHS 1	1	1	
Staphylococcus arlettae		41.31	0.18		99	99	
Staphylococcus kloosii		14.83	0.20		99	99	
065540	**Very good identification**						**065540**
Staphylococcus kloosii		100.0	0.58	MAN 99			
065542	**Very good identification**						**065542**
					NOV	YEP	
Staphylococcus kloosii		97.83	0.59	MAN 99	99	99	
Aerococcus viridans		2.17	0.24		1	1	
065550	**Excellent identification**						**065550**
Staphylococcus kloosii		100.0	0.99				
065551	**Good identification**						**065551**
					NOV	YEP	
Staphylococcus kloosii		92.11	0.58	MNS 1	99	99	
Staphylococcus cohnii subsp. urealyticum		6.16	0.22		99	64	
Aerococcus viridans		1.73	0.22		1	1	
065552	**Excellent identification**						**065552**
					NOV	YEP	
Staphylococcus kloosii		99.91	1.00		99	99	
Aerococcus viridans		0.07	0.34		1	1	
Staphylococcus arlettae		0.03	0.15		99	99	
065553	**Low discrimination**						**065553**
					NOV	YEP	
Staphylococcus kloosii		55.04	0.59	MNS 1	99	99	
Staphylococcus cohnii subsp. urealyticum		34.38	0.43		99	64	
Aerococcus viridans		8.56	0.42		1	1	
065560	**Good identification**						**065560**
Staphylococcus kloosii		100.0	0.46	MAN 99			
065562	**Low discrimination**						**065562**
					NOV	YEP	
Staphylococcus kloosii		83.37	0.47	MAN 99	99	99	
Staphylococcus arlettae		9.22	0.15		99	99	
Aerococcus viridans		7.41	0.24		1	1	
065563	**Low discrimination**						**065563**
					NOV	YEP	
Aerococcus viridans		58.60	0.32	PHS 1	1	1	
Staphylococcus arlettae		41.40	0.18		99	99	
065570	**Very good identification**						**065570**
Staphylococcus kloosii		99.88	0.87				
Staphylococcus arlettae		0.12	0.15				
065571	**Low discrimination**						**065571**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus kloosii		81.06	0.46	MNS 1	99	99		
Staphylococcus arlettae		12.87	0.18		99	99		
Aerococcus viridans		6.07	0.22		1	1		
065572	**Low discrimination**							**065572**
					NOV	YEP		
Staphylococcus kloosii		89.83	0.88		99	99		
Staphylococcus arlettae		9.93	0.56		99	99		
Aerococcus viridans		0.24	0.34		1	1		
065573	**Low discrimination**							**065573**
					NOV	YEP		
Staphylococcus arlettae		89.94	0.59	SUC 99	99	99		
Staphylococcus kloosii		6.20	0.47		99	99		
Aerococcus viridans		3.86	0.42		1	1		
065610	**Good identification**							**065610**
Staphylococcus kloosii		100.0	0.46	SUC 9				
065612	**Acceptable identification**							**065612**
					NOV	YEP		
Staphylococcus kloosii		94.09	0.47	SUC 9	99	99		
Aerococcus viridans		5.91	0.22	MLT 95	1	1		
065613	**Good identification**							**065613**
Aerococcus viridans		100.0	0.29	PHS 1				
065630	**Good identification**							**065630**
Staphylococcus kloosii		100.0	0.34	SUC 9				
065632	**Low discrimination**							**065632**
					NOV	YEP		
Staphylococcus arlettae		85.33	0.41	MLT 99	99	99		
Staphylococcus kloosii		11.73	0.35		99	99		
Aerococcus viridans		2.95	0.22		1	1		
065633	**Acceptable identification**							**065633**
					NOV	YEP		
Staphylococcus arlettae		94.27	0.43	MLT 99	99	99		
Aerococcus viridans		5.73	0.29		1	1		
065640	**Good identification**							**065640**
Staphylococcus kloosii		100.0	0.32	SUC 9				
065642	**Low discrimination**							**065642**
					NOV	YEP		
Staphylococcus kloosii		69.63	0.32	SUC 9	99	99		
Aerococcus viridans		30.37	0.24	MAN 99	1	1		
065643	**Good identification**							**065643**
Aerococcus viridans		100.0	0.32	PHS 1				
065650	**Very good identification**							**065650**
Staphylococcus kloosii		100.0	0.73	SUC 9				
065651	**Low discrimination**							**065651**
					OXI	NOV	YEP	
Staphylococcus kloosii		62.45	0.32	SUC 9	1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		22.96	0.22	MNS 1	1	1	1	
Staphylococcus sciuri		14.59	0.02		99	99	57	
065652	**Good identification**							**065652**
					NOV	YEP		
Staphylococcus kloosii		90.17	0.73	SUC 9	99	99		
Staphylococcus arlettae		8.63	0.41		99	99		
Aerococcus viridans		1.19	0.34		1	1		
065653	**Low discrimination**							**065653**
					NOV	YEP		
Staphylococcus arlettae		71.79	0.43	XYL 99	99	99		
Aerococcus viridans		17.45	0.42		1	1		
Staphylococcus kloosii		5.71	0.32		99	99		
065662	**Acceptable identification**							**065662**
					NOV	YEP		
Staphylococcus arlettae		93.25	0.41	MAN 99	99	99		
Aerococcus viridans		4.29	0.24		1	1		
Staphylococcus kloosii		2.46	0.20		99	99		
065663	**Acceptable identification**							**065663**
					NOV	YEP		
Staphylococcus arlettae		92.51	0.43	MAN 99	99	99		
Aerococcus viridans		7.49	0.32		1	1		
065670	**Low discrimination**							**065670**
Staphylococcus kloosii		70.68	0.60	SUC 9				
Staphylococcus arlettae		29.33	0.41					
065671	**Acceptable identification**							**065671**
					NOV	YEP	PYR	
Staphylococcus arlettae		92.29	0.43	LAC 99	99	99	1	
Staphylococcus equorum		3.52	0.06		99	1	5	
Aerococcus viridans		2.49	0.22		1	1	95	
065672	**Very good identification**							**065672**
					NOV	YEP		
Staphylococcus arlettae		97.30	0.82		99	99		
Staphylococcus kloosii		2.57	0.61		99	99		
Aerococcus viridans		0.14	0.34		1	1		
065673	**Very good identification**							**065673**
					NOV	YEP	PYR	
Staphylococcus arlettae		99.62	0.85		99	99	1	
Aerococcus viridans		0.25	0.42		1	1	95	
Staphylococcus equorum		0.10	0.15		99	1	5	
065710	**Good identification**							**065710**
Staphylococcus kloosii		100.0	0.52	SUC 9				
065712	**Good identification**							**065712**
					NOV	YEP		
Staphylococcus kloosii		92.08	0.53	SUC 9	99	99		
Staphylococcus arlettae		4.94	0.15	MLT 95	99	99		
Aerococcus viridans		2.98	0.22		1	1		
065713	**Low discrimination**							**065713**
					NOV	YEP		
Aerococcus viridans		51.49	0.29	PHS 1	1	1		
Staphylococcus arlettae		48.51	0.18		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
065730	**Low discrimination**							**065730**
Staphylococcus kloosii		81.14	0.40	SUC 9				
Staphylococcus arlettae		18.86	0.15	MLT 95				
065732	**Good identification**							**065732**
					NOV	YEP		
Staphylococcus arlettae		94.95	0.56	MLT 99	99	99		
Staphylococcus kloosii		4.47	0.41		99	99		
Aerococcus viridans		0.58	0.22		1	1		
065733	**Very good identification**							**065733**
					NOV	YEP	PYR	
Staphylococcus arlettae		97.27	0.59	MLT 99	99	99	1	
Staphylococcus equorum		1.69	0.15		99	1	5	
Aerococcus viridans		1.04	0.29		1	1	95	
065740	**Good identification**							**065740**
Staphylococcus kloosii		100.0	0.38	SUC 9				
065742	**Low discrimination**							**065742**
					NOV	YEP		
Staphylococcus kloosii		66.48	0.38	SUC 9	99	99		
Staphylococcus arlettae		18.58	0.15	MAN 99	99	99		
Aerococcus viridans		14.94	0.24		1	1		
065743	**Low discrimination**							**065743**
					NOV	YEP		
Aerococcus viridans		58.60	0.32	PHS 1	1	1		
Staphylococcus arlettae		41.40	0.18		99	99		
065750	**Very good identification**							**065750**
Staphylococcus kloosii		99.70	0.79	SUC 9				
Staphylococcus arlettae		0.31	0.15					
065751	**Low discrimination**							**065751**
					OXI			
Staphylococcus kloosii		54.81	0.38	SUC 9	1			
Staphylococcus arlettae		22.00	0.18	MNS 1	1			
Staphylococcus sciuri		12.81	0.08		99			
065752	**Low discrimination**							**065752**
					NOV	YEP		
Staphylococcus kloosii		77.74	0.79	SUC 9	99	99		
Staphylococcus arlettae		21.73	0.56		99	99		
Aerococcus viridans		0.53	0.34		1	1		
065753	**Good identification**							**065753**
					NOV	YEP		
Staphylococcus arlettae		92.67	0.59	XYL 99	99	99		
Aerococcus viridans		3.97	0.42		1	1		
Staphylococcus kloosii		2.53	0.38		99	99		
065762	**Very good identification**							**065762**
					NOV	YEP		
Staphylococcus arlettae		98.31	0.56	MAN 99	99	99		
Staphylococcus kloosii		0.89	0.26		99	99		
Aerococcus viridans		0.80	0.24		1	1		
065763	**Very good identification**							**065763**
					NOV	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		96.52	0.59	MAN 99	99	1	
Staphylococcus gallinarum		2.11	0.14		99	12	
Aerococcus viridans		1.38	0.32		1	95	
065770	**Low discrimination**						**065770**
Staphylococcus arlettae		54.78	0.56	LAC 99			
Staphylococcus kloosii		45.22	0.66				
065771	**Low discrimination**						**065771**
					YEP		
Staphylococcus arlettae		84.57	0.59	LAC 99	99		
Staphylococcus gallinarum		7.62	0.27		78		
Staphylococcus equorum		5.76	0.27		1		
065772	**Excellent identification**						**065772**
					NOV	YEP	
Staphylococcus arlettae		99.08	0.98		99	99	
Staphylococcus kloosii		0.90	0.67		99	99	
Aerococcus viridans		0.02	0.34		1	1	
065773	**Excellent identification**						**065773**
					YEP		
Staphylococcus arlettae		99.52	1.00		99		
Staphylococcus gallinarum		0.22	0.35		78		
Staphylococcus equorum		0.18	0.35		1		
066303	**Acceptable identification**						**066303**
					PYR		
Staphylococcus hyicus		90.39	0.28	ARG 96	1		
Staphylococcus intermedius		9.61	0.14	BGA 10	90		
066412	**Good identification**						**066412**
Staphylococcus kloosii		100.0	0.27	NIT 1			
066413	**Low discrimination**						**066413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		72.35	0.28	URE 99	99		
Staphylococcus carnosus subsp. carnosus		17.08	0.07		1		
Staphylococcus simulans		10.57	0.05		1		
066450	**Acceptable identification**						**066450**
Staphylococcus kloosii		94.59	0.52	NIT 1			
Staphylococcus saprophyticus subsp. bovis		5.41	0.09				
066452	**Very good identification**						**066452**
Staphylococcus kloosii		100.0	0.53	NIT 1			
066453	**Good identification**						**066453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	URE 99			
066470	**Good identification**						**066470**
Staphylococcus kloosii		100.0	0.40	NIT 1			
066472	**Good identification**						**066472**
Staphylococcus kloosii		100.0	0.41	NIT 1			
066510	**Good identification**						**066510**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.32	NIT 1			
066512	**Good identification**						**066512**
Staphylococcus kloosii		100.0	0.33	NIT 1			
066550	**Low discrimination**						**066550**
Staphylococcus kloosii		89.45	0.58	NIT 1			
Staphylococcus saprophyticus subsp. bovis		10.55	0.22				
066552	**Very good identification**						**066552**
Staphylococcus kloosii		97.35	0.59	NIT 1			
Staphylococcus saprophyticus subsp. bovis		2.65	0.09				
066553	**Low discrimination**						**066553**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		84.31	0.36	URE 99	1	99	
Staphylococcus intermedius		8.68	0.08		99	1	
Staphylococcus xylosus		4.01	-.01		1	99	
066570	**Good identification**						**066570**
Staphylococcus kloosii		100.0	0.46	NIT 1			
066572	**Acceptable identification**						**066572**
Staphylococcus kloosii		90.04	0.47	NIT 1			
Staphylococcus arlettae		9.96	0.15				
066603	**Low discrimination**						**066603**
Staphylococcus hyicus		83.13	0.28	ARG 96			
Staphylococcus simulans		16.87	0.21	BGA 10			
066613	**Good identification**						**066613**
					COA		
Staphylococcus simulans		95.46	0.34	URE 98	1		
Staphylococcus intermedius		4.54	0.05	ARG 92	99		
066620	**Good identification**						**066620**
Staphylococcus muscae		100.0	0.39	BGA 1			
066672	**Good identification**						**066672**
Staphylococcus arlettae		97.43	0.41	NIT 1			
Staphylococcus kloosii		2.57	0.20				
066673	**Low discrimination**						**066673**
Staphylococcus arlettae		68.74	0.43	NIT 1			
Staphylococcus xylosus		31.26	0.27				
066701	**Low discrimination**						**066701**
					PYR		
Staphylococcus hyicus		81.31	0.28	ARG 96	1		
Staphylococcus intermedius		18.70	0.21	BGA 10	90		
066703	**Low discrimination**						**066703**
					COA	PYR	
Staphylococcus hyicus		88.94	0.50	ARG 96	24	1	
Staphylococcus intermedius		10.92	0.37	BGA 10	99	90	
Staphylococcus simulans		0.14	-.01		1	70	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
066711	**Good identification**							**066711**
Staphylococcus intermedius		100.0	0.29	URE 90				
066713	**Acceptable identification**							**066713**
					COA	PYR		
Staphylococcus intermedius		93.68	0.46	URE 90	99	90		
Staphylococcus hyicus		2.85	0.09	GLR 3	24	1		
Staphylococcus simulans		1.73	0.12		1	70		
066743	**Acceptable identification**							**066743**
					NOV	PYR	COA	
Staphylococcus intermedius		93.45	0.40	URE 90	1	90	99	
Staphylococcus hyicus		5.12	0.09	GLR 3	1	1	24	
Staphylococcus xylosus		1.43	0.01		99	65	1	
066750	**Low discrimination**							**066750**
Staphylococcus saprophyticus subsp. bovis		82.67	0.34	URE 99				
Staphylococcus kloosii		17.33	0.38					
066751	**Low discrimination**							**066751**
					COA	NOV		
Staphylococcus intermedius		69.18	0.33	URE 90	99	1		
Staphylococcus saprophyticus subsp. bovis		16.02	0.22	GLR 3	1	99		
Staphylococcus xylosus		13.24	0.16	LAC 86	1	99		
066753	**Low discrimination**							**066753**
					COA	NOV		
Staphylococcus intermedius		88.97	0.49	URE 90	99	1		
Staphylococcus xylosus		7.89	0.26	GLR 3	1	99		
Staphylococcus arlettae		1.13	0.18		1	99		
066771	**Low discrimination**							**066771**
Staphylococcus xylosus		72.51	0.38	URE 99				
Staphylococcus gallinarum		22.91	0.27					
Staphylococcus arlettae		2.57	0.18					
066772	**Very good identification**							**066772**
Staphylococcus arlettae		98.05	0.56	NIT 1				
Staphylococcus xylosus		1.07	0.07					
Staphylococcus kloosii		0.89	0.26					
066773	**Low discrimination**							**066773**
Staphylococcus arlettae		47.35	0.59	NIT 1				
Staphylococcus xylosus		38.42	0.48					
Staphylococcus gallinarum		10.44	0.35					
067371	**Low discrimination**							**067371**
					OXI			
Staphylococcus gallinarum		83.65	0.27	URE 92	1			
Staphylococcus equorum		7.35	0.08	TRE 99	1			
Staphylococcus sciuri		6.24	0.09		99			
067373	**Low discrimination**							**067373**
					YEP			
Staphylococcus gallinarum		83.82	0.35	URE 92	78			
Staphylococcus equorum		7.73	0.16	TRE 99	1			
Staphylococcus arlettae		3.84	0.18		99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
067412	**Good identification**						**067412**
Staphylococcus kloosii		100.0	0.27	NIT 1			
067450	**Good identification**						**067450**
Staphylococcus kloosii		100.0	0.52	NIT 1			
067452	**Very good identification**						**067452**
Staphylococcus kloosii		100.0	0.53	NIT 1			
067453	**Good identification**						**067453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	URE 99			
067470	**Good identification**						**067470**
Staphylococcus kloosii		100.0	0.40	NIT 1			
067472	**Good identification**						**067472**
Staphylococcus kloosii		100.0	0.41	NIT 1			
067510	**Good identification**						**067510**
Staphylococcus kloosii		100.0	0.32	NIT 1			
067512	**Good identification**						**067512**
Staphylococcus kloosii		100.0	0.33	NIT 1			
067550	**Very good identification**						**067550**
Staphylococcus kloosii		100.0	0.58	NIT 1			
067552	**Very good identification**						**067552**
Staphylococcus kloosii		100.0	0.59	NIT 1			
067553	**Low discrimination**						**067553**
					OXI	YEP	
Staphylococcus cohnii subsp. urealyticum		82.47	0.27	URE 99	1	64	
Staphylococcus lentus		9.97	0.05		99	1	
Staphylococcus kloosii		7.56	0.18		1	99	
067570	**Good identification**						**067570**
Staphylococcus kloosii		100.0	0.46	NIT 1			
067571	**Low discrimination**						**067571**
Staphylococcus gallinarum		88.54	0.27	URE 92			
Staphylococcus equorum		7.78	0.08	SUC 99			
Staphylococcus xylosus		3.69	-.01				
067572	**Acceptable identification**						**067572**
Staphylococcus kloosii		90.04	0.47	NIT 1			
Staphylococcus arlettae		9.96	0.15				
067573	**Low discrimination**						**067573**
Staphylococcus gallinarum		83.12	0.35	URE 92			
Staphylococcus equorum		7.67	0.16	SUC 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		4.03	0.09				
067651	**Good identification**						**067651**
Staphylococcus sciuri		100.0	0.37	BGA 1			
067652	**Low discrimination**						**067652**
					OXI		
Staphylococcus kloosii		82.47	0.32	NIT 1	1		
Staphylococcus sciuri		17.53	0.02	SUC 9	99		
067671	**Low discrimination**						**067671**
					OXI		
Staphylococcus sciuri		39.07	0.34	BGA 1	99		
Staphylococcus gallinarum		31.44	0.27	GLR 14	1		
Staphylococcus equorum		27.03	0.28		1		
067672	**Good identification**						**067672**
Staphylococcus arlettae		97.43	0.41	NIT 1			
Staphylococcus kloosii		2.57	0.20				
067673	**Low discrimination**						**067673**
					YEP		
Staphylococcus gallinarum		34.56	0.35	URE 92	78		
Staphylococcus equorum		31.21	0.36	GAL 99	1		
Staphylococcus arlettae		27.66	0.43		99		
067711	**Low discrimination**						**067711**
Staphylococcus sciuri		60.26	0.31	BGA 1			
Staphylococcus lentus		39.74	0.29	GLR 14			
067713	**Acceptable identification**						**067713**
					OXI	COA	
Staphylococcus lentus		90.95	0.40	GLR 1	99	1	
Staphylococcus sciuri		6.33	0.15		99	1	
Staphylococcus intermedius		2.72	0.05		1	99	
067731	**Low discrimination**						**067731**
					OXI		
Staphylococcus gallinarum		32.04	0.27	URE 92	1		
Staphylococcus equorum		27.55	0.28	MLT 99	1		
Staphylococcus sciuri		20.55	0.28		99		
067733	**Low discrimination**						**067733**
					OXI		
Staphylococcus gallinarum		34.81	0.35	URE 92	1		
Staphylococcus equorum		31.44	0.36	MLT 99	1		
Staphylococcus lentus		28.26	0.40		99		
067751	**Low discrimination**						**067751**
					OXI		
Staphylococcus sciuri		58.22	0.43	BGA 1	99		
Staphylococcus lentus		20.72	0.35	GLR 14	99		
Staphylococcus gallinarum		18.20	0.27		1		
067752	**Low discrimination**						**067752**
					OXI		
Staphylococcus kloosii		61.47	0.38	NIT 1	1		
Staphylococcus arlettae		17.18	0.15	SUC 9	1		
Staphylococcus sciuri		13.07	0.08		99		
067753	**Low discrimination**						**067753**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI		
Staphylococcus lentus		52.91	0.47	GLR 1	99		
Staphylococcus gallinarum		32.10	0.35		1		
Staphylococcus sciuri		6.83	0.27		99		
067761	**Good identification**						**067761**
					OXI		
Staphylococcus gallinarum		97.87	0.47	URE 92	1		
Staphylococcus equorum		0.88	0.08	MAN 91	1		
Staphylococcus sciuri		0.75	0.09		99		
067763	**Very good identification**						**067763**
Staphylococcus gallinarum		97.87	0.55	URE 92			
Staphylococcus equorum		0.92	0.16	MAN 91			
Staphylococcus xylosus		0.59	0.11				
067770	**Low discrimination**						**067770**
					OXI		
Staphylococcus gallinarum		64.91	0.27	URE 92	1		
Staphylococcus sciuri		19.36	0.22	MNS 99	99		
Staphylococcus equorum		5.70	0.08		1		
067771	**Low discrimination**						**067771**
					OXI		
Staphylococcus gallinarum		88.28	0.68	URE 92	1		
Staphylococcus equorum		7.75	0.49		1		
Staphylococcus sciuri		2.15	0.40		99		
067772	**Low discrimination**						**067772**
					YEP		
Staphylococcus arlettae		74.69	0.56	NIT 1	99		
Staphylococcus gallinarum		21.83	0.35		78		
Staphylococcus equorum		2.01	0.16		1		
067773	**Low discrimination**						**067773**
					YEP		
Staphylococcus gallinarum		85.79	0.76	URE 92	78		
Staphylococcus equorum		7.91	0.57		1		
Staphylococcus arlettae		3.93	0.59		99		
070672	**Good identification**						**070672**
Staphylococcus arlettae		100.0	0.37	ORN 1			
070673	**Good identification**						**070673**
					NOV	YEP	
Staphylococcus arlettae		96.11	0.40	ORN 1	99	99	
Aerococcus viridans		3.89	0.22		1	1	
070772	**Good identification**						**070772**
Staphylococcus arlettae		100.0	0.53	ORN 1			
070773	**Very good identification**						**070773**
					NOV	YEP	
Staphylococcus arlettae		98.76	0.55	ORN 1	99	99	
Aerococcus viridans		0.71	0.22		1	1	
Staphylococcus xylosus		0.53	-.01		99	40	
071413	**Good identification**						**071413**
Aerococcus viridans		100.0	0.29	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
071433	**Good identification**						**071433**
Aerococcus viridans		100.0	0.29	ORN 1			
071443	**Good identification**						**071443**
Aerococcus viridans		100.0	0.32	ORN 1			
071452	**Good identification**						**071452**
Aerococcus viridans		100.0	0.34	ORN 1			
071453	**Acceptable identification**						**071453**
					NOV		
Aerococcus viridans		94.07	0.42	ORN 1	1		
Staphylococcus cohnii subsp. urealyticum		5.93	0.06		99		
071463	**Good identification**						**071463**
Aerococcus viridans		100.0	0.32	ORN 1			
071472	**Good identification**						**071472**
Aerococcus viridans		100.0	0.34	ORN 1			
071473	**Good identification**						**071473**
Aerococcus viridans		100.0	0.42	ORN 1			
071513	**Good identification**						**071513**
Aerococcus viridans		100.0	0.29	ORN 1			
071533	**Good identification**						**071533**
Aerococcus viridans		100.0	0.29	ORN 1			
071543	**Good identification**						**071543**
Aerococcus viridans		100.0	0.32	ORN 1			
071552	**Low discrimination**						**071552**
					NOV	YEP	
Aerococcus viridans		86.84	0.34	ORN 1	1	1	
Staphylococcus kloosii		13.16	0.18		99	99	
071553	**Good identification**						**071553**
Aerococcus viridans		100.0	0.42	ORN 1			
071563	**Good identification**						**071563**
Aerococcus viridans		100.0	0.32	ORN 1			
071572	**Low discrimination**						**071572**
					NOV	YEP	
Aerococcus viridans		78.35	0.34	ORN 1	1	1	
Staphylococcus arlettae		21.65	0.12		99	99	
071573	**Low discrimination**						**071573**
					NOV	YEP	
Aerococcus viridans		86.43	0.42	ORN 1	1	1	
Staphylococcus arlettae		13.57	0.14		99	99	
071613	**Good identification**						**071613**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.29	ORN 1			
071633	**Good identification**						**071633**
Aerococcus viridans		100.0	0.29	ORN 1			
071643	**Good identification**						**071643**
Aerococcus viridans		100.0	0.32	ORN 1			
071652	**Good identification**						**071652**
Aerococcus viridans		100.0	0.34	ORN 1			
071653	**Good identification**						**071653**
Aerococcus viridans		100.0	0.42	ORN 1			
071663	**Good identification**						**071663**
Aerococcus viridans		100.0	0.32	ORN 1			
071672	**Low discrimination**						**071672**
					NOV	YEP	
Staphylococcus arlettae		82.84	0.37	ORN 1	99	99	
Aerococcus viridans		17.16	0.34		1	1	
071673	**Low discrimination**						**071673**
					NOV	YEP	
Staphylococcus arlettae		73.29	0.40	ORN 1	99	99	
Aerococcus viridans		26.72	0.42		1	1	
071713	**Good identification**						**071713**
Aerococcus viridans		100.0	0.29	ORN 1			
071733	**Low discrimination**						**071733**
					NOV	YEP	
Aerococcus viridans		61.42	0.29	ORN 1	1	1	
Staphylococcus arlettae		38.58	0.14		99	99	
071743	**Good identification**						**071743**
Aerococcus viridans		100.0	0.32	ORN 1			
071752	**Low discrimination**						**071752**
					NOV	YEP	
Aerococcus viridans		78.35	0.34	ORN 1	1	1	
Staphylococcus arlettae		21.65	0.12		99	99	
071753	**Low discrimination**						**071753**
					NOV	YEP	
Aerococcus viridans		86.43	0.42	ORN 1	1	1	
Staphylococcus arlettae		13.57	0.14		99	99	
071763	**Low discrimination**						**071763**
					NOV	YEP	
Aerococcus viridans		67.98	0.32	ORN 1	1	1	
Staphylococcus arlettae		32.02	0.14		99	99	
071772	**Good identification**						**071772**
					NOV	YEP	
Staphylococcus arlettae		96.47	0.53	ORN 1	99	99	
Aerococcus viridans		3.53	0.34		1	1	
071773	**Good identification**						**071773**

Code Taxon		ID score	T index	tests against	Differentiation tests NOV	 YEP	Code
Staphylococcus arlettae		93.96	0.55	ORN 1	99	99	
Aerococcus viridans		6.04	0.42		1	1	
074412	**Good identification**						**074412**
Staphylococcus kloosii		100.0	0.27	ORN 1			
074450	**Good identification**						**074450**
Staphylococcus kloosii		100.0	0.52	ORN 1			
074452	**Very good identification**						**074452**
Staphylococcus kloosii		100.0	0.53	ORN 1			
074470	**Good identification**						**074470**
Staphylococcus kloosii		100.0	0.40	ORN 1			
074472	**Good identification**						**074472**
Staphylococcus kloosii		100.0	0.41	ORN 1			
074510	**Good identification**						**074510**
Staphylococcus kloosii		100.0	0.32	ORN 1			
074512	**Good identification**						**074512**
Staphylococcus kloosii		100.0	0.33	ORN 1			
074550	**Very good identification**						**074550**
Staphylococcus kloosii		100.0	0.58	ORN 1			
074552	**Very good identification**						**074552**
Staphylococcus kloosii		100.0	0.59	ORN 1			
074570	**Good identification**						**074570**
Staphylococcus kloosii		100.0	0.46	ORN 1			
074572	**Acceptable identification**						**074572**
Staphylococcus kloosii		90.04	0.47	ORN 1			
Staphylococcus arlettae		9.96	0.15				
074650	**Good identification**						**074650**
Staphylococcus kloosii		100.0	0.32	ORN 1			
074652	**Good identification**						**074652**
Staphylococcus kloosii		100.0	0.32	ORN 1			
074672	**Good identification**						**074672**
Staphylococcus arlettae		97.43	0.41	ORN 1			
Staphylococcus kloosii		2.57	0.20				
074673	**Good identification**						**074673**
Staphylococcus arlettae		100.0	0.43	ORN 1			
074750	**Good identification**						**074750**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.38	ORN 1		
074752	**Low discrimination**					**074752**
Staphylococcus kloosii		78.16	0.38	ORN 1		
Staphylococcus arlettae		21.85	0.15	SUC 9		
074772	**Very good identification**					**074772**
Staphylococcus arlettae		99.11	0.56	ORN 1		
Staphylococcus kloosii		0.90	0.26			
074773	**Very good identification**					**074773**
Staphylococcus arlettae		100.0	0.59	ORN 1		
075412	**Good identification**					**075412**
Staphylococcus kloosii		100.0	0.27	ORN 1		
075450	**Good identification**					**075450**
Staphylococcus kloosii		100.0	0.52	ORN 1		
075452	**Very good identification**					**075452**
Staphylococcus kloosii		100.0	0.53	ORN 1		
075470	**Good identification**					**075470**
Staphylococcus kloosii		100.0	0.40	ORN 1		
075472	**Good identification**					**075472**
Staphylococcus kloosii		100.0	0.41	ORN 1		
075510	**Good identification**					**075510**
Staphylococcus kloosii		100.0	0.32	ORN 1		
075512	**Good identification**					**075512**
Staphylococcus kloosii		100.0	0.33	ORN 1		
075550	**Very good identification**					**075550**
Staphylococcus kloosii		100.0	0.58	ORN 1		
075552	**Very good identification**					**075552**
Staphylococcus kloosii		100.0	0.59	ORN 1		
075570	**Good identification**					**075570**
Staphylococcus kloosii		100.0	0.46	ORN 1		
075572	**Acceptable identification**					**075572**
Staphylococcus kloosii		90.04	0.47	ORN 1		
Staphylococcus arlettae		9.96	0.15			
075650	**Good identification**					**075650**
Staphylococcus kloosii		100.0	0.32	ORN 1		
075652	**Good identification**					**075652**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.32	ORN 1				
075672	**Good identification**							**075672**
Staphylococcus arlettae		97.43	0.41	ORN 1				
Staphylococcus kloosii		2.57	0.20					
075673	**Good identification**							**075673**
Staphylococcus arlettae		100.0	0.43	ORN 1				
075750	**Good identification**							**075750**
Staphylococcus kloosii		100.0	0.38	ORN 1				
075752	**Low discrimination**							**075752**
Staphylococcus kloosii		78.16	0.38	ORN 1				
Staphylococcus arlettae		21.85	0.15	SUC 9				
075772	**Very good identification**							**075772**
Staphylococcus arlettae		99.11	0.56	ORN 1				
Staphylococcus kloosii		0.90	0.26					
075773	**Very good identification**							**075773**
Staphylococcus arlettae		100.0	0.59	ORN 1				
077771	**Acceptable identification**							**077771**
Staphylococcus gallinarum		91.93	0.27	URE 92				
Staphylococcus equorum		8.07	0.08	ORN 1				
077773	**Low discrimination**							**077773**
					YEP			
Staphylococcus gallinarum		86.60	0.35	URE 92	78			
Staphylococcus equorum		7.99	0.16	ORN 1	1			
Staphylococcus arlettae		3.97	0.18		99			
100000	**Low discrimination**							**100000**
Micrococcus sp.		49.77	0.69	VPT 3				
Dermacoccus nishinomiyaensis		30.08	0.74					
Kocuria rosea		18.28	0.68					
100001	**Low discrimination**							**100001**
					OXI			
Staphylococcus capitis subsp. capitis		26.15	0.45	NIT 89	1			
Micrococcus sp.		25.91	0.34	MAN 93	89			
Kocuria rosea		24.54	0.42		35			
100002	**Low discrimination**							**100002**
					OXI			
Micrococcus sp.		36.35	0.28	VPT 3	89			
Kocuria varians		27.36	0.42	LAC 1	13			
Dermacoccus nishinomiyaensis		21.97	0.33		93			
100010	**Low discrimination**							**100010**
					NOV	YEP	OXI	
Kocuria rosea		43.07	0.48	NIT 90	1	1	35	
Macrococcus bovicus		23.62	0.48	MAN 10	99	99	99	
Staphylococcus cohnii subsp. cohnii		10.86	0.47		99	12	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
100011	**Very good identification**							**100011**
					NOV			
Staphylococcus capitis subsp. capitis		95.23	0.68	NIT 89	1			
Staphylococcus cohnii subsp. cohnii		3.88	0.48		99			
Kocuria rosea		0.75	0.22		1			
100013	**Good identification**							**100013**
					NOV			
Staphylococcus capitis subsp. capitis		98.03	0.33	NIT 89	1			
Staphylococcus cohnii subsp. cohnii		1.98	0.07	LAC 2	99			
100020	**Low discrimination**							**100020**
					OXI			
Kocuria rosea		85.94	0.60	NIT 90	35			
Micrococcus sp.		5.52	0.28		89			
Kocuria varians		5.17	0.44		13			
100021	**Low discrimination**							**100021**
Kocuria rosea		86.20	0.34	NIT 90				
Kocuria varians		8.57	0.22	MNS 5				
Staphylococcus capitis subsp. capitis		2.17	0.04					
100022	**Low discrimination**							**100022**
					OXI			
Kocuria varians		61.22	0.33	VPT 14	13			
Kocuria rosea		36.44	0.19		35			
Micrococcus sp.		2.34	-.13		89			
100030	**Good identification**							**100030**
					NOV	YEP	OXI	
Kocuria rosea		97.07	0.41	NIT 90	1	1	35	
Macrococcus bovicus		1.26	0.07	MAN 10	99	99	99	
Staphylococcus cohnii subsp. cohnii		0.58	0.06		99	12	1	
100031	**Low discrimination**							**100031**
					NOV			
Staphylococcus capitis subsp. capitis		72.80	0.27	NIT 89	1			
Kocuria rosea		24.24	0.14	XYL 1	1			
Staphylococcus cohnii subsp. cohnii		2.96	0.07		99			
100040	**Low discrimination**							**100040**
					OXI	NOV		
Micrococcus sp.		60.14	0.46	VPT 3	89	1		
Staphylococcus cohnii subsp. cohnii		20.44	0.55	MLT 7	1	99		
Dermacoccus nishinomiyaensis		4.88	0.33		93	80		
100041	**Low discrimination**							**100041**
					NOV	OXI		
Staphylococcus cohnii subsp. cohnii		75.29	0.56	TRE 99	99	1		
Kocuria kristinae		16.85	0.33		1	99		
Micrococcus sp.		4.17	0.11		1	89		
100042	**Low discrimination**							**100042**
					OXI			
Kocuria varians		53.40	0.29	VPT 14	13			
Micrococcus sp.		24.33	0.05		89			
Staphylococcus auricularis		14.01	0.16		1			
100050	**Low discrimination**							**100050**
					OXI	YEP		
Macrococcus bovicus		57.36	0.55	VPT 1	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		35.77	0.58		1	12		
Macrococcus equipersicus		5.59	0.36		99	99		
100051	**Good identification**							**100051**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		90.86	0.58	TRE 99	1	99	12	
Staphylococcus capitis subsp. capitis		7.12	0.27		1	1	1	
Macrococcus bovicus		1.36	0.14		99	99	99	
100060	**Low discrimination**							**100060**
					OXI			
Kocuria varians		44.35	0.31	VPT 14	13			
Kocuria rosea		33.92	0.19		35			
Micrococcus sp.		16.22	0.05		89			
100100	**Low discrimination**							**100100**
					OXI			
Dermacoccus nishinomiyaensis		67.47	0.59	VPT 5	93			
Kocuria varians		23.69	0.57		13			
Micrococcus sp.		5.92	0.28		89			
100101	**Low discrimination**							**100101**
					OXI	NOV		
Kocuria varians		51.37	0.35	VPT 14	13	1		
Dermacoccus nishinomiyaensis		17.00	0.18	MNS 8	93	80		
Staphylococcus cohnii subsp. cohnii		14.21	0.26		1	99		
100102	**Acceptable identification**							**100102**
					OXI			
Kocuria varians		90.01	0.46	VPT 14	13			
Dermacoccus nishinomiyaensis		9.18	0.18		93			
Micrococcus sp.		0.81	-.13		89			
100110	**Low discrimination**							**100110**
					OXI	YEP		
Macrococcus bovicus		70.39	0.40	VPT 1	99	99		
Dermacoccus nishinomiyaensis		8.54	0.18		93	1		
Staphylococcus cohnii subsp. cohnii		8.39	0.28		1	12		
100111	**Low discrimination**							**100111**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		67.44	0.27	NIT 89	1	1	1	
Staphylococcus cohnii subsp. cohnii		30.20	0.28	GAL 1	1	99	12	
Macrococcus bovicus		2.36	-.01		99	99	99	
100120	**Low discrimination**							**100120**
					OXI			
Kocuria varians		83.27	0.48	VPT 14	13			
Kocuria rosea		9.32	0.19		35			
Dermacoccus nishinomiyaensis		6.82	0.18		93			
100122	**Good identification**							**100122**
Kocuria varians		100.0	0.37	VPT 14				
100140	**Low discrimination**							**100140**
					NOV	OXI	YEP	
Kocuria varians		56.27	0.44	VPT 14	1	13	80	
Staphylococcus cohnii subsp. cohnii		18.03	0.36		99	1	12	
Macrococcus caseolyticus		7.51	0.14		99	99	99	
100141	**Low discrimination**							**100141**
					NOV	OXI	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		52.03	0.36	GAL 10	99	1	1		
Kocuria kristinae		34.94	0.23	TRE 99	1	99	93		
Kocuria varians		13.04	0.22		1	13	10		
100142	**Low discrimination**								**100142**
					NOV	OXI	YEP		
Kocuria varians		57.28	0.32	VPT 14	1	13	80		
Macrococcus caseolyticus		40.67	0.18		99	99	99		
Staphylococcus hominis subsp. novobiosepticus		2.05	-.07		99	1	1		
100150	**Low discrimination**								**100150**
					OXI	YEP			
Macrococcus bovicus		79.17	0.48	VPT 1	99	99			
Staphylococcus cohnii subsp. cohnii		12.80	0.38		1	12			
Macrococcus equipersicus		7.71	0.28		99	99			
100151	**Acceptable identification**								**100151**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		92.62	0.39	GAL 10	1	99	1	12	
Macrococcus bovicus		5.34	0.07	TRE 99	99	99	1	99	
Kocuria kristinae		2.04	-.06		99	1	93	15	
100160	**Good identification**								**100160**
Kocuria varians		100.0	0.34	VPT 14					
100200	**Low discrimination**								**100200**
Micrococcus sp.		52.33	0.48	VPT 3					
Kocuria rosea		21.59	0.48	SUC 9					
Dermacoccus nishinomiyaensis		20.41	0.49						
100201	**Low discrimination**								**100201**
					PYR	OXI			
Kocuria kristinae		61.14	0.54	TRE 95	93	99			
Staphylococcus capitis subsp. capitis		34.26	0.50	MLT 90	1	1			
Kocuria rosea		1.92	0.22		1	35			
100203	**Good identification**								**100203**
					OXI	PYR			
Kocuria kristinae		95.14	0.42	TRE 95	99	93			
Staphylococcus capitis subsp. capitis		4.35	0.15	MLT 90	1	1			
Kocuria varians		0.50	0.03		13	10			
100210	**Low discrimination**								**100210**
					NOV	YEP			
Macrococcus bovicus		28.91	0.35	VPT 1	99	99			
Staphylococcus capitis subsp. capitis		26.53	0.32		1	1			
Kocuria rosea		23.43	0.29		1	1			
100211	**Very good identification**								**100211**
					PYR	OXI			
Staphylococcus capitis subsp. capitis		99.38	0.73	NIT 89	1	1			
Kocuria kristinae		0.56	0.26		93	99			
Kocuria rosea		0.05	0.02		1	35			
100213	**Acceptable identification**								**100213**
					OXI	PYR			
Staphylococcus capitis subsp. capitis		93.59	0.39	NIT 89	1	1			
Kocuria kristinae		6.42	0.13	LAC 2	99	93			
100220	**Low discrimination**								**100220**
					OXI				

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria rosea		86.71	0.41	NIT 90	35			
Kocuria varians		6.40	0.26	SUC 10	13			
Micrococcus sp.		4.95	0.07		89			
100230	**Low discrimination**							**100230**
					NOV			
Staphylococcus vitulinus		74.45	0.29	VPT 1	99			
Kocuria rosea		25.55	0.21	NIT 91	1			
100231	**Good identification**							**100231**
Staphylococcus capitis subsp. capitis		98.05	0.32	NIT 89				
Kocuria rosea		1.95	-.05	XYL 1				
100240	**Good identification**							**100240**
					OXI			
Staphylococcus petrasii subsp. pragensis		92.91	0.59	TRE 99	1			
Micrococcus sp.		1.82	0.25		89			
Kocuria kristinae		1.52	0.33		99			
100241	**Very good identification**							**100241**
					OXI	NOV	PYR	
Kocuria kristinae		98.94	0.74	TRE 95	99	1	93	
Staphylococcus petrasii subsp. pragensis		0.62	0.18		1	1	99	
Staphylococcus hominis subsp. novobiosepticus		0.22	0.17		1	99	1	
100243	**Very good identification**							**100243**
					OXI	NOV	PYR	
Kocuria kristinae		99.24	0.61	TRE 95	99	1	93	
Staphylococcus hominis subsp. novobiosepticus		0.48	0.12		1	99	1	
Staphylococcus epidermidis		0.28	0.15		1	1	1	
100250	**Low discrimination**							**100250**
					OXI	PYR		
Macrococcus bovicus		70.89	0.43	VPT 1	99	1		
Staphylococcus petrasii subsp. pragensis		20.84	0.18		1	99		
Macrococcus equipersicus		2.73	0.15		99	1		
100251	**Low discrimination**							**100251**
					OXI	NOV	PYR	
Kocuria kristinae		81.99	0.45	TRE 95	99	1	93	
Staphylococcus capitis subsp. capitis		16.45	0.32	MAN 4	1	1	1	
Staphylococcus cohnii subsp. cohnii		1.14	0.17		1	99	1	
100253	**Good identification**							**100253**
					OXI	PYR		
Kocuria kristinae		96.18	0.33	TRE 95	99	93		
Staphylococcus capitis subsp. ureolyticus		2.25	-.09	MAN 4	1	1		
Staphylococcus capitis subsp. capitis		1.58	-.03		1	1		
100261	**Good identification**							**100261**
Kocuria kristinae		100.0	0.33	TRE 95				
100300	**Low discrimination**							**100300**
					OXI			
Dermacoccus nishinomiyaensis		48.82	0.34	VPT 5	93			
Kocuria varians		36.62	0.39	SUC 6	13			
Micrococcus sp.		6.64	0.07		89			
100301	**Good identification**							**100301**
					OXI	PYR		
Kocuria kristinae		95.98	0.44	TRE 95	99	93		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		2.02	0.18	MLT 90	13	10		
Staphylococcus capitis subsp. capitis		1.63	0.09		1	1		
100302	**Acceptable identification**							**100302**
					NOV			
Kocuria varians		94.57	0.28	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		5.43	-.07	SUC 12	99			
100303	**Good identification**							**100303**
					OXI	PYR		
Kocuria kristinae		97.68	0.32	TRE 95	99	93		
Kocuria varians		2.32	0.06	MLT 90	13	10		
100310	**Acceptable identification**							**100310**
Micrococcus bovicus		91.78	0.28	VPT 1				
Micrococcus carouselicus		5.17	-.02					
Micrococcus equipersicus		3.05	-.01					
100311	**Low discrimination**							**100311**
					OXI	PYR		
Staphylococcus capitis subsp. capitis		84.41	0.32	NIT 89	1	1		
Kocuria kristinae		15.59	0.16	GAL 1	99	93		
100320	**Acceptable identification**							**100320**
Kocuria varians		91.64	0.30	VPT 14				
Kocuria rosea		8.36	0.00	SUC 12				
100340	**Low discrimination**							**100340**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		63.90	0.39	URE 99	1	99	1	
Staphylococcus petrasii subsp. pragensis		14.92	0.18	NIT 90	1	1	99	
Kocuria kristinae		8.04	0.23		99	1	93	
100341	**Very good identification**							**100341**
					OXI	NOV	PYR	
Kocuria kristinae		97.53	0.64	TRE 95	99	1	93	
Staphylococcus hominis subsp. novobiosepticus		1.96	0.27		1	99	1	
Staphylococcus epidermidis		0.37	0.20		1	1	1	
100342	**Low discrimination**							**100342**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. novobiosepticus		70.58	0.34	URE 99	1	99	1	
Staphylococcus epidermidis		10.59	0.25	NIT 90	1	1	1	
Micrococcus caseolyticus		9.45	0.14		99	99	99	
100343	**Acceptable identification**							**100343**
					OXI	NOV	PYR	
Kocuria kristinae		91.63	0.52	TRE 95	99	1	93	
Staphylococcus epidermidis		4.41	0.30		1	1	1	
Staphylococcus hominis subsp. novobiosepticus		3.96	0.21		1	99	1	
100350	**Low discrimination**							**100350**
					OXI	PYR		
Micrococcus bovicus		88.28	0.35	VPT 1	99	1		
Micrococcus equipersicus		3.40	0.08		99	1		
Staphylococcus haemolyticus		3.03	0.09		1	98		
100351	**Good identification**							**100351**
Kocuria kristinae		100.0	0.35	TRE 95				
100400	**Good identification**							**100400**

Code Taxon	ID score	T index	tests against	Differentiation tests		Code
				NOV	OXI	
Staphylococcus cohnii subsp. cohnii	94.09	0.86		99	1	
Staphylococcus auricularis	2.28	0.49		1	1	
Micrococcus sp.	1.19	0.28		1	89	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
100401	**Very good identification**								**100401**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		98.04	0.87		1	99			
Kocuria kristinae		1.48	0.39		99	1			
Staphylococcus auricularis		0.24	0.30		1	1			
100402	**Low discrimination**								**100402**
					NOV				
Staphylococcus cohnii subsp. cohnii		74.05	0.45	LAC 1	99				
Staphylococcus auricularis		19.77	0.30		1				
Kocuria varians		5.24	0.19		1				
100403	**Low discrimination**								**100403**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		70.81	0.46	LAC 1	1	99			
Kocuria kristinae		26.49	0.27		99	1			
Staphylococcus auricularis		1.94	0.10		1	1			
100410	**Very good identification**								**100410**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		96.35	0.88		1	12			
Macrococcus bovicus		3.18	0.51		99	99			
Macrococcus carouselicus		0.23	0.24		99	67			
100411	**Very good identification**								**100411**
					OXI	NOV	PYR		
Staphylococcus cohnii subsp. cohnii		99.42	0.89		1	99	1		
Staphylococcus capitis subsp. capitis		0.50	0.33		1	1	1		
Kocuria kristinae		0.05	0.11		99	1	93		
100412	**Good identification**								**100412**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		95.72	0.47	LAC 1	1	99	1	12	
Macrococcus bovicus		3.16	0.10		99	99	1	99	
Aerococcus viridans		1.12	0.19		1	1	95	1	
100413	**Good identification**								**100413**
					OXI	NOV	PYR		
Staphylococcus cohnii subsp. cohnii		95.45	0.48	LAC 1	1	99	1		
Aerococcus viridans		2.41	0.27		1	1	95		
Kocuria kristinae		1.17	-.02		99	1	93		
100420	**Low discrimination**								**100420**
					NOV				
Staphylococcus cohnii subsp. cohnii		75.65	0.45	XYL 1	99				
Kocuria rosea		14.88	0.19		1				
Kocuria varians		6.67	0.21		1				
100421	**Good identification**								**100421**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		97.59	0.46	XYL 1	99	1			
Kocuria kristinae		1.48	-.02		1	93			
Kocuria rosea		0.93	-.07		1	1			
100430	**Good identification**								**100430**
					NOV	YEP	OXI		
Staphylococcus cohnii subsp. cohnii		95.23	0.47	XYL 1	99	12	1		
Macrococcus bovicus		3.14	0.10		99	99	99		
Kocuria rosea		1.64	0.00		1	1	35		
100431	**Good identification**								**100431**

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		100.0	0.48	XYL 1					
100433	**Low discrimination**								**100433**
					NOV	PYR			
Aerococcus viridans		71.38	0.27	VPT 1	1	95			
Staphylococcus cohnii subsp. cohnii		28.62	0.07	ESL 90	99	1			
100440	**Low discrimination**								**100440**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		85.45	0.96		99	1			
Staphylococcus petrasii subsp. pragensis		12.81	0.59		1	99			
Staphylococcus auricularis		1.53	0.57		1	40			
100441	**Very good identification**								**100441**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		95.59	0.97		1	99			
Kocuria kristinae		4.11	0.59		99	1			
Staphylococcus auricularis		0.18	0.37		1	1			
100442	**Low discrimination**								**100442**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		68.55	0.55	LAC 1	99	1			
Staphylococcus auricularis		13.48	0.37		1	40			
Staphylococcus petrasii subsp. pragensis		10.28	0.18		1	99			
100443	**Low discrimination**								**100443**
					OXI	NOV			
Kocuria kristinae		50.70	0.47	SUC 99	99	1			
Staphylococcus cohnii subsp. cohnii		47.69	0.56		1	99			
Staphylococcus auricularis		0.96	0.18		1	1			
100450	**Very good identification**								**100450**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		97.31	0.99		1	12			
Macrococcus bovicus		2.36	0.59		99	99			
Macrococcus equipersicus		0.15	0.36		99	99			
100451	**Excellent identification**								**100451**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		99.83	0.99		1	99	1	12	
Kocuria kristinae		0.14	0.31		99	1	93	15	
Macrococcus bovicus		0.02	0.18		99	99	1	99	
100452	**Low discrimination**								**100452**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		84.75	0.58	LAC 1	1	12			
Staphylococcus saprophyticus subsp. saprophyticus		10.37	0.23		1	16			
Macrococcus bovicus		2.06	0.18		99	99			
100453	**Good identification**								**100453**
					OXI	NOV	PYR		
Staphylococcus cohnii subsp. cohnii		93.52	0.58	LAC 1	1	99	1		
Kocuria kristinae		3.26	0.18		99	1	93		
Aerococcus viridans		2.98	0.39		1	1	95		
100460	**Low discrimination**								**100460**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		85.18	0.55	XYL 1	99	1			
Staphylococcus petrasii subsp. pragensis		12.77	0.18		1	99			
Staphylococcus auricularis		1.52	0.16		1	40			
100461	**Very good identification**								**100461**
					OXI	NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		95.88	0.56	XYL 1	1	99			
Kocuria kristinae		4.12	0.18		99	1			
100463	**Low discrimination**								**100463**
					OXI	NOV	PYR		
Aerococcus viridans		39.31	0.30	VPT 1	1	1	95		
Kocuria kristinae		31.27	0.05	ESL 90	99	1	93		
Staphylococcus cohnii subsp. cohnii		29.41	0.15		1	99	1		
100470	**Very good identification**								**100470**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		97.28	0.58	XYL 1	1	12			
Macrococcus bovicus		2.36	0.18		99	99			
Staphylococcus kloosii		0.36	0.20		1	99			
100471	**Very good identification**								**100471**
					NOV	PYR			
Staphylococcus cohnii subsp. cohnii		99.65	0.58	XYL 1	99	1			
Aerococcus viridans		0.35	0.20		1	95			
100472	**Low discrimination**								**100472**
					NOV	YEP	PYR		
Aerococcus viridans		51.24	0.32	VPT 1	1	1	95		
Staphylococcus cohnii subsp. cohnii		35.03	0.16	ESL 90	99	12	1		
Staphylococcus kloosii		13.73	0.21		99	99	52		
100473	**Low discrimination**								**100473**
					NOV	PYR			
Aerococcus viridans		75.91	0.39	VPT 1	1	95			
Staphylococcus cohnii subsp. cohnii		24.10	0.17	ESL 90	99	1			
100500	**Good identification**								**100500**
					NOV				
Staphylococcus cohnii subsp. cohnii		90.41	0.67	GAL 10	99				
Kocuria varians		3.10	0.34		1				
Staphylococcus auricularis		2.19	0.30		1				
100501	**Good identification**								**100501**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		94.12	0.67	GAL 10	1	99			
Kocuria kristinae		4.27	0.30		99	1			
Staphylococcus schleiferi subsp. schleiferi		1.03	0.19		1	1			
100510	**Low discrimination**								**100510**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		87.45	0.69	GAL 10	1	12			
Macrococcus bovicus		11.12	0.44		99	99			
Macrococcus carouselicus		0.81	0.16		99	67			
100511	**Very good identification**								**100511**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		99.73	0.69	GAL 10	1	99	1	12	
Kocuria kristinae		0.15	0.01		99	1	93	15	
Macrococcus bovicus		0.12	0.03		99	99	1	99	
100512	**Low discrimination**								**100512**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		81.14	0.28	GAL 10	1	99	1	12	
Macrococcus bovicus		10.32	0.03	LAC 1	99	99	1	99	
Aerococcus viridans		8.54	0.19		1	1	95	1	
100513	**Low discrimination**								**100513**
					NOV	PYR			

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		81.52	0.28	GAL 10	99	1			
Aerococcus viridans		18.48	0.27	LAC 1	1	95			
100530	**Low discrimination**								**100530**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		88.72	0.28	GAL 10	1	12			
Macrococcus bovicus		11.28	0.03	XYL 1	99	99			
100531	**Good identification**								**100531**
Staphylococcus cohnii subsp. cohnii		100.0	0.28	GAL 10					
100533	**Good identification**								**100533**
Aerococcus viridans		100.0	0.27	VPT 1					
100540	**Low discrimination**								**100540**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		68.77	0.77	GAL 10	1	99	12		
Macrococcus caseolyticus		28.66	0.55		99	99	99		
Staphylococcus auricularis		1.23	0.37		1	1	1		
100541	**Low discrimination**								**100541**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		88.09	0.78	GAL 10	1	99	12		
Kocuria kristinae		11.35	0.49		99	1	15		
Macrococcus caseolyticus		0.34	0.14		99	99	99		
100542	**Very good identification**								**100542**
					OXI	NOV	YEP		
Macrococcus caseolyticus		97.86	0.59	NIT 99	99	99	99		
Staphylococcus cohnii subsp. cohnii		1.58	0.36		1	99	12		
Staphylococcus auricularis		0.31	0.18		1	1	1		
100543	**Low discrimination**								**100543**
					OXI	NOV	YEP		
Kocuria kristinae		65.25	0.37	SUC 99	99	1	15		
Staphylococcus cohnii subsp. cohnii		20.46	0.36		1	99	12		
Macrococcus caseolyticus		11.80	0.18		99	99	99		
100550	**Good identification**								**100550**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		90.42	0.79	GAL 10	1	12			
Macrococcus bovicus		8.47	0.51		99	99			
Macrococcus equipersicus		0.55	0.28		99	99			
100551	**Very good identification**								**100551**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		99.46	0.80	GAL 10	1	99	1	12	
Kocuria kristinae		0.42	0.21		99	1	93	15	
Macrococcus bovicus		0.09	0.10		99	99	1	99	
100552	**Low discrimination**								**100552**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		47.84	0.38	GAL 10	1	12			
Macrococcus caseolyticus		23.50	0.18	LAC 1	99	99			
Staphylococcus kloosii		13.24	0.39		1	99			
100553	**Low discrimination**								**100553**
					OXI	NOV	PYR		
Staphylococcus cohnii subsp. cohnii		71.90	0.39	GAL 10	1	99	1		
Aerococcus viridans		20.59	0.39	LAC 1	1	1	95		
Kocuria kristinae		7.51	0.08		99	1	93		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
100560	**Low discrimination**							**100560**
					NOV	OXI	YEP	
Staphylococcus cohnii subsp. cohnii		66.70	0.36	GAL 10	99	1	12	
Macrococcus caseolyticus		27.79	0.14	XYL 1	99	99	99	
Kocuria varians		5.51	0.11		1	13	80	
100561	**Low discrimination**							**100561**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		88.58	0.36	GAL 10	1	99		
Kocuria kristinae		11.42	0.08	XYL 1	99	1		
100563	**Low discrimination**							**100563**
					OXI			
Aerococcus viridans		79.04	0.30	VPT 1	1			
Kocuria kristinae		20.96	-.04	ESL 90	99			
100570	**Low discrimination**							**100570**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		86.39	0.38	GAL 10	1	12		
Macrococcus bovicus		8.10	0.10	XYL 1	99	99		
Staphylococcus kloosii		5.52	0.26		1	99		
100571	**Good identification**							**100571**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		96.92	0.39	GAL 10	99	1		
Aerococcus viridans		3.08	0.20	XYL 1	1	95		
100572	**Low discrimination**							**100572**
					NOV	YEP		
Aerococcus viridans		65.79	0.32	VPT 1	1	1		
Staphylococcus kloosii		34.22	0.27	ESL 90	99	99		
100573	**Good identification**							**100573**
Aerococcus viridans		100.0	0.39	VPT 1				
100600	**Low discrimination**							**100600**
					OXI			
Staphylococcus petrasii subsp. pragensis		88.25	0.59	MLT 99	1			
Staphylococcus auricularis		4.16	0.49		1			
Kocuria kristinae		3.04	0.39		99			
100601	**Very good identification**							**100601**
					OXI	NOV	PYR	
Kocuria kristinae		98.82	0.80	MLT 90	99	1	93	
Staphylococcus cohnii subsp. cohnii		0.67	0.46		1	99	1	
Staphylococcus petrasii subsp. pragensis		0.29	0.18		1	1	99	
100603	**Very good identification**							**100603**
					OXI			
Kocuria kristinae		99.88	0.68	MLT 90	99			
Staphylococcus auricularis		0.07	0.09		1			
Aerococcus viridans		0.03	0.17		1			
100610	**Low discrimination**							**100610**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		34.39	0.47	SUC 1	1	12		
Macrococcus bovicus		28.06	0.39		99	99		
Macrococcus carouselicus		16.50	0.30		99	67		
100611	**Low discrimination**							**100611**
					OXI	NOV	PYR	
Kocuria kristinae		71.44	0.52	MAN 4	99	1	93	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		14.73	0.48	MLT 90	1	99	1		
Staphylococcus capitis subsp. capitis		13.71	0.39		1	1	1		
100613	**Good identification**								**100613**
					OXI	PYR			
Kocuria kristinae		95.71	0.40	MAN 4	99	93			
Aerococcus viridans		1.99	0.27	MLT 90	1	95			
Staphylococcus capitis subsp. capitis		1.50	0.04		1	1			
100621	**Good identification**								**100621**
					OXI	NOV			
Kocuria kristinae		99.33	0.39	XYL 1	99	1			
Staphylococcus cohnii subsp. cohnii		0.67	0.05	MLT 90	1	99			
100623	**Good identification**								**100623**
					OXI				
Kocuria kristinae		97.22	0.27	XYL 1	99				
Aerococcus viridans		2.78	0.17	MLT 90	1				
100633	**Low discrimination**								**100633**
					OXI				
Aerococcus viridans		67.29	0.27	VPT 1	1				
Kocuria kristinae		32.71	-.02	ESL 90	99				
100640	**Excellent identification**								**100640**
					OXI				
Staphylococcus petrasii subsp. pragensis		99.15	1.00		1				
Staphylococcus petrasii subsp. jettensis		0.33	0.51		1				
Kocuria kristinae		0.31	0.59		99				
100641	**Very good identification**								**100641**
					OXI	NOV	PYR		
Kocuria kristinae		96.57	1.00		99	1	93		
Staphylococcus petrasii subsp. pragensis		3.14	0.59		1	1	99		
Staphylococcus cohnii subsp. cohnii		0.23	0.56		1	99	1		
100642	**Low discrimination**								**100642**
					OXI				
Staphylococcus petrasii subsp. pragensis		85.53	0.59	LAC 1	1				
Kocuria kristinae		6.64	0.47		99				
Staphylococcus petrasii subsp. jettensis		4.95	0.36		1				
100643	**Very good identification**								**100643**
					OXI				
Kocuria kristinae		99.80	0.88		99				
Staphylococcus petrasii subsp. pragensis		0.13	0.18		1				
Staphylococcus auricularis		0.02	0.17		1				
100650	**Low discrimination**								**100650**
					OXI	PYR	NOV	YEP	
Staphylococcus petrasii subsp. pragensis		83.73	0.59	MAN 1	1	99	1	1	
Staphylococcus cohnii subsp. cohnii		7.18	0.58		1	1	99	12	
Macrococcus bovicus		4.32	0.47		99	1	99	99	
100651	**Good identification**								**100651**
					OXI	NOV	PYR		
Kocuria kristinae		92.46	0.72	MAN 4	99	1	93		
Staphylococcus cohnii subsp. cohnii		6.71	0.58		1	99	1		
Staphylococcus petrasii subsp. pragensis		0.73	0.18		1	1	99		
100652	**Low discrimination**								**100652**
					NOV	PYR			
Staphylococcus saprophyticus subsp. saprophyticus		81.23	0.43	URE 99	99	7			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus petrasii subsp. pragensis		7.74	0.18		1	99		
Staphylococcus warneri		3.36	0.36		1	8		
100653	**Very good identification**							**100653**
					OXI	NOV	PYR	
Kocuria kristinae		98.47	0.59	MAN 4	99	1	93	
Aerococcus viridans		0.91	0.39		1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		0.33	0.02		1	99	7	
100660	**Very good identification**							**100660**
					OXI			
Staphylococcus petrasii subsp. pragensis		99.18	0.59	XYL 1	1			
Staphylococcus petrasii subsp. jettensis		0.33	0.10		1			
Kocuria kristinae		0.31	0.18		99			
100661	**Very good identification**							**100661**
					OXI	NOV	PYR	
Kocuria kristinae		96.63	0.59	XYL 1	99	1	93	
Staphylococcus petrasii subsp. pragensis		3.15	0.18		1	1	99	
Staphylococcus cohnii subsp. cohnii		0.23	0.15		1	99	1	
100663	**Good identification**							**100663**
					OXI			
Kocuria kristinae		98.75	0.47	XYL 1	99			
Aerococcus viridans		1.25	0.30		1			
100671	**Acceptable identification**							**100671**
					OXI	NOV	PYR	
Kocuria kristinae		91.08	0.31	MAN 4	99	1	93	
Staphylococcus cohnii subsp. cohnii		6.61	0.17	XYL 1	1	99	1	
Aerococcus viridans		2.31	0.20		1	1	95	
100672	**Low discrimination**							**100672**
					NOV	PYR		
Aerococcus viridans		41.82	0.32	VPT 1	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		35.37	0.02	ESL 90	99	7		
Staphylococcus arlettae		22.82	0.15		99	1		
100700	**Low discrimination**							**100700**
					OXI	NOV	PYR	
Kocuria kristinae		23.49	0.30	MLT 90	99	1	93	
Staphylococcus petrasii subsp. pragensis		20.64	0.18	MNS 99	1	1	99	
Micrococcus carouselicus		17.79	0.27		99	99	1	
100701	**Very good identification**							**100701**
					OXI	NOV		
Kocuria kristinae		97.78	0.71	MLT 90	99	1		
Rothia mucilaginosa		1.95	0.32		1	1		
Staphylococcus cohnii subsp. cohnii		0.22	0.26		1	99		
100703	**Very good identification**							**100703**
					OXI			
Kocuria kristinae		99.83	0.58	MLT 90	99			
Aerococcus viridans		0.09	0.17		1			
Rothia mucilaginosa		0.08	-.09		1			
100710	**Low discrimination**							**100710**
					OXI	YEP		
Micrococcus bovicus		49.96	0.31	VPT 1	99	99		
Micrococcus carouselicus		29.37	0.23		99	67		
Staphylococcus cohnii subsp. cohnii		15.88	0.28		1	12		
100711	**Acceptable identification**							**100711**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI	NOV	PYR		
Kocuria kristinae		92.65	0.42	MAN 4	99	1	93		
Staphylococcus cohnii subsp. cohnii		6.37	0.28	MLT 90	1	99	1		
Staphylococcus capitis subsp. capitis		0.54	-.03		1	1	1		
100713	**Acceptable identification**								**100713**
					OXI				
Kocuria kristinae		94.13	0.30	MAN 4	99				
Aerococcus viridans		5.87	0.27	MLT 90	1				
100721	**Good identification**								**100721**
					OXI				
Kocuria kristinae		98.05	0.30	XYL 1	99				
Rothia mucilaginosa		1.95	-.09	MLT 90	1				
100733	**Good identification**								**100733**
Aerococcus viridans		100.0	0.27	VPT 1					
100740	**Low discrimination**								**100740**
					OXI	NOV			
Staphylococcus petrasii subsp. pragensis		73.05	0.59	GAL 1	1	1			
Macrococcus caseolyticus		15.04	0.52		99	99			
Kocuria kristinae		7.56	0.49		99	1			
100741	**Very good identification**								**100741**
					OXI				
Kocuria kristinae		99.27	0.90		99				
Rothia mucilaginosa		0.51	0.39		1				
Staphylococcus petrasii subsp. pragensis		0.10	0.18		1				
100742	**Low discrimination**								**100742**
					OXI	NOV			
Macrococcus caseolyticus		86.45	0.55	NIT 99	99	99			
Kocuria kristinae		7.24	0.37		99	1			
Staphylococcus petrasii subsp. pragensis		2.83	0.18		1	1			
100743	**Very good identification**								**100743**
					OXI	NOV			
Kocuria kristinae		99.75	0.78		99	1			
Macrococcus caseolyticus		0.12	0.14		99	99			
Staphylococcus lugdunensis		0.07	0.14		1	1			
100750	**Low discrimination**								**100750**
					OXI	PYR	NOV	YEP	
Macrococcus bovicus		35.81	0.39	VPT 1	99	1	99	99	
Staphylococcus petrasii subsp. pragensis		16.37	0.18		1	99	1	1	
Staphylococcus cohnii subsp. cohnii		15.44	0.38		1	1	99	12	
100751	**Very good identification**								**100751**
					OXI	NOV			
Kocuria kristinae		97.35	0.62	MAN 4	99	1			
Staphylococcus cohnii subsp. cohnii		2.35	0.39		1	99			
Rothia mucilaginosa		0.12	-.02		1	1			
100753	**Good identification**								**100753**
					OXI				
Kocuria kristinae		97.30	0.49	MAN 4	99				
Aerococcus viridans		2.70	0.39		1				
100761	**Good identification**								**100761**
					OXI				
Kocuria kristinae		99.49	0.49	XYL 1	99				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Rothia mucilaginosa		0.51	-.02		1			
100763	**Good identification**							**100763**
					OXI			
Kocuria kristinae		96.33	0.37	XYL 1	99			
Aerococcus viridans		3.67	0.30		1			
100772	**Low discrimination**							**100772**
					NOV	YEP		
Staphylococcus arlettae		75.56	0.31	VPT 1	99	99		
Aerococcus viridans		24.44	0.32	GLR 92	1	1		
100773	**Low discrimination**							**100773**
					OXI	NOV	YEP	
Staphylococcus arlettae		56.28	0.34	VPT 1	1	99	99	
Aerococcus viridans		32.04	0.39	GLR 92	1	1	1	
Kocuria kristinae		11.68	0.08		99	1	15	
101000	**Low discrimination**							**101000**
					NOV			
Micrococcus sp.		41.60	0.28	VPT 3	1			
Dermacoccus nishinomiyaensis		25.15	0.33	ESL 1	80			
Macrococcus carouselicus		15.65	0.28		99			
101010	**Low discrimination**							**101010**
Macrococcus equipersicus		41.93	0.34	VPT 1				
Macrococcus bovicus		26.09	0.28	URE 86				
Macrococcus carouselicus		25.50	0.24					
101011	**Good identification**							**101011**
					NOV			
Staphylococcus capitis subsp. capitis		96.09	0.27	ESL 1	1			
Staphylococcus cohnii subsp. cohnii		3.91	0.07	NIT 89	99			
101050	**Low discrimination**							**101050**
					OXI	YEP		
Macrococcus equipersicus		63.80	0.43	VPT 1	99	99		
Macrococcus bovicus		34.26	0.36	URE 86	99	99		
Staphylococcus cohnii subsp. cohnii		1.94	0.16		1	12		
101053	**Good identification**							**101053**
Aerococcus viridans		100.0	0.33	VPT 1				
101073	**Good identification**							**101073**
Aerococcus viridans		100.0	0.33	VPT 1				
101150	**Low discrimination**							**101150**
Macrococcus equipersicus		65.07	0.35	VPT 1				
Macrococcus bovicus		34.93	0.28	URE 86				
101153	**Good identification**							**101153**
Aerococcus viridans		100.0	0.33	VPT 1				
101173	**Good identification**							**101173**
Aerococcus viridans		100.0	0.33	VPT 1				
101200	**Low discrimination**							**101200**
					NOV			
Macrococcus carouselicus		79.64	0.34	VPT 1	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Micrococcus sp.		10.31	0.07	TRE 94	1			
Kocuria rosea		4.25	0.07		1			
101201	**Good identification**							**101201**
					OXI	PYR		
Kocuria kristinae		97.79	0.42	TRE 95	99	93		
Staphylococcus capitis subsp. capitis		2.21	0.09	MLT 90	1	1		
101203	**Good identification**							**101203**
Kocuria kristinae		100.0	0.29	TRE 95				
101210	**Low discrimination**							**101210**
Staphylococcus vitulinus		48.75	0.29	VPT 1				
Macrococcus carouselicus		42.49	0.30	NIT 91				
Macrococcus bovicus		5.35	0.16					
101211	**Low discrimination**							**101211**
					OXI	PYR		
Staphylococcus capitis subsp. capitis		87.84	0.32	ESL 1	1	1		
Kocuria kristinae		12.16	0.13	NIT 89	99	93		
101230	**Good identification**							**101230**
Staphylococcus vitulinus		100.0	0.38	VPT 1				
101241	**Very good identification**							**101241**
					OXI	NOV		
Kocuria kristinae		99.92	0.61	TRE 95	99	1		
Staphylococcus hominis subsp. novobiosepticus		0.08	-.05		1	99		
101243	**Good identification**							**101243**
					OXI			
Kocuria kristinae		99.54	0.49	TRE 95	99			
Aerococcus viridans		0.46	0.23		1			
101251	**Good identification**							**101251**
Kocuria kristinae		100.0	0.33	TRE 95				
101273	**Good identification**							**101273**
Aerococcus viridans		100.0	0.33	VPT 1				
101300	**Acceptable identification**							**101300**
					OXI	NOV		
Macrococcus carouselicus		92.69	0.27	VPT 1	99	99		
Rothia mucilaginosa		7.32	0.00	TRE 94	1	1		
101301	**Low discrimination**							**101301**
					OXI			
Kocuria kristinae		87.90	0.32	TRE 95	99			
Rothia mucilaginosa		12.10	0.10	MLT 90	1			
101341	**Good identification**							**101341**
					OXI	NOV		
Kocuria kristinae		95.91	0.52	TRE 95	99	1		
Rothia mucilaginosa		3.42	0.18		1	1		
Staphylococcus hominis subsp. novobiosepticus		0.67	0.05		1	99		
101343	**Good identification**							**101343**
					OXI	NOV	PYR	
Kocuria kristinae		97.19	0.39	TRE 95	99	1	93	
Staphylococcus hominis subsp. novobiosepticus		1.46	-.01		1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		1.35	0.23		1	1	95	
101373	**Good identification**							**101373**
Aerococcus viridans		100.0	0.33	VPT 1				
101400	**Low discrimination**							**101400**
					OXI	YEP		
Macrococcus carouselicus		87.39	0.53	VPT 1	99	67		
Staphylococcus cohnii subsp. cohnii		11.86	0.45		1	12		
Macrococcus equipersicus		0.31	0.08		99	99		
101401	**Low discrimination**							**101401**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		68.80	0.46	ESL 1	1	99	1	
Kocuria kristinae		25.74	0.27		99	1	93	
Macrococcus carouselicus		4.73	0.12		99	99	1	
101403	**Low discrimination**							**101403**
					OXI	NOV	PYR	
Aerococcus viridans		48.14	0.37	VPT 1	1	1	95	
Kocuria kristinae		46.81	0.15		99	1	93	
Staphylococcus cohnii subsp. cohnii		5.06	0.05		1	99	1	
101410	**Low discrimination**							**101410**
					OXI	YEP		
Macrococcus carouselicus		67.74	0.49	VPT 1	99	67		
Staphylococcus cohnii subsp. cohnii		18.30	0.47		1	12		
Macrococcus equipersicus		7.11	0.34		99	99		
101411	**Acceptable identification**							**101411**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		93.29	0.48	ESL 1	1	99	1	
Macrococcus carouselicus		3.22	0.08		99	99	1	
Aerococcus viridans		2.35	0.27		1	1	95	
101412	**Low discrimination**							**101412**
					OXI	NOV	PYR	
Aerococcus viridans		68.91	0.39	VPT 1	1	1	95	
Macrococcus carouselicus		24.48	0.08		99	99	1	
Staphylococcus cohnii subsp. cohnii		6.61	0.06		1	99	1	
101413	**Good identification**							**101413**
					NOV	PYR		
Aerococcus viridans		95.74	0.47	VPT 1	1	95		
Staphylococcus cohnii subsp. cohnii		4.27	0.07		99	1		
101422	**Good identification**							**101422**
Aerococcus viridans		100.0	0.29	VPT 1				
101423	**Good identification**							**101423**
Aerococcus viridans		100.0	0.37	VPT 1				
101431	**Low discrimination**							**101431**
					NOV	PYR		
Aerococcus viridans		71.38	0.27	VPT 1	1	95		
Staphylococcus cohnii subsp. cohnii		28.62	0.07	LAC 90	99	1		
101432	**Good identification**							**101432**
Aerococcus viridans		100.0	0.39	VPT 1				
101433	**Good identification**							**101433**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.47	VPT 1				
101440	**Low discrimination**							**101440**
					OXI	PYR	NOV	
Staphylococcus cohnii subsp. cohnii		80.98	0.55	ESL 1	1	1	99	
Staphylococcus petrasii subsp. pragensis		12.14	0.18		1	99	1	
Macrococcus carouselicus		1.90	0.12		99	1	99	
101441	**Low discrimination**							**101441**
					OXI	NOV	PYR	
Kocuria kristinae		51.20	0.47	SUC 99	99	1	93	
Staphylococcus cohnii subsp. cohnii		48.15	0.56		1	99	1	
Aerococcus viridans		0.65	0.30		1	1	95	
101442	**Low discrimination**							**101442**
					NOV	PYR		
Aerococcus viridans		84.81	0.42	VPT 1	1	95		
Staphylococcus cohnii subsp. cohnii		15.19	0.14		99	1		
101443	**Low discrimination**							**101443**
					OXI	NOV	PYR	
Kocuria kristinae		66.89	0.34	SUC 99	99	1	93	
Aerococcus viridans		30.57	0.49		1	1	95	
Staphylococcus cohnii subsp. cohnii		2.54	0.15		1	99	1	
101450	**Low discrimination**							**101450**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		60.97	0.58	ESL 1	1	12		
Macrococcus equipersicus		20.23	0.43		99	99		
Macrococcus bovicus		16.29	0.39		99	99		
101451	**Good identification**							**101451**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		93.27	0.58	ESL 1	1	99	1	
Kocuria kristinae		3.25	0.18		99	1	93	
Aerococcus viridans		2.97	0.39		1	1	95	
101452	**Low discrimination**							**101452**
					NOV	YEP	PYR	
Aerococcus viridans		80.61	0.51	VPT 1	1	1	95	
Staphylococcus kloosii		9.60	0.33		99	99	52	
Staphylococcus cohnii subsp. cohnii		6.12	0.16		99	12	1	
101453	**Good identification**							**101453**
					OXI	NOV	PYR	
Aerococcus viridans		93.84	0.59	VPT 1	1	1	95	
Staphylococcus cohnii subsp. cohnii		3.31	0.17		1	99	1	
Kocuria kristinae		2.85	0.06		99	1	93	
101461	**Low discrimination**							**101461**
					OXI	NOV	PYR	
Aerococcus viridans		39.31	0.30	VPT 1	1	1	95	
Kocuria kristinae		31.27	0.05	LAC 90	99	1	93	
Staphylococcus cohnii subsp. cohnii		29.42	0.15		1	99	1	
101462	**Good identification**							**101462**
Aerococcus viridans		100.0	0.42	VPT 1				
101463	**Good identification**							**101463**
Aerococcus viridans		100.0	0.49	VPT 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
101470	**Low discrimination**							**101470**
					NOV	YEP	PYR	
Aerococcus viridans		42.72	0.32	VPT 1	1	1	95	
Staphylococcus cohnii subsp. cohnii		29.21	0.16	LAC 90	99	12	1	
Staphylococcus kloosii		10.57	0.20		99	99	52	
101471	**Low discrimination**							**101471**
					NOV	PYR		
Aerococcus viridans		75.91	0.39	VPT 1	1	95		
Staphylococcus cohnii subsp. cohnii		24.10	0.17	LAC 90	99	1		
101472	**Good identification**							**101472**
					NOV	YEP		
Aerococcus viridans		97.11	0.51	VPT 1	1	1		
Staphylococcus kloosii		2.89	0.21		99	99		
101473	**Very good identification**							**101473**
Aerococcus viridans		100.0	0.59	VPT 1				
101500	**Good identification**							**101500**
					OXI	NOV		
Macrococcus carouselicus		95.33	0.45	VPT 1	99	99		
Staphylococcus cohnii subsp. cohnii		3.35	0.25		1	99		
Rothia mucilaginosa		0.98	0.00		1	1		
101502	**Low discrimination**							**101502**
					OXI	NOV		
Aerococcus viridans		58.33	0.29	VPT 1	1	1		
Macrococcus carouselicus		41.67	0.04		99	99		
101503	**Low discrimination**							**101503**
					OXI			
Aerococcus viridans		75.52	0.37	VPT 1	1			
Kocuria kristinae		24.48	0.05		99			
101510	**Low discrimination**							**101510**
Macrococcus carouselicus		78.10	0.41	VPT 1				
Macrococcus equipersicus		8.20	0.26					
Macrococcus bovicus		7.65	0.24					
101511	**Low discrimination**							**101511**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		73.54	0.28	ESL 1	1	99	1	
Aerococcus viridans		16.67	0.27	GAL 10	1	1	95	
Macrococcus carouselicus		9.79	0.00		99	99	1	
101512	**Low discrimination**							**101512**
					OXI	NOV		
Aerococcus viridans		86.79	0.39	VPT 1	1	1		
Macrococcus carouselicus		13.21	0.00		99	99		
101513	**Good identification**							**101513**
Aerococcus viridans		100.0	0.47	VPT 1				
101522	**Good identification**							**101522**
Aerococcus viridans		100.0	0.29	VPT 1				
101523	**Good identification**							**101523**
Aerococcus viridans		100.0	0.37	VPT 1				

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
101531	**Good identification**							**101531**
Aerococcus viridans		100.0	0.27	VPT 1				
101532	**Good identification**							**101532**
Aerococcus viridans		100.0	0.39	VPT 1				
101533	**Good identification**							**101533**
Aerococcus viridans		100.0	0.47	VPT 1				
101540	**Low discrimination**							**101540**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		52.67	0.36	ESL 1	1	99	12	
Macrococcus caseolyticus		21.95	0.14	GAL 10	99	99	99	
Rothia mucilaginosa		11.28	0.08		1	1	1	
101541	**Low discrimination**							**101541**
					OXI	NOV		
Kocuria kristinae		65.04	0.37	SUC 99	99	1		
Staphylococcus cohnii subsp. cohnii		20.39	0.36		1	99		
Rothia mucilaginosa		12.10	0.18		1	1		
101543	**Low discrimination**							**101543**
					OXI			
Aerococcus viridans		57.83	0.49	VPT 1	1			
Kocuria kristinae		42.17	0.24		99			
101550	**Low discrimination**							**101550**
					OXI	YEP		
Macrococcus equipersicus		34.18	0.35	VPT 1	99	99		
Macrococcus bovicus		27.53	0.32	URE 86	99	99		
Staphylococcus cohnii subsp. cohnii		26.71	0.38		1	12		
101551	**Low discrimination**							**101551**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		71.90	0.39	ESL 1	1	99	1	
Aerococcus viridans		20.59	0.39	GAL 10	1	1	95	
Kocuria kristinae		7.51	0.08		99	1	93	
101552	**Low discrimination**							**101552**
					NOV	YEP		
Aerococcus viridans		81.22	0.51	VPT 1	1	1		
Staphylococcus kloosii		18.78	0.39		99	99		
101553	**Very good identification**							**101553**
					OXI			
Aerococcus viridans		99.00	0.59	VPT 1	1			
Kocuria kristinae		1.00	-.04		99			
101561	**Low discrimination**							**101561**
					OXI			
Aerococcus viridans		79.04	0.30	VPT 1	1			
Kocuria kristinae		20.96	-.04	LAC 90	99			
101562	**Good identification**							**101562**
Aerococcus viridans		100.0	0.42	VPT 1				
101563	**Good identification**							**101563**
Aerococcus viridans		100.0	0.49	VPT 1				
101570	**Low discrimination**							**101570**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Aerococcus viridans		67.56	0.32	VPT 1	1	1		
Staphylococcus kloosii		32.44	0.26	LAC 90	99	99		
101571	**Good identification**							**101571**
Aerococcus viridans		100.0	0.39	VPT 1				
101572	**Acceptable identification**							**101572**
					NOV	YEP		
Aerococcus viridans		94.54	0.51	VPT 1	1	1		
Staphylococcus kloosii		5.46	0.27		99	99		
101573	**Very good identification**							**101573**
Aerococcus viridans		100.0	0.59	VPT 1				
101600	**Good identification**							**101600**
					OXI	NOV	PYR	
Macrococcus carouselicus		94.06	0.59	VPT 1	99	99	1	
Staphylococcus petrasii subsp. pragensis		2.99	0.18		1	1	99	
Kocuria kristinae		2.55	0.27		99	1	93	
101601	**Very good identification**							**101601**
					OXI	NOV	PYR	
Kocuria kristinae		99.33	0.68	MLT 90	99	1	93	
Macrococcus carouselicus		0.37	0.18		99	99	1	
Rothia mucilaginosa		0.24	0.10		1	1	75	
101603	**Very good identification**							**101603**
					OXI			
Kocuria kristinae		98.97	0.56	MLT 90	99			
Aerococcus viridans		1.03	0.37		1			
101610	**Good identification**							**101610**
Macrococcus carouselicus		94.07	0.55	VPT 1				
Staphylococcus vitulinus		3.88	0.24					
Macrococcus bovicus		1.14	0.19					
101611	**Acceptable identification**							**101611**
					OXI	NOV	PYR	
Kocuria kristinae		90.75	0.40	MAN 4	99	1	93	
Macrococcus carouselicus		5.25	0.14	MLT 90	99	99	1	
Aerococcus viridans		1.89	0.27		1	1	95	
101612	**Low discrimination**							**101612**
					OXI	NOV		
Aerococcus viridans		58.10	0.39	VPT 1	1	1		
Macrococcus carouselicus		41.90	0.14		99	99		
101613	**Low discrimination**							**101613**
					OXI			
Kocuria kristinae		57.21	0.27	MAN 4	99			
Aerococcus viridans		42.79	0.47	MLT 90	1			
101621	**Good identification**							**101621**
					OXI			
Kocuria kristinae		97.22	0.27	XYL 1	99			
Aerococcus viridans		2.78	0.17	MLT 90	1			
101622	**Good identification**							**101622**
Aerococcus viridans		100.0	0.29	VPT 1				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
101623	**Low discrimination**							**101623**
					OXI			
Aerococcus viridans		50.70	0.37	VPT 1	1			
Kocuria kristinae		49.30	0.15		99			
101630	**Acceptable identification**							**101630**
					OXI	NOV		
Staphylococcus vitulinus		90.53	0.33	VPT 1	99	99		
Macrococcus carouselicus		8.21	0.14	NIT 91	99	99		
Aerococcus viridans		1.27	0.19		1	1		
101631	**Low discrimination**							**101631**
					OXI	NOV	PYR	
Aerococcus viridans		55.68	0.27	VPT 1	1	1	95	
Kocuria kristinae		27.07	-.02	LAC 90	99	1	93	
Staphylococcus vitulinus		17.25	-.08		99	99	1	
101632	**Acceptable identification**							**101632**
					OXI	NOV		
Aerococcus viridans		92.56	0.39	VPT 1	1	1		
Staphylococcus vitulinus		7.44	-.08		99	99		
101633	**Good identification**							**101633**
Aerococcus viridans		100.0	0.47	VPT 1				
101640	**Good identification**							**101640**
					OXI			
Staphylococcus petrasii subsp. pragensis		91.92	0.59	ESL 1	1			
Kocuria kristinae		7.13	0.47		99			
Staphylococcus petrasii subsp. jettensis		0.31	0.10		1			
101641	**Very good identification**							**101641**
					OXI			
Kocuria kristinae		99.78	0.88		99			
Staphylococcus petrasii subsp. pragensis		0.13	0.18		1			
Rothia mucilaginosa		0.06	0.18		1			
101642	**Low discrimination**							**101642**
					OXI			
Kocuria kristinae		57.25	0.34	MNS 99	99			
Staphylococcus petrasii subsp. pragensis		29.81	0.18		1			
Aerococcus viridans		11.22	0.42		1			
101643	**Very good identification**							**101643**
					OXI			
Kocuria kristinae		99.54	0.75		99			
Aerococcus viridans		0.46	0.49		1			
101651	**Very good identification**							**101651**
					OXI	NOV	PYR	
Kocuria kristinae		98.50	0.59	MAN 4	99	1	93	
Aerococcus viridans		0.91	0.39		1	1	95	
Staphylococcus sciuri		0.30	0.13		99	99	1	
101652	**Low discrimination**							**101652**
					OXI	NOV	PYR	
Aerococcus viridans		85.85	0.51	VPT 1	1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		8.07	0.02		1	99	7	
Kocuria kristinae		6.09	0.06		99	1	93	
101653	**Low discrimination**							**101653**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		75.05	0.47	MAN 4	99			
Aerococcus viridans		24.95	0.59		1			
101661	**Good identification**							**101661**
					OXI			
Kocuria kristinae		98.75	0.47	XYL 1	99			
Aerococcus viridans		1.25	0.30		1			
101662	**Good identification**							**101662**
Aerococcus viridans		100.0	0.42	VPT 1				
101663	**Low discrimination**							**101663**
					OXI			
Kocuria kristinae		68.63	0.34	XYL 1	99			
Aerococcus viridans		31.37	0.49		1			
101670	**Low discrimination**							**101670**
					OXI	NOV		
Aerococcus viridans		84.69	0.32	VPT 1	1	1		
Staphylococcus vitulinus		15.31	-.08	LAC 90	99	99		
101672	**Acceptable identification**							**101672**
					NOV	YEP		
Aerococcus viridans		94.28	0.51	VPT 1	1	1		
Staphylococcus arlettae		5.72	0.15		99	99		
101673	**Good identification**							**101673**
					OXI	NOV	YEP	
Aerococcus viridans		93.91	0.59	VPT 1	1	1	1	
Staphylococcus arlettae		3.23	0.18		1	99	99	
Kocuria kristinae		2.85	0.06		99	1	15	
101700	**Low discrimination**							**101700**
					OXI	NOV	PYR	
Macrococcus carouselicus		65.80	0.51	VPT 1	99	99	1	
Rothia mucilaginosa		32.81	0.42		1	1	75	
Kocuria kristinae		1.39	0.17		99	1	93	
101701	**Low discrimination**							**101701**
					OXI	NOV	PYR	
Kocuria kristinae		58.04	0.58	MLT 90	99	1	93	
Rothia mucilaginosa		41.63	0.51		1	1	75	
Macrococcus carouselicus		0.28	0.10		99	99	1	
101703	**Acceptable identification**							**101703**
					OXI			
Kocuria kristinae		94.33	0.46	MLT 90	99			
Aerococcus viridans		2.94	0.37		1			
Rothia mucilaginosa		2.73	0.10		1			
101710	**Good identification**							**101710**
					OXI	NOV		
Macrococcus carouselicus		97.21	0.47	VPT 1	99	99		
Macrococcus bovicus		1.17	0.12		99	99		
Rothia mucilaginosa		0.77	0.00		1	1		
101711	**Low discrimination**							**101711**
					OXI	NOV	PYR	
Kocuria kristinae		68.12	0.30	MAN 4	99	1	93	
Rothia mucilaginosa		11.85	0.10	MLT 90	1	1	75	
Staphylococcus lentus		7.82	0.18		99	99	6	
101712	**Low discrimination**							**101712**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Aerococcus viridans		76.39	0.39	VPT 1	1	1		
Macrococcus carouselicus		23.61	0.06		99	99		
101713	**Low discrimination**							**101713**
					OXI	NOV	PYR	
Aerococcus viridans		46.05	0.47	VPT 1	1	1	95	
Staphylococcus lentus		33.43	0.30		99	99	6	
Kocuria kristinae		20.52	0.17		99	1	93	
101722	**Good identification**							**101722**
Aerococcus viridans		100.0	0.29	VPT 1				
101723	**Low discrimination**							**101723**
					OXI			
Aerococcus viridans		75.52	0.37	VPT 1	1			
Kocuria kristinae		24.48	0.05		99			
101732	**Good identification**							**101732**
Aerococcus viridans		100.0	0.39	VPT 1				
101733	**Low discrimination**							**101733**
					OXI	NOV		
Aerococcus viridans		57.94	0.47	VPT 1	1	1		
Staphylococcus lentus		42.06	0.30		99	99		
101740	**Low discrimination**							**101740**
					OXI			
Rothia mucilaginosa		79.93	0.49	NIT 99	1			
Kocuria kristinae		13.03	0.37		99			
Staphylococcus petrasii subsp. pragensis		5.09	0.18		1			
101741	**Low discrimination**							**101741**
					OXI			
Kocuria kristinae		84.29	0.78		99			
Rothia mucilaginosa		15.68	0.59		1			
Aerococcus viridans		0.03	0.30		1			
101743	**Very good identification**							**101743**
					OXI			
Kocuria kristinae		97.91	0.65		99			
Aerococcus viridans		1.36	0.49		1			
Rothia mucilaginosa		0.74	0.18		1			
101751	**Low discrimination**							**101751**
					OXI			
Kocuria kristinae		89.57	0.49	MAN 4	99			
Rothia mucilaginosa		4.04	0.18		1			
Aerococcus viridans		2.48	0.39		1			
101752	**Good identification**							**101752**
					OXI	NOV	YEP	
Aerococcus viridans		95.56	0.51	VPT 1	1	1	1	
Kocuria kristinae		2.26	-.04		99	1	15	
Staphylococcus kloosii		2.19	0.18		1	99	99	
101753	**Low discrimination**							**101753**
					OXI	NOV	PYR	
Kocuria kristinae		42.08	0.37	MAN 4	99	1	93	
Aerococcus viridans		41.97	0.59		1	1	95	
Staphylococcus lentus		15.46	0.36		99	99	6	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
101761	**Low discrimination**							**101761**
					OXI			
Kocuria kristinae		81.70	0.37	XYL 1	99			
Rothia mucilaginosa		15.19	0.18		1			
Aerococcus viridans		3.11	0.30		1			
101762	**Good identification**							**101762**
Aerococcus viridans		100.0	0.42	VPT 1				
101763	**Low discrimination**							**101763**
					OXI			
Aerococcus viridans		57.83	0.49	VPT 1	1			
Kocuria kristinae		42.17	0.24		99			
101770	**Good identification**							**101770**
Aerococcus viridans		100.0	0.32	VPT 1				
101771	**Low discrimination**							**101771**
					OXI	NOV		
Aerococcus viridans		35.91	0.39	VPT 1	1	1		
Staphylococcus lentus		33.58	0.25	LAC 90	99	99		
Staphylococcus sciuri		17.41	0.17		99	99		
101772	**Low discrimination**							**101772**
					NOV	YEP		
Aerococcus viridans		74.43	0.51	VPT 1	1	1		
Staphylococcus arlettae		25.57	0.31		99	99		
101773	**Low discrimination**							**101773**
					OXI	NOV	YEP	
Aerococcus viridans		63.19	0.59	VPT 1	1	1	1	
Staphylococcus lentus		23.28	0.36		99	99	1	
Staphylococcus arlettae		12.33	0.34		1	99	99	
102000	**Low discrimination**							**102000**
					OXI			
Kocuria rosea		85.90	0.88		35			
Dermacoccus nishinomiyaensis		8.84	0.69		93			
Kocuria varians		3.42	0.68		13			
102001	**Low discrimination**							**102001**
Kocuria rosea		48.04	0.61	MNS 5				
Staphylococcus capitis subsp. capitis		46.03	0.63					
Kocuria varians		3.16	0.46					
102002	**Low discrimination**							**102002**
					OXI			
Kocuria varians		48.74	0.56	VPT 14	13			
Kocuria rosea		43.84	0.47		35			
Dermacoccus nishinomiyaensis		4.51	0.28		93			
102003	**Low discrimination**							**102003**
Staphylococcus capitis subsp. capitis		39.67	0.29	MAN 93				
Kocuria varians		37.65	0.35	LAC 2				
Kocuria rosea		20.49	0.20					
102010	**Good identification**							**102010**
					OXI			
Kocuria rosea		92.15	0.68	MAN 10	35			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus capitis subsp. capitis		5.61	0.45		1		
Dermacoccus nishinomiyaensis		0.86	0.28		93		
102011	**Very good identification**						**102011**
					NOV		
Staphylococcus capitis subsp. capitis		99.09	0.87		1		
Kocuria rosea		0.87	0.42		1		
Staphylococcus cohnii subsp. cohnii		0.04	0.25		99		
102012	**Low discrimination**						**102012**
Kocuria rosea		81.69	0.27	MAN 10			
Staphylococcus capitis subsp. capitis		10.05	0.11	LAC 1			
Kocuria varians		8.26	0.15				
102013	**Good identification**						**102013**
Staphylococcus capitis subsp. capitis		99.35	0.52	LAC 2			
Kocuria rosea		0.43	0.01				
Staphylococcus carnosus subsp. carnosus		0.22	-.02				
102020	**Very good identification**						**102020**
					OXI		
Kocuria rosea		96.57	0.80		35		
Kocuria varians		3.15	0.58		13		
Dermacoccus nishinomiyaensis		0.23	0.28		93		
102021	**Good identification**						**102021**
Kocuria rosea		92.89	0.54	MNS 5			
Kocuria varians		5.01	0.37				
Staphylococcus capitis subsp. capitis		2.10	0.22				
102022	**Low discrimination**						**102022**
Kocuria rosea		52.32	0.39	LAC 1			
Kocuria varians		47.68	0.47				
102030	**Very good identification**						**102030**
					NOV	OXI	
Kocuria rosea		99.20	0.60	MAN 10	1	35	
Staphylococcus vitulinus		0.36	0.09		99	99	
Kocuria varians		0.29	0.17		1	13	
102031	**Low discrimination**						**102031**
Staphylococcus capitis subsp. capitis		72.97	0.45	XYL 1			
Kocuria rosea		27.03	0.34				
102040	**Low discrimination**						**102040**
Staphylococcus auricularis		32.36	0.55	VPT 12			
Kocuria rosea		30.47	0.47	TRE 92			
Kocuria varians		26.37	0.54				
102041	**Low discrimination**						**102041**
					NOV		
Staphylococcus auricularis		31.14	0.35	VPT 12	1		
Kocuria varians		19.86	0.32	TRE 92	1		
Staphylococcus cohnii subsp. cohnii		16.49	0.33	MNS 10	99		
102042	**Low discrimination**						**102042**
					NOV	OXI	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		55.22	0.43	VPT 14	1	13		
Staphylococcus auricularis		26.70	0.35		1	1		
Macrococcus caseolyticus		12.04	0.18		99	99		
102050	**Low discrimination**							**102050**
					NOV	YEP	OXI	
Macrococcus bovicus		44.60	0.36	VPT 1	99	99	99	
Kocuria rosea		28.52	0.27	NIT 10	1	1	35	
Staphylococcus cohnii subsp. cohnii		18.84	0.34		99	12	1	
102051	**Low discrimination**							**102051**
					NOV			
Staphylococcus capitis subsp. capitis		88.70	0.45	MLT 1	1			
Staphylococcus cohnii subsp. cohnii		10.53	0.35		99			
Kocuria rosea		0.77	0.01		1			
102060	**Low discrimination**							**102060**
Kocuria rosea		57.66	0.39	MLT 1				
Kocuria varians		40.90	0.45					
Staphylococcus auricularis		1.44	0.14					
102062	**Good identification**							**102062**
Kocuria varians		95.19	0.34	VPT 14				
Kocuria rosea		4.81	-.02					
102100	**Low discrimination**							**102100**
					OXI			
Kocuria varians		66.03	0.71	VPT 14	13			
Dermacoccus nishinomiyaensis		21.67	0.54		93			
Kocuria rosea		11.17	0.47		35			
102101	**Low discrimination**							**102101**
Kocuria varians		69.47	0.50	VPT 14				
Staphylococcus schleiferi subsp. schleiferi		12.84	0.29	MNS 8				
Kocuria rosea		7.11	0.20					
102102	**Good identification**							**102102**
					OXI	YEP	NOV	
Kocuria varians		94.96	0.60	VPT 14	13	80	1	
Macrococcus caseolyticus		3.03	0.18		99	99	99	
Dermacoccus nishinomiyaensis		1.12	0.13		93	1	80	
102103	**Good identification**							**102103**
Kocuria varians		100.0	0.38	VPT 14				
102110	**Low discrimination**							**102110**
					NOV	YEP	OXI	
Kocuria rosea		47.96	0.27	GAL 1	1	1	35	
Kocuria varians		25.78	0.30	MAN 10	1	80	13	
Macrococcus bovicus		13.78	0.21		99	99	99	
102111	**Good identification**							**102111**
Staphylococcus capitis subsp. capitis		97.98	0.45	GAL 1				
Kocuria rosea		0.86	0.01					
Kocuria varians		0.76	0.09					
102120	**Low discrimination**							**102120**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		82.26	0.62	VPT 14	13			
Kocuria rosea		16.97	0.39		35			
Dermacoccus nishinomiyaensis		0.78	0.13		93			
102121	**Low discrimination**							**102121**
Kocuria varians		88.90	0.40	VPT 14				
Kocuria rosea		11.10	0.13	MNS 8				
102122	**Good identification**							**102122**
Kocuria varians		99.27	0.51	VPT 14				
Kocuria rosea		0.73	-.02					
102123	**Good identification**							**102123**
Kocuria varians		100.0	0.29	VPT 14				
102140	**Low discrimination**							**102140**
					NOV	OXI		
Macrococcus caseolyticus		69.89	0.55	TRE 99	99	99		
Kocuria varians		25.82	0.58		1	13		
Staphylococcus auricularis		2.35	0.35		1	1		
102141	**Low discrimination**							**102141**
					NOV	OXI		
Kocuria varians		57.02	0.36	VPT 14	1	13		
Macrococcus caseolyticus		17.93	0.14	MNS 8	99	99		
Staphylococcus auricularis		6.62	0.16		1	1		
102142	**Good identification**							**102142**
					NOV	OXI	YEP	
Macrococcus caseolyticus		92.53	0.59	TRE 99	99	99	99	
Kocuria varians		6.43	0.47		1	13	80	
Staphylococcus hominis subsp. novobiosepticus		0.42	0.12		99	1	1	
102150	**Low discrimination**							**102150**
					NOV	OXI		
Macrococcus bovicus		54.37	0.28	VPT 1	99	99		
Macrococcus caseolyticus		25.91	0.14	NIT 10	99	99		
Kocuria varians		9.57	0.17		1	13		
102160	**Acceptable identification**							**102160**
					NOV	YEP	OXI	
Kocuria varians		91.97	0.49	VPT 14	1	80	13	
Macrococcus caseolyticus		7.16	0.14		99	99	99	
Kocuria rosea		0.87	-.02		1	1	35	
102161	**Good identification**							**102161**
Kocuria varians		100.0	0.27	VPT 14				
102162	**Low discrimination**							**102162**
					NOV	OXI		
Kocuria varians		70.73	0.37	VPT 14	1	13		
Macrococcus caseolyticus		29.27	0.18		99	99		
102200	**Low discrimination**							**102200**
					OXI			
Kocuria rosea		85.25	0.68	SUC 10	35			
Dermacoccus nishinomiyaensis		5.04	0.44		93			
Kocuria varians		4.17	0.50		13			
102201	**Good identification**							**102201**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR	OXI	
Staphylococcus capitis subsp. capitis		91.67	0.69	MAN 93	1	1	
Kocuria rosea		5.72	0.42		1	35	
Kocuria kristinae		1.52	0.31		93	99	
102202	**Low discrimination**						**102202**
Kocuria varians		45.59	0.39	VPT 14			
Kocuria rosea		33.41	0.27	SUC 12			
Staphylococcus auricularis		14.04	0.27				
102203	**Low discrimination**						**102203**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		73.87	0.34	MAN 93	1	1	
Kocuria kristinae		15.02	0.19	LAC 2	99	93	
Kocuria varians		5.15	0.17		13	10	
102210	**Low discrimination**						**102210**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		39.62	0.51	MNS 99	1	1	
Kocuria rosea		38.92	0.48		1	35	
Staphylococcus vitulinus		20.19	0.41		99	99	
102211	**Very good identification**						**102211**
					PYR	OXI	
Staphylococcus capitis subsp. capitis		99.94	0.92		1	1	
Kocuria rosea		0.05	0.22		1	35	
Kocuria kristinae		0.01	0.03		93	99	
102213	**Very good identification**						**102213**
Staphylococcus capitis subsp. capitis		99.86	0.57	LAC 2			
Staphylococcus capitis subsp. ureolyticus		0.14	-.11				
102220	**Very good identification**						**102220**
					NOV	OXI	
Kocuria rosea		95.60	0.60	SUC 10	1	35	
Kocuria varians		3.83	0.41		1	13	
Staphylococcus vitulinus		0.35	0.09		99	99	
102221	**Low discrimination**						**102221**
Kocuria rosea		69.27	0.34	SUC 10			
Staphylococcus capitis subsp. capitis		26.15	0.28	MNS 5			
Kocuria varians		4.59	0.19				
102222	**Low discrimination**						**102222**
Kocuria varians		52.80	0.29	VPT 14			
Kocuria rosea		47.20	0.19	SUC 12			
102230	**Low discrimination**						**102230**
					NOV	OXI	
Staphylococcus vitulinus		76.17	0.50	VPT 1	99	99	
Kocuria rosea		23.27	0.41		1	35	
Staphylococcus capitis subsp. capitis		0.56	0.10		1	1	
102231	**Good identification**						**102231**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		96.52	0.51	XYL 1	1	1	
Kocuria rosea		2.14	0.14		1	35	
Staphylococcus vitulinus		1.34	0.09		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
102240	**Low discrimination**							**102240**
					NOV	PYR		
Staphylococcus hominis subsp. novobiosepticus		27.64	0.49	URE 99	99	1		
Staphylococcus petrasii subsp. pragensis		23.67	0.39		1	99		
Staphylococcus auricularis		18.35	0.54		1	40		
102241	**Low discrimination**							**102241**
					OXI	NOV	PYR	
Kocuria kristinae		64.19	0.51	NIT 7	99	1	93	
Staphylococcus hominis subsp. novobiosepticus		17.11	0.37	TRE 95	1	99	1	
Staphylococcus epidermidis		6.16	0.35		1	1	1	
102242	**Low discrimination**							**102242**
					PYR	NOV		
Staphylococcus hominis subsp. novobiosepticus		38.21	0.44	URE 99	1	99		
Staphylococcus haemolyticus		23.22	0.49		98	2		
Staphylococcus hominis subsp. hominis		17.60	0.48		1	1		
102243	**Low discrimination**							**102243**
					OXI	NOV	PYR	
Staphylococcus epidermidis		40.98	0.46	URE 92	1	1	1	
Kocuria kristinae		33.69	0.38		99	1	93	
Staphylococcus hominis subsp. novobiosepticus		19.34	0.31		1	99	1	
102250	**Low discrimination**							**102250**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		75.66	0.49	ARG 92	1	98	2	
Staphylococcus hominis subsp. hominis		6.99	0.30	TRE 93	1	1	1	
Macrococcus bovicus		6.52	0.23		99	1	99	
102251	**Acceptable identification**							**102251**
					OXI	PYR		
Staphylococcus capitis subsp. capitis		93.35	0.51	MLT 1	1	1		
Kocuria kristinae		4.33	0.22		99	93		
Staphylococcus haemolyticus		1.17	0.18		1	98		
102252	**Low discrimination**							**102252**
					PYR	NOV		
Staphylococcus haemolyticus		81.90	0.44	ARG 92	98	2		
Staphylococcus hominis subsp. hominis		15.79	0.31	TRE 93	1	1		
Staphylococcus hominis subsp. novobiosepticus		2.32	0.03		1	99		
102300	**Low discrimination**							**102300**
					OXI			
Kocuria varians		64.01	0.54	VPT 14	13			
Dermacoccus nishinomiyaensis		9.83	0.29	SUC 12	93			
Kocuria rosea		8.82	0.27		35			
102301	**Low discrimination**							**102301**
Rothia mucilaginosa		51.30	0.32	ESL 90				
Staphylococcus capitis subsp. capitis		18.62	0.28	TRE 99				
Kocuria varians		13.94	0.32					
102302	**Low discrimination**							**102302**
					NOV	OXI	YEP	
Kocuria varians		72.27	0.42	VPT 14	1	13	80	
Macrococcus caseolyticus		11.28	0.14	SUC 12	99	99	99	
Staphylococcus hominis subsp. novobiosepticus		7.66	0.12		99	1	1	
102311	**Good identification**							**102311**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus capitis subsp. capitis		99.79	0.51	GAL 1					
Rothia mucilaginosa		0.21	-.09						
102320	**Low discrimination**								**102320**
Kocuria varians		85.61	0.44	VPT 14					
Kocuria rosea		14.39	0.19	SUC 12					
102322	**Good identification**								**102322**
Kocuria varians		100.0	0.33	VPT 14					
102340	**Low discrimination**								**102340**
					OXI	NOV	YEP	PYR	
Staphylococcus hominis subsp. novobiosepticus		49.28	0.59	URE 99	1	99	1	1	
Micrococcus caseolyticus		26.06	0.52		99	99	99	80	
Staphylococcus haemolyticus		10.78	0.55		1	2	20	98	
102341	**Low discrimination**								**102341**
					NOV				
Staphylococcus hominis subsp. novobiosepticus		36.76	0.47	URE 99	99				
Staphylococcus epidermidis		25.02	0.51		1				
Rothia mucilaginosa		19.98	0.39		1				
102342	**Low discrimination**								**102342**
					OXI	NOV	YEP		
Micrococcus caseolyticus		42.92	0.55	TRE 99	99	99	99		
Staphylococcus hominis subsp. novobiosepticus		29.13	0.53		1	99	1		
Staphylococcus epidermidis		15.70	0.56		1	1	1		
102343	**Low discrimination**								**102343**
					OXI	NOV	PYR		
Staphylococcus epidermidis		74.82	0.61	URE 92	1	1	1		
Staphylococcus hominis subsp. novobiosepticus		18.69	0.41		1	99	1		
Kocuria kristinae		3.62	0.28		99	1	93		
102350	**Low discrimination**								**102350**
					PYR	NOV			
Staphylococcus haemolyticus		79.70	0.50	ARG 92	98	2			
Staphylococcus hominis subsp. hominis		7.07	0.30	TRE 93	1	1			
Staphylococcus hominis subsp. novobiosepticus		6.27	0.18		1	99			
102352	**Low discrimination**								**102352**
					OXI	PYR	NOV		
Staphylococcus haemolyticus		70.82	0.45	ARG 92	1	98	2		
Staphylococcus hominis subsp. hominis		13.12	0.32	TRE 93	1	1	1		
Micrococcus caseolyticus		7.86	0.14		99	80	99		
102360	**Low discrimination**								**102360**
					NOV	OXI	YEP		
Kocuria varians		42.05	0.31	VPT 14	1	13	80		
Staphylococcus hominis subsp. novobiosepticus		30.25	0.18	SUC 12	99	1	1		
Micrococcus caseolyticus		16.00	0.11		99	99	99		
102400	**Low discrimination**								**102400**
					NOV				
Staphylococcus auricularis		60.75	0.69	VPT 12	1				
Staphylococcus cohnii subsp. cohnii		20.93	0.63		99				
Kocuria rosea		11.61	0.47		1				
102401	**Low discrimination**								**102401**
					NOV				
Staphylococcus cohnii subsp. cohnii		64.42	0.64	NIT 7	99				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		19.17	0.49		1			
Staphylococcus schleiferi subsp. schleiferi		9.40	0.39		1			
102402	**Low discrimination**							**102402**
					NOV	OXI		
Staphylococcus auricularis		77.87	0.49	VPT 12	1	1		
Kocuria varians		11.21	0.33	LAC 10	1	13		
Macrococcus caseolyticus		7.13	0.18		99	99		
102403	**Low discrimination**							**102403**
					OXI	NOV		
Staphylococcus auricularis		63.42	0.30	VPT 12	1	1		
Staphylococcus cohnii subsp. cohnii		19.37	0.23	MNS 10	1	99		
Kocuria kristinae		7.25	0.04	LAC 10	99	1		
102410	**Low discrimination**							**102410**
					NOV	YEP	OXI	
Staphylococcus cohnii subsp. cohnii		88.42	0.65	NIT 7	99	12	1	
Macrococcus bovicus		4.30	0.32		99	99	99	
Kocuria rosea		4.28	0.27		1	1	35	
102411	**Low discrimination**							**102411**
					NOV			
Staphylococcus cohnii subsp. cohnii		64.51	0.66	NIT 7	99			
Staphylococcus capitis subsp. capitis		35.11	0.52		1			
Staphylococcus auricularis		0.15	0.08		1			
102420	**Low discrimination**							**102420**
Kocuria rosea		70.97	0.39	TRE 1				
Kocuria varians		17.26	0.35					
Staphylococcus auricularis		8.76	0.28					
102440	**Low discrimination**							**102440**
					OXI	NOV	YEP	
Staphylococcus auricularis		53.10	0.77	VPT 12	1	1	1	
Staphylococcus cohnii subsp. cohnii		24.83	0.73		1	99	12	
Macrococcus caseolyticus		15.28	0.55		99	99	99	
102441	**Low discrimination**							**102441**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		78.55	0.74	NIT 7	1	99		
Staphylococcus auricularis		17.23	0.57		1	1		
Kocuria kristinae		3.38	0.36		99	1		
102442	**Low discrimination**							**102442**
					OXI	NOV		
Macrococcus caseolyticus		77.41	0.59	GAL 99	99	99		
Staphylococcus auricularis		19.93	0.57		1	1		
Staphylococcus haemolyticus		0.95	0.31		1	2		
102443	**Low discrimination**							**102443**
					OXI	NOV		
Staphylococcus auricularis		42.64	0.37	VPT 12	1	1		
Kocuria kristinae		18.79	0.23	MNS 10	99	1		
Staphylococcus cohnii subsp. cohnii		17.68	0.33	LAC 10	1	99		
102450	**Good identification**							**102450**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		93.28	0.75	NIT 7	1	99	12	
Macrococcus bovicus		3.35	0.39		99	99	99	
Staphylococcus auricularis		1.58	0.36		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
102451	**Very good identification**								**102451**
					NOV				
Staphylococcus cohnii subsp. cohnii		99.42	0.76	NIT 7	99				
Staphylococcus auricularis		0.17	0.16		1				
Staphylococcus capitis subsp. capitis		0.17	0.11		1				
102452	**Low discrimination**								**102452**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		39.44	0.34	NIT 7	1	99	1	12	
Micrococcus caseolyticus		28.59	0.18	LAC 1	99	99	80	99	
Staphylococcus haemolyticus		20.45	0.26		1	2	98	20	
102453	**Good identification**								**102453**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		96.64	0.35	NIT 7	1	99			
Kocuria kristinae		3.36	-.05	LAC 1	99	1			
102460	**Low discrimination**								**102460**
					OXI	NOV	YEP		
Staphylococcus auricularis		48.09	0.36	VPT 12	1	1	1		
Staphylococcus cohnii subsp. cohnii		22.49	0.32	XYL 1	1	99	12		
Micrococcus caseolyticus		13.84	0.14		99	99	99		
102461	**Low discrimination**								**102461**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		79.22	0.33	NIT 7	1	99			
Staphylococcus auricularis		17.37	0.16	XYL 1	1	1			
Kocuria kristinae		3.40	-.05		99	1			
102470	**Good identification**								**102470**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		96.54	0.34	NIT 7	1	12			
Micrococcus bovicus		3.46	-.02	XYL 1	99	99			
102471	**Good identification**								**102471**
Staphylococcus cohnii subsp. cohnii		100.0	0.35	NIT 7					
102500	**Low discrimination**								**102500**
					NOV	OXI			
Micrococcus caseolyticus		72.26	0.55	MLT 99	99	99			
Staphylococcus auricularis		11.96	0.49		1	1			
Kocuria varians		9.15	0.48		1	13			
102501	**Low discrimination**								**102501**
					NOV				
Staphylococcus schleiferi subsp. schleiferi		31.37	0.39	ARG 86	1				
Rothia mucilaginosa		28.39	0.32	PHS 99	1				
Staphylococcus cohnii subsp. cohnii		23.89	0.44		99				
102502	**Very good identification**								**102502**
					NOV	OXI			
Micrococcus caseolyticus		96.52	0.59	MLT 99	99	99			
Kocuria varians		2.30	0.37		1	13			
Staphylococcus auricularis		1.18	0.30		1	1			
102510	**Low discrimination**								**102510**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		73.13	0.46	NIT 7	1	12			
Micrococcus bovicus		13.73	0.24	GAL 10	99	99			
Micrococcus caseolyticus		10.18	0.14		99	99			
102511	**Acceptable identification**								**102511**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV			
Staphylococcus cohnii subsp. cohnii		93.52	0.46	NIT 7	99			
Staphylococcus capitis subsp. capitis		4.63	0.11	GAL 10	1			
Staphylococcus schleiferi subsp. schleiferi		0.98	-.02		1			
102520	**Low discrimination**							**102520**
					NOV	OXI		
Kocuria varians		77.39	0.39	VPT 14	1	13		
Macrococcus caseolyticus		17.56	0.14	TRE 7	99	99		
Staphylococcus auricularis		2.91	0.08		1	1		
102540	**Excellent identification**							**102540**
					OXI	NOV	YEP	
Macrococcus caseolyticus		99.30	0.96		99	99	99	
Staphylococcus auricularis		0.39	0.57		1	1	1	
Staphylococcus cohnii subsp. cohnii		0.18	0.54		1	99	12	
102541	**Low discrimination**							**102541**
					OXI	NOV	YEP	
Macrococcus caseolyticus		69.47	0.55	MNS 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		13.59	0.54		1	99	12	
Rothia mucilaginosa		11.90	0.39		1	1	1	
102542	**Excellent identification**							**102542**
					OXI	NOV		
Macrococcus caseolyticus		99.95	1.00		99	99		
Staphylococcus auricularis		0.03	0.37		1	1		
Staphylococcus haemolyticus		0.01	0.32		1	2		
102543	**Very good identification**							**102543**
					OXI	NOV		
Macrococcus caseolyticus		98.60	0.59	MNS 1	99	99		
Kocuria kristinae		0.41	0.14		99	1		
Staphylococcus lugdunensis		0.39	0.14		1	1		
102550	**Low discrimination**							**102550**
					OXI	YEP		
Macrococcus caseolyticus		76.31	0.55	MAN 1	99	99		
Staphylococcus cohnii subsp. cohnii		17.54	0.56		1	12		
Macrococcus bovicus		2.43	0.32		99	99		
102551	**Good identification**							**102551**
					OXI	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		92.79	0.57	NIT 7	1	1	12	
Macrococcus caseolyticus		3.76	0.14	GAL 10	99	80	99	
Staphylococcus saprophyticus subsp. bovis		2.16	0.18		1	99	80	
102552	**Very good identification**							**102552**
					OXI	NOV		
Macrococcus caseolyticus		98.66	0.59	MAN 1	99	99		
Staphylococcus haemolyticus		0.80	0.27		1	2		
Staphylococcus saprophyticus subsp. bovis		0.38	0.18		1	99		
102560	**Very good identification**							**102560**
					NOV	OXI		
Macrococcus caseolyticus		98.47	0.55	XYL 1	99	99		
Kocuria varians		0.96	0.25		1	13		
Staphylococcus auricularis		0.38	0.16		1	1		
102562	**Very good identification**							**102562**
					NOV	OXI		
Macrococcus caseolyticus		99.82	0.59	XYL 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		0.18	0.14		1	13		
102600	**Low discrimination**							**102600**
Staphylococcus auricularis		72.68	0.68	VPT 12				
Staphylococcus petrasii subsp. pragensis		19.02	0.39					
Staphylococcus haemolyticus		1.73	0.36					
102601	**Low discrimination**							**102601**
					OXI			
Kocuria kristinae		73.91	0.57	NIT 7	99			
Staphylococcus auricularis		13.57	0.49	MLT 90	1			
Rothia mucilaginosa		6.52	0.32		1			
102602	**Low discrimination**							**102602**
					OXI	NOV		
Staphylococcus auricularis		77.14	0.49	VPT 12	1	1		
Staphylococcus haemolyticus		9.32	0.31	LAC 10	1	2		
Macrococcus caseolyticus		5.10	0.14		99	99		
102603	**Low discrimination**							**102603**
					OXI			
Kocuria kristinae		89.39	0.45	NIT 7	99			
Staphylococcus auricularis		7.29	0.29	MLT 90	1			
Staphylococcus lugdunensis		2.52	0.14		1			
102610	**Low discrimination**							**102610**
					OXI	NOV		
Staphylococcus vitulinus		60.70	0.36	VPT 1	99	99		
Staphylococcus haemolyticus		11.52	0.31		1	2		
Staphylococcus auricularis		8.31	0.27		1	1		
102611	**Good identification**							**102611**
					OXI	NOV	PYR	
Staphylococcus capitis subsp. capitis		94.07	0.57	TRE 2	1	1	1	
Kocuria kristinae		4.56	0.29		99	1	93	
Staphylococcus cohnii subsp. cohnii		0.94	0.25		1	99	1	
102630	**Good identification**							**102630**
					NOV			
Staphylococcus vitulinus		99.45	0.45	VPT 1	99			
Kocuria rosea		0.55	0.00		1			
102640	**Low discrimination**							**102640**
Staphylococcus petrasii subsp. pragensis		77.97	0.80	NIT 10				
Staphylococcus haemolyticus		7.11	0.77					
Staphylococcus auricularis		7.02	0.76					
102641	**Low discrimination**							**102641**
					OXI			
Kocuria kristinae		87.00	0.77	NIT 7	99			
Staphylococcus petrasii subsp. pragensis		4.18	0.39		1			
Staphylococcus auricularis		4.14	0.56		1			
102642	**Low discrimination**							**102642**
					OXI	NOV		
Staphylococcus haemolyticus		42.03	0.72	ARG 92	1	2		
Macrococcus caseolyticus		23.00	0.55		99	99		
Staphylococcus petrasii subsp. jettensis		10.06	0.43		1	1		
102643	**Low discrimination**							**102643**
					OXI			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		72.48	0.65	NIT 7	99			
Staphylococcus lugdunensis		22.51	0.55		1			
Staphylococcus haemolyticus		2.19	0.41		1			
102650	**Low discrimination**							**102650**
					PYR			
Staphylococcus haemolyticus		77.46	0.72	ARG 92	98			
Staphylococcus petrasii subsp. pragensis		14.61	0.39		99			
Staphylococcus hominis subsp. hominis		1.71	0.40		1			
102651	**Low discrimination**							**102651**
					OXI	NOV	PYR	
Kocuria kristinae		74.44	0.49	NIT 7	99	1	93	
Staphylococcus haemolyticus		14.07	0.41	MAN 4	1	2	98	
Staphylococcus cohnii subsp. cohnii		5.40	0.35		1	99	1	
102652	**Good identification**							**102652**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		92.35	0.67	ARG 92	1	98	2	
Staphylococcus hominis subsp. hominis		4.24	0.41		1	1	1	
Macrococcus caseolyticus		0.87	0.14		99	80	99	
102653	**Low discrimination**							**102653**
					OXI			
Kocuria kristinae		65.27	0.36	NIT 7	99			
Staphylococcus haemolyticus		27.76	0.36	MAN 4	1			
Staphylococcus lugdunensis		4.91	0.14		1			
102660	**Low discrimination**							**102660**
Staphylococcus petrasii subsp. pragensis		78.00	0.39	NIT 10				
Staphylococcus haemolyticus		7.11	0.36	XYL 1				
Staphylococcus auricularis		7.03	0.35					
102661	**Low discrimination**							**102661**
					OXI			
Kocuria kristinae		88.34	0.36	NIT 7	99			
Staphylococcus petrasii subsp. pragensis		4.25	-.02	XYL 1	1			
Staphylococcus auricularis		4.21	0.15		1			
102662	**Low discrimination**							**102662**
					OXI	NOV		
Staphylococcus haemolyticus		42.38	0.31	ARG 92	1	2		
Macrococcus caseolyticus		23.20	0.14	XYL 1	99	99		
Staphylococcus petrasii subsp. jettensis		10.15	0.02		1	1		
102670	**Low discrimination**							**102670**
					OXI	PYR		
Staphylococcus haemolyticus		75.05	0.31	ARG 92	1	98		
Staphylococcus petrasii subsp. pragensis		14.16	-.02	XYL 1	1	99		
Staphylococcus vitulinus		10.80	0.04		99	1		
102700	**Low discrimination**							**102700**
					OXI	NOV		
Rothia mucilaginosa		73.25	0.63	ESL 90	1	1		
Macrococcus caseolyticus		20.15	0.52		99	99		
Staphylococcus auricularis		4.62	0.49		1	1		
102701	**Very good identification**							**102701**
					OXI			
Rothia mucilaginosa		95.98	0.73	ESL 90	1			
Kocuria kristinae		3.66	0.47		99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		0.22	0.29		1			
102702	**Good identification**							**102702**
					OXI	NOV		
Macrococcus caseolyticus		93.33	0.55	MLT 99	99	99		
Rothia mucilaginosa		2.28	0.22		1	1		
Staphylococcus haemolyticus		1.94	0.32		1	2		
102703	**Low discrimination**							**102703**
					OXI	NOV		
Rothia mucilaginosa		45.48	0.32	ESL 90	1	1		
Kocuria kristinae		42.96	0.35	LAC 1	99	1		
Macrococcus caseolyticus		6.26	0.14		99	99		
102711	**Low discrimination**							**102711**
					OXI	PYR		
Rothia mucilaginosa		74.25	0.32	ESL 90	1	75		
Kocuria kristinae		11.69	0.19	MAN 1	99	93		
Staphylococcus capitis subsp. capitis		7.31	0.16		1	1		
102712	**Low discrimination**							**102712**
					OXI	NOV		
Staphylococcus haemolyticus		54.75	0.27	ARG 92	1	2		
Macrococcus caseolyticus		45.25	0.14	MLT 99	99	99		
102713	**Low discrimination**							**102713**
					OXI	NOV	PYR	
Staphylococcus lentus		81.15	0.30	VPT 7	99	99	6	
Kocuria kristinae		15.00	0.07	ESL 99	99	1	93	
Rothia mucilaginosa		3.85	-.09		1	1	75	
102721	**Good identification**							**102721**
					OXI			
Rothia mucilaginosa		96.32	0.32	ESL 90	1			
Kocuria kristinae		3.68	0.06	XYL 1	99			
102733	**Good identification**							**102733**
Staphylococcus lentus		100.0	0.30	VPT 7				
102740	**Low discrimination**							**102740**
					OXI	NOV		
Macrococcus caseolyticus		86.37	0.93		99	99		
Rothia mucilaginosa		7.40	0.71		1	1		
Staphylococcus haemolyticus		4.79	0.78		1	2		
102741	**Low discrimination**							**102741**
					OXI	NOV		
Rothia mucilaginosa		83.28	0.80	ESL 90	1	1		
Kocuria kristinae		12.26	0.67		99	1		
Macrococcus caseolyticus		3.27	0.52		99	99		
102742	**Very good identification**							**102742**
					OXI	PYR	NOV	
Macrococcus caseolyticus		97.49	0.96		99	80	99	
Staphylococcus haemolyticus		2.03	0.73		1	98	2	
Staphylococcus hominis subsp. hominis		0.35	0.59		1	1	1	
102743	**Low discrimination**							**102743**
					OXI	NOV		
Macrococcus caseolyticus		40.58	0.55	MNS 1	99	99		
Kocuria kristinae		25.33	0.55		99	1		
Staphylococcus lugdunensis		23.60	0.55		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
102750	**Low discrimination**							**102750**
					OXI	NOV		
Staphylococcus haemolyticus		70.82	0.73	ARG 92	1	2		
Macrococcus caseolyticus		21.95	0.52		99	99		
Staphylococcus saprophyticus subsp. bovis		3.06	0.43		1	99		
102751	**Low discrimination**							**102751**
					OXI			
Rothia mucilaginosa		44.16	0.39	ESL 90	1			
Kocuria kristinae		26.81	0.39	MAN 1	99			
Staphylococcus haemolyticus		17.15	0.42		1			
102752	**Low discrimination**							**102752**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		52.87	0.68	ARG 92	1	98	2	
Macrococcus caseolyticus		43.70	0.55		99	80	99	
Staphylococcus hominis subsp. hominis		2.33	0.42		1	1	1	
102753	**Low discrimination**							**102753**
					OXI	NOV	PYR	
Staphylococcus haemolyticus		31.05	0.37	ARG 92	1	2	98	
Staphylococcus lentus		26.33	0.36	MNS 3	99	99	6	
Kocuria kristinae		21.58	0.26		99	1	93	
102760	**Low discrimination**							**102760**
					OXI	NOV		
Macrococcus caseolyticus		86.31	0.52	XYL 1	99	99		
Rothia mucilaginosa		7.39	0.30		1	1		
Staphylococcus haemolyticus		4.79	0.37		1	2		
102761	**Low discrimination**							**102761**
					OXI	NOV		
Rothia mucilaginosa		83.81	0.39	ESL 90	1	1		
Kocuria kristinae		12.34	0.26	XYL 1	99	1		
Macrococcus caseolyticus		3.29	0.11		99	99		
102762	**Very good identification**							**102762**
					OXI	PYR	NOV	
Macrococcus caseolyticus		97.62	0.55	XYL 1	99	80	99	
Staphylococcus haemolyticus		2.03	0.32		1	98	2	
Staphylococcus hominis subsp. hominis		0.35	0.18		1	1	1	
102770	**Low discrimination**							**102770**
					OXI	NOV		
Staphylococcus haemolyticus		73.91	0.32	ARG 92	1	2		
Macrococcus caseolyticus		22.91	0.11	XYL 1	99	99		
Staphylococcus saprophyticus subsp. bovis		3.19	0.02		1	99		
102772	**Low discrimination**							**102772**
					OXI	NOV		
Staphylococcus haemolyticus		54.75	0.27	ARG 92	1	2		
Macrococcus caseolyticus		45.25	0.14	XYL 1	99	99		
102773	**Acceptable identification**							**102773**
					OXI			
Staphylococcus lentus		94.67	0.36	VPT 7	99			
Staphylococcus xylosus		5.33	0.04	ESL 99	1			
103000	**Low discrimination**							**103000**
					OXI			
Kocuria rosea		85.43	0.47	ESL 1	35			
Dermacoccus nishinomiyaensis		8.79	0.28		93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		3.40	0.27		13		
103010	**Low discrimination**						**103010**
					NOV	YEP	
Kocuria rosea		75.02	0.27	ESL 1	1	1	
Staphylococcus vitulinus		11.69	0.09	MAN 10	99	75	
Macrococcus bovicus		5.59	0.08		99	99	
103011	**Good identification**						**103011**
Staphylococcus capitis subsp. capitis		99.14	0.45	ESL 1			
Kocuria rosea		0.87	0.01				
103020	**Good identification**						**103020**
Kocuria rosea		96.84	0.39	ESL 1			
Kocuria varians		3.16	0.17				
103100	**Low discrimination**						**103100**
					OXI		
Kocuria varians		59.51	0.30	VPT 14	13		
Dermacoccus nishinomiyaensis		19.53	0.13	ESL 1	93		
Kocuria rosea		10.07	0.05		35		
103200	**Low discrimination**						**103200**
					NOV		
Kocuria rosea		66.44	0.27	ESL 1	1		
Staphylococcus vitulinus		10.36	0.09	SUC 10	99		
Macrococcus carouselicus		8.82	0.10		99		
103201	**Low discrimination**						**103201**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		58.02	0.28	ESL 1	1	1	
Kocuria kristinae		23.83	0.19	MAN 93	99	93	
Rothia mucilaginosa		14.53	0.10		1	75	
103210	**Good identification**						**103210**
					NOV	OXI	
Staphylococcus vitulinus		98.04	0.50	VPT 1	99	99	
Staphylococcus capitis subsp. capitis		0.72	0.10		1	1	
Kocuria rosea		0.71	0.07		1	35	
103211	**Good identification**						**103211**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		98.28	0.51	ESL 1	1	1	
Staphylococcus vitulinus		1.37	0.09		99	99	
Staphylococcus sciuri		0.35	0.05		99	99	
103230	**Very good identification**						**103230**
					NOV		
Staphylococcus vitulinus		99.89	0.59	VPT 1	99		
Kocuria rosea		0.11	0.00		1		
103240	**Low discrimination**						**103240**
					NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		69.54	0.27	URE 99	99	1	
Rothia mucilaginosa		9.88	0.08	ESL 8	1	75	
Staphylococcus petrasii subsp. pragensis		6.92	-.02		1	99	
103241	**Low discrimination**						**103241**
					OXI	NOV	
Kocuria kristinae		79.95	0.38	NIT 7	99	1	
Rothia mucilaginosa		12.64	0.18	TRE 95	1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. novobiosepticus		7.41	0.15		1	99		
103300	**Good identification**							**103300**
					NOV	OXI		
Rothia mucilaginosa		97.68	0.42	TRE 99	1	1		
Kocuria varians		1.03	0.13		1	13		
Macrococcus carouselicus		0.80	0.02		99	99		
103301	**Good identification**							**103301**
					OXI			
Rothia mucilaginosa		99.45	0.51	TRE 99	1			
Kocuria kristinae		0.55	0.09		99			
103313	**Good identification**							**103313**
Staphylococcus lentus		100.0	0.30	VPT 7				
103333	**Good identification**							**103333**
Staphylococcus lentus		100.0	0.30	VPT 7				
103340	**Low discrimination**							**103340**
					OXI	NOV	YEP	
Rothia mucilaginosa		80.91	0.49	TRE 99	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		17.26	0.37		1	99	1	
Macrococcus caseolyticus		1.06	0.11		99	99	99	
103341	**Very good identification**							**103341**
					OXI	NOV		
Rothia mucilaginosa		96.11	0.59	TRE 99	1	1		
Kocuria kristinae		2.05	0.28		99	1		
Staphylococcus hominis subsp. novobiosepticus		1.71	0.25		1	99		
103342	**Low discrimination**							**103342**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. novobiosepticus		73.12	0.32	URE 99	1	99	1	
Macrococcus caseolyticus		12.51	0.14	ESL 8	99	99	99	
Rothia mucilaginosa		6.43	0.08		1	1	1	
103353	**Good identification**							**103353**
Staphylococcus lentus		95.44	0.36	VPT 7				
Staphylococcus sciuri		4.56	0.07	TRE 99				
103373	**Good identification**							**103373**
Staphylococcus lentus		96.52	0.36	VPT 7				
Staphylococcus sciuri		3.48	0.04	TRE 99				
103400	**Low discrimination**							**103400**
					OXI	NOV		
Macrococcus carouselicus		55.31	0.28	VPT 1	99	99		
Staphylococcus auricularis		25.68	0.28	NIT 6	1	1		
Staphylococcus cohnii subsp. cohnii		8.85	0.22		1	99		
103440	**Low discrimination**							**103440**
					OXI	NOV	YEP	
Staphylococcus auricularis		49.83	0.36	VPT 12	1	1	1	
Staphylococcus cohnii subsp. cohnii		23.31	0.32	ESL 1	1	99	12	
Macrococcus caseolyticus		14.34	0.14		99	99	99	
103450	**Low discrimination**							**103450**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		69.49	0.34	ESL 1	1	12		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		27.41	0.20	NIT 7	99	99		
Macrococcus equipersicus		3.09	0.02		99	99		
103451	**Acceptable identification**							**103451**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		90.57	0.35	ESL 1	1	99	1	
Staphylococcus sciuri		6.27	0.07	NIT 7	99	99	1	
Kocuria kristinae		3.15	-.05		99	1	93	
103500	**Low discrimination**							**103500**
					OXI	NOV		
Rothia mucilaginosa		89.10	0.42	SUC 99	1	1		
Macrococcus carouselicus		5.62	0.20		99	99		
Macrococcus caseolyticus		4.09	0.14		99	99		
103501	**Good identification**							**103501**
					OXI			
Rothia mucilaginosa		99.77	0.51	SUC 99	1			
Staphylococcus schleiferi subsp. schleiferi		0.12	-.02		1			
Kocuria kristinae		0.11	-.06		99			
103513	**Good identification**							**103513**
Staphylococcus lentus		100.0	0.30	VPT 7				
103533	**Good identification**							**103533**
Staphylococcus lentus		100.0	0.30	VPT 7				
103540	**Low discrimination**							**103540**
					OXI	NOV		
Macrococcus caseolyticus		65.80	0.55	ESL 1	99	99		
Rothia mucilaginosa		33.82	0.49		1	1		
Staphylococcus auricularis		0.26	0.16		1	1		
103541	**Very good identification**							**103541**
					OXI	NOV		
Rothia mucilaginosa		98.82	0.59	SUC 99	1	1		
Macrococcus caseolyticus		0.65	0.14		99	99		
Kocuria kristinae		0.40	0.14		99	1		
103542	**Very good identification**							**103542**
					OXI	NOV		
Macrococcus caseolyticus		99.66	0.59	ESL 1	99	99		
Rothia mucilaginosa		0.35	0.08		1	1		
103553	**Good identification**							**103553**
					OXI	NOV		
Staphylococcus lentus		97.33	0.36	VPT 7	99	99		
Aerococcus viridans		2.67	0.18	SUC 99	1	1		
103573	**Good identification**							**103573**
					OXI	NOV		
Staphylococcus lentus		97.33	0.36	VPT 7	99	99		
Aerococcus viridans		2.67	0.18	SUC 99	1	1		
103600	**Low discrimination**							**103600**
					OXI	NOV		
Rothia mucilaginosa		71.41	0.42	GAL 99	1	1		
Macrococcus carouselicus		21.33	0.34		99	99		
Staphylococcus auricularis		4.50	0.27		1	1		
103601	**Low discrimination**							**103601**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Rothia mucilaginosa		75.81	0.51	GAL 99	1			
Kocuria kristinae		23.87	0.45		99			
Staphylococcus auricularis		0.18	0.08		1			
103603	**Low discrimination**							**103603**
					OXI			
Kocuria kristinae		88.63	0.32	NIT 7	99			
Rothia mucilaginosa		11.38	0.10	MLT 90	1			
103610	**Low discrimination**							**103610**
Staphylococcus vitulinus		84.52	0.45	VPT 1				
Macrococcus carouselicus		12.95	0.30					
Staphylococcus sciuri		1.57	0.18					
103611	**Low discrimination**							**103611**
					OXI	NOV	PYR	
Staphylococcus sciuri		58.11	0.36	VPT 1	99	99	1	
Kocuria kristinae		12.21	0.16		99	1	93	
Staphylococcus capitis subsp. capitis		10.18	0.16		1	1	1	
103613	**Low discrimination**							**103613**
					NOV	PYR		
Staphylococcus lentus		63.76	0.30	VPT 7	99	6		
Staphylococcus sciuri		27.40	0.20	GAL 99	99	1		
Kocuria kristinae		8.84	0.04		1	93		
103630	**Very good identification**							**103630**
Staphylococcus vitulinus		99.48	0.54	VPT 1				
Staphylococcus sciuri		0.52	0.15					
103631	**Low discrimination**							**103631**
Staphylococcus sciuri		72.34	0.34	VPT 1				
Staphylococcus vitulinus		17.42	0.13					
Staphylococcus lentus		10.24	0.18					
103633	**Low discrimination**							**103633**
Staphylococcus lentus		75.52	0.30	VPT 7				
Staphylococcus sciuri		24.49	0.18	GAL 99				
103640	**Low discrimination**							**103640**
Rothia mucilaginosa		51.78	0.49	GAL 99				
Staphylococcus petrasii subsp. pragensis		36.25	0.39					
Staphylococcus haemolyticus		3.31	0.36					
103641	**Low discrimination**							**103641**
					OXI	NOV	PYR	
Kocuria kristinae		54.63	0.65	NIT 7	99	1	93	
Rothia mucilaginosa		44.99	0.59		1	1	75	
Staphylococcus sciuri		0.14	0.17		99	99	1	
103642	**Low discrimination**							**103642**
					OXI	NOV		
Staphylococcus haemolyticus		34.52	0.31	ARG 92	1	2		
Macrococcus caseolyticus		18.89	0.14	ESL 1	99	99		
Rothia mucilaginosa		9.71	0.08		1	1		
103643	**Good identification**							**103643**
					OXI			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		95.47	0.52	NIT 7	99			
Rothia mucilaginosa		3.18	0.18		1			
Staphylococcus lugdunensis		1.20	0.14		1			
103650	**Low discrimination**							**103650**
					OXI	NOV		
Staphylococcus haemolyticus		36.42	0.31	ARG 92	1	2		
Staphylococcus sciuri		36.27	0.30	ESL 1	99	99		
Rothia mucilaginosa		9.81	0.08		1	1		
103651	**Low discrimination**							**103651**
					OXI	NOV	PYR	
Staphylococcus sciuri		60.22	0.48	VPT 1	99	99	1	
Kocuria kristinae		30.26	0.36		99	1	93	
Rothia mucilaginosa		6.04	0.18		1	1	75	
103653	**Low discrimination**							**103653**
					NOV	PYR		
Staphylococcus lentus		41.01	0.36	VPT 7	99	6		
Staphylococcus sciuri		32.66	0.32	GAL 99	99	1		
Kocuria kristinae		25.21	0.24		1	93		
103670	**Low discrimination**							**103670**
Staphylococcus sciuri		65.88	0.27	VPT 1				
Staphylococcus vitulinus		34.12	0.13	MNS 89				
103671	**Acceptable identification**							**103671**
					NOV	PYR		
Staphylococcus sciuri		92.32	0.46	VPT 1	99	1		
Staphylococcus lentus		7.06	0.25		99	6		
Kocuria kristinae		0.62	-.05		1	93		
103673	**Low discrimination**							**103673**
					OXI	NOV		
Staphylococcus lentus		61.42	0.36	VPT 7	99	99		
Staphylococcus sciuri		36.90	0.30	GAL 99	99	99		
Aerococcus viridans		1.68	0.18		1	1		
103700	**Very good identification**							**103700**
					OXI	NOV		
Rothia mucilaginosa		99.83	0.83		1	1		
Macrococcus carouselicus		0.13	0.27		99	99		
Macrococcus caseolyticus		0.03	0.11		99	99		
103701	**Very good identification**							**103701**
					OXI	NOV	PYR	
Rothia mucilaginosa		99.88	0.92		1	1	75	
Kocuria kristinae		0.11	0.35		99	1	93	
Staphylococcus lentus		0.01	0.18		99	99	6	
103702	**Good identification**							**103702**
					OXI	NOV		
Rothia mucilaginosa		95.62	0.42	LAC 1	1	1		
Macrococcus caseolyticus		4.38	0.14		99	99		
103703	**Good identification**							**103703**
					OXI	NOV	PYR	
Rothia mucilaginosa		95.25	0.51	LAC 1	1	1	75	
Kocuria kristinae		2.50	0.23		99	1	93	
Staphylococcus lentus		2.25	0.30		99	99	6	
103710	**Low discrimination**							**103710**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Rothia mucilaginosa		86.25	0.42	MAN 1	1	1		
Macrococcus carouselicus		7.06	0.23		99	99		
Staphylococcus sciuri		3.88	0.24		99	99		
103711	**Low discrimination**							**103711**
					OXI	NOV		
Rothia mucilaginosa		55.99	0.51	MAN 1	1	1		
Staphylococcus lentus		36.98	0.59		99	99		
Staphylococcus sciuri		6.79	0.42		99	99		
103712	**Low discrimination**							**103712**
					OXI	NOV		
Staphylococcus lentus		80.29	0.30	VPT 7	99	99		
Rothia mucilaginosa		11.43	0.00	MNS 99	1	1		
Staphylococcus sciuri		8.28	0.07		99	99		
103713	**Very good identification**							**103713**
					OXI	NOV		
Staphylococcus lentus		98.70	0.71	VPT 7	99	99		
Staphylococcus sciuri		0.83	0.26		99	99		
Rothia mucilaginosa		0.43	0.10		1	1		
103720	**Good identification**							**103720**
Rothia mucilaginosa		100.0	0.42	XYL 1				
103721	**Good identification**							**103721**
					OXI	NOV		
Rothia mucilaginosa		98.96	0.51	XYL 1	1	1		
Staphylococcus lentus		0.66	0.18		99	99		
Staphylococcus sciuri		0.28	0.09		99	99		
103723	**Low discrimination**							**103723**
					OXI	NOV		
Staphylococcus lentus		70.07	0.30	VPT 7	99	99		
Rothia mucilaginosa		29.93	0.10	MAN 99	1	1		
103731	**Low discrimination**							**103731**
					OXI	NOV		
Staphylococcus lentus		86.67	0.59	VPT 7	99	99		
Staphylococcus sciuri		12.00	0.40		99	99		
Rothia mucilaginosa		1.33	0.10		1	1		
103732	**Acceptable identification**							**103732**
Staphylococcus lentus		92.78	0.30	VPT 7				
Staphylococcus sciuri		7.22	0.05	MNS 99				
103733	**Very good identification**							**103733**
Staphylococcus lentus		99.37	0.71	VPT 7				
Staphylococcus sciuri		0.63	0.24					
103740	**Very good identification**							**103740**
					OXI	NOV		
Rothia mucilaginosa		98.60	0.90		1	1		
Macrococcus caseolyticus		1.29	0.52		99	99		
Staphylococcus haemolyticus		0.07	0.37		1	2		
103741	**Excellent identification**							**103741**
					OXI	NOV	PYR	
Rothia mucilaginosa		99.58	1.00		1	1	75	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		0.41	0.55		99	1	93	
Staphylococcus lentus		0.01	0.25		99	99	6	
103742	**Low discrimination**							**103742**
					OXI	NOV		
Macrococcus caseolyticus		64.93	0.55	ESL 1	99	99		
Rothia mucilaginosa		33.38	0.49		1	1		
Staphylococcus haemolyticus		1.35	0.32		1	2		
103743	**Low discrimination**							**103743**
					OXI	NOV	PYR	
Rothia mucilaginosa		88.25	0.59	LAC 1	1	1	75	
Kocuria kristinae		8.93	0.42		99	1	93	
Staphylococcus lentus		1.82	0.36		99	99	6	
103750	**Low discrimination**							**103750**
					OXI	NOV		
Rothia mucilaginosa		87.05	0.49	MAN 1	1	1		
Staphylococcus sciuri		6.31	0.36		99	99		
Staphylococcus haemolyticus		3.68	0.32		1	2		
103751	**Low discrimination**							**103751**
					OXI	NOV		
Rothia mucilaginosa		55.95	0.59	MAN 1	1	1		
Staphylococcus lentus		32.15	0.66		99	99		
Staphylococcus sciuri		10.93	0.54		99	99		
103752	**Low discrimination**							**103752**
					OXI	YEP	NOV	
Staphylococcus lentus		48.59	0.36	VPT 7	99	1	99	
Staphylococcus haemolyticus		18.71	0.27	MNS 99	1	20	2	
Macrococcus caseolyticus		15.47	0.14		99	99	99	
103753	**Very good identification**							**103753**
					OXI	NOV		
Staphylococcus lentus		97.76	0.77	VPT 7	99	99		
Staphylococcus sciuri		1.53	0.38		99	99		
Rothia mucilaginosa		0.49	0.18		1	1		
103760	**Good identification**							**103760**
					OXI	NOV		
Rothia mucilaginosa		98.54	0.49	XYL 1	1	1		
Macrococcus caseolyticus		1.29	0.11		99	99		
Staphylococcus sciuri		0.17	0.02		99	99		
103761	**Very good identification**							**103761**
					OXI	NOV		
Rothia mucilaginosa		98.58	0.59	XYL 1	1	1		
Staphylococcus lentus		0.57	0.25		99	99		
Staphylococcus sciuri		0.45	0.21		99	99		
103763	**Low discrimination**							**103763**
					OXI	NOV	PYR	
Staphylococcus lentus		63.43	0.36	VPT 7	99	99	6	
Rothia mucilaginosa		31.13	0.18	MAN 99	1	1	75	
Kocuria kristinae		3.15	0.01		99	1	93	
103770	**Low discrimination**							**103770**
					OXI	NOV		
Staphylococcus sciuri		66.52	0.33	VPT 1	99	99		
Staphylococcus lentus		21.19	0.25	MNS 89	99	99		
Rothia mucilaginosa		12.29	0.08		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
103771	**Low discrimination**							**103771**
					OXI	NOV		
Staphylococcus lentus		78.48	0.66	VPT 7	99	99		
Staphylococcus sciuri		20.14	0.52		99	99		
Rothia mucilaginosa		1.38	0.18		1	1		
103772	**Low discrimination**							**103772**
Staphylococcus lentus		87.40	0.36	VPT 7				
Staphylococcus sciuri		12.60	0.17	MNS 99				
103773	**Very good identification**							**103773**
					OXI	NOV		
Staphylococcus lentus		98.81	0.77	VPT 7	99	99		
Staphylococcus sciuri		1.16	0.35		99	99		
Aerococcus viridans		0.03	0.18		1	1		
104000	**Low discrimination**							**104000**
Dermacoccus nishinomiyaensis		67.62	0.47	VPT 5				
Micrococcus sp.		21.47	0.28	PHS 5				
Kocuria rosea		7.88	0.27					
104001	**Low discrimination**							**104001**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		89.41	0.51	ARG 86	1			
Staphylococcus capitis subsp. capitis		6.52	0.26	NIT 90	1			
Staphylococcus cohnii subsp. cohnii		2.22	0.25		99			
104010	**Low discrimination**							**104010**
					NOV			
Staphylococcus cohnii subsp. cohnii		42.02	0.27	PHS 9	99			
Kocuria rosea		17.03	0.07	TRE 99	1			
Staphylococcus capitis subsp. capitis		14.12	0.08		1			
104011	**Good identification**							**104011**
					NOV			
Staphylococcus capitis subsp. capitis		95.88	0.49	NIT 89	1			
Staphylococcus cohnii subsp. cohnii		3.12	0.27	PHS 11	99			
Staphylococcus schleiferi subsp. schleiferi		1.00	0.10		1			
104040	**Low discrimination**							**104040**
					COA	NOV	OXI	
Staphylococcus cohnii subsp. cohnii		43.55	0.35	PHS 9	1	99	1	
Staphylococcus aureus subsp. anaerobius		37.84	0.16	TRE 99	99	1	1	
Micrococcus sp.		13.08	0.05		1	1	89	
104041	**Low discrimination**							**104041**
					NOV			
Staphylococcus cohnii subsp. cohnii		88.61	0.35	PHS 9	99			
Staphylococcus schleiferi subsp. schleiferi		11.40	0.10	TRE 99	1			
104050	**Low discrimination**							**104050**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		76.25	0.37	PHS 9	1	12		
Macrococcus bovicus		12.49	0.14	TRE 99	99	99		
Staphylococcus kloosii		11.27	0.32		1	99		
104051	**Acceptable identification**							**104051**
					NOV			
Staphylococcus cohnii subsp. cohnii		91.08	0.38	PHS 9	99			
Staphylococcus capitis subsp. capitis		8.92	0.08	TRE 99	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
104052	**Good identification**							**104052**
Staphylococcus kloosii		100.0	0.33	VPT 10				
104100	**Low discrimination**							**104100**
					OXI			
Dermacoccus nishinomiyaensis		86.43	0.33	VPT 5	93			
Kocuria varians		5.82	0.16	PHS 5	13			
Staphylococcus schleiferi subsp. schleiferi		5.03	0.10		1			
104101	**Good identification**							**104101**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		99.73	0.51	ARG 86	1			
Staphylococcus cohnii subsp. cohnii		0.28	0.05	NIT 90	99			
104150	**Low discrimination**							**104150**
					OXI	YEP		
Staphylococcus kloosii		61.28	0.38	VPT 10	1	99		
Staphylococcus cohnii subsp. cohnii		23.73	0.17	TRE 99	1	12		
Macrococcus bovicus		14.99	0.07		99	99		
104152	**Good identification**							**104152**
Staphylococcus kloosii		100.0	0.39	VPT 10				
104172	**Good identification**							**104172**
Staphylococcus kloosii		100.0	0.27	VPT 10				
104201	**Low discrimination**							**104201**
					OXI	PYR		
Staphylococcus capitis subsp. capitis		81.94	0.32	NIT 89	1	1		
Kocuria kristinae		11.95	0.13	PHS 11	99	93		
Staphylococcus schleiferi subsp. schleiferi		6.11	0.10	MAN 93	1	37		
104211	**Very good identification**							**104211**
Staphylococcus capitis subsp. capitis		100.0	0.55	NIT 89				
104240	**Very good identification**							**104240**
					PYR	COA		
Staphylococcus aureus subsp. anaerobius		98.08	0.57	VPT 1	1	99		
Staphylococcus petrasii subsp. pragensis		1.73	0.18		99	1		
Staphylococcus epidermidis		0.19	0.14		1	1		
104241	**Low discrimination**							**104241**
					OXI	COA	PYR	
Kocuria kristinae		67.58	0.33	PHS 1	99	1	93	
Staphylococcus aureus subsp. anaerobius		23.95	0.16	TRE 95	1	99	1	
Staphylococcus epidermidis		8.47	0.20		1	1	1	
104243	**Low discrimination**							**104243**
					OXI	PYR		
Staphylococcus epidermidis		61.37	0.30	URE 92	1	1		
Kocuria kristinae		38.63	0.20	NIT 97	99	93		
104340	**Low discrimination**							**104340**
					COA	NOV		
Staphylococcus epidermidis		50.10	0.30	URE 92	1	1		
Staphylococcus aureus subsp. anaerobius		46.40	0.16	NIT 97	99	1		
Staphylococcus hominis subsp. novobiosepticus		3.50	-.02		1	99		
104341	**Low discrimination**							**104341**
					OXI	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		68.07	0.35	URE 92	1	1		
Kocuria kristinae		31.93	0.23	NIT 97	99	93		
104342	**Good identification**							**104342**
					NOV			
Staphylococcus epidermidis		98.83	0.40	URE 92	1			
Staphylococcus hominis subsp. novobiosepticus		1.17	-.07	NIT 97	99			
104343	**Good identification**							**104343**
					OXI	PYR		
Staphylococcus epidermidis		96.43	0.46	URE 92	1	1		
Kocuria kristinae		3.57	0.10	NIT 97	99	93		
104400	**Very good identification**							**104400**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		97.92	0.65	PHS 9	1	99		
Staphylococcus schleiferi subsp. schleiferi		1.31	0.19		1	1		
Dermacoccus nishinomiyaensis		0.40	0.06		93	80		
104401	**Low discrimination**							**104401**
					OXI	NOV		
Staphylococcus schleiferi subsp. schleiferi		54.95	0.61	ARG 86	1	1		
Staphylococcus cohnii subsp. cohnii		44.98	0.66	NIT 90	1	99		
Kocuria kristinae		0.07	-.02		99	1		
104410	**Very good identification**							**104410**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		97.28	0.68	PHS 9	1	12		
Staphylococcus kloosii		2.40	0.47		1	99		
Macrococcus bovicus		0.33	0.10		99	99		
104411	**Very good identification**							**104411**
					NOV			
Staphylococcus cohnii subsp. cohnii		98.42	0.68	PHS 9	99			
Staphylococcus schleiferi subsp. schleiferi		0.95	0.19		1			
Staphylococcus capitis subsp. capitis		0.62	0.14		1			
104412	**Low discrimination**							**104412**
					YEP			
Staphylococcus kloosii		72.54	0.48	VPT 10	99			
Staphylococcus cohnii subsp. cohnii		27.46	0.27	MLT 95	12			
104413	**Good identification**							**104413**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	PHS 9				
104430	**Low discrimination**							**104430**
					YEP			
Staphylococcus cohnii subsp. cohnii		62.13	0.27	PHS 9	12			
Staphylococcus kloosii		37.87	0.35	XYL 1	99			
104431	**Good identification**							**104431**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	PHS 9				
104432	**Good identification**							**104432**
Staphylococcus kloosii		100.0	0.35	VPT 10				
104440	**Very good identification**							**104440**
					NOV	YEP	PYR	
Staphylococcus cohnii subsp. cohnii		98.13	0.76	PHS 9	99	12	1	
Staphylococcus petrasii subsp. pragensis		1.50	0.18		1	1	99	
Staphylococcus kloosii		0.19	0.32		99	99	52	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
104441	**Very good identification**						**104441**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		99.18	0.76	PHS 9	1	99	
Kocuria kristinae		0.44	0.18		99	1	
Staphylococcus schleiferi subsp. schleiferi		0.39	0.19		1	1	
104442	**Low discrimination**						**104442**
					YEP		
Staphylococcus cohnii subsp. cohnii		83.07	0.35	PHS 9	12		
Staphylococcus kloosii		16.93	0.33	LAC 1	99		
104443	**Acceptable identification**						**104443**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		90.21	0.35	PHS 9	1	99	
Kocuria kristinae		9.80	0.05	LAC 1	99	1	
104450	**Low discrimination**						**104450**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		86.94	0.78	PHS 9	1	12	
Staphylococcus kloosii		12.85	0.73		1	99	
Macrococcus bovicus		0.22	0.18		99	99	
104451	**Very good identification**						**104451**
					YEP		
Staphylococcus cohnii subsp. cohnii		99.86	0.79	PHS 9	12		
Staphylococcus kloosii		0.14	0.32		99		
104452	**Good identification**						**104452**
					YEP		
Staphylococcus kloosii		94.07	0.74	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		5.94	0.37		12		
104453	**Low discrimination**						**104453**
					YEP		
Staphylococcus cohnii subsp. cohnii		84.20	0.38	PHS 9	12		
Staphylococcus kloosii		12.44	0.33	LAC 1	99		
Staphylococcus cohnii subsp. urealyticum		3.35	0.10		64		
104460	**Good identification**						**104460**
					YEP		
Staphylococcus cohnii subsp. cohnii		95.51	0.35	PHS 9	12		
Staphylococcus kloosii		4.49	0.20	XYL 1	99		
104461	**Good identification**						**104461**
Staphylococcus cohnii subsp. cohnii		100.0	0.35	PHS 9			
104470	**Low discrimination**						**104470**
					YEP		
Staphylococcus kloosii		78.53	0.61	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		21.47	0.37		12		
104471	**Good identification**						**104471**
					YEP		
Staphylococcus cohnii subsp. cohnii		96.70	0.38	PHS 9	12		
Staphylococcus kloosii		3.30	0.20	XYL 1	99		
104472	**Very good identification**						**104472**
Staphylococcus kloosii		100.0	0.62	VPT 10			
104500	**Low discrimination**						**104500**
					NOV		
Staphylococcus cohnii subsp. cohnii		89.26	0.46	PHS 9	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. schleiferi		10.74	0.19	GAL 10	1		
104501	**Good identification**						**104501**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		91.66	0.61	ARG 86	1		
Staphylococcus cohnii subsp. cohnii		8.34	0.47	NIT 90	99		
104510	**Low discrimination**						**104510**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		69.29	0.48	PHS 9	1	12	
Staphylococcus kloosii		29.81	0.53	GAL 10	1	99	
Macrococcus bovicus		0.90	0.03		99	99	
104511	**Acceptable identification**						**104511**
					NOV		
Staphylococcus cohnii subsp. cohnii		91.97	0.49	PHS 9	99		
Staphylococcus schleiferi subsp. schleiferi		8.03	0.19	GAL 10	1		
104512	**Very good identification**						**104512**
					YEP		
Staphylococcus kloosii		97.88	0.54	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		2.12	0.07	MLT 95	12		
104530	**Acceptable identification**						**104530**
					YEP		
Staphylococcus kloosii		91.42	0.41	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		8.58	0.07	MLT 95	12		
104532	**Good identification**						**104532**
Staphylococcus kloosii		100.0	0.41	VPT 10			
104540	**Good identification**						**104540**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		92.96	0.56	PHS 9	1	12	
Macrococcus caseolyticus		3.96	0.14	GAL 10	99	99	
Staphylococcus kloosii		3.09	0.38		1	99	
104541	**Very good identification**						**104541**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		95.40	0.57	PHS 9	1	99	
Staphylococcus schleiferi subsp. schleiferi		3.35	0.19	GAL 10	1	1	
Kocuria kristinae		1.26	0.08		99	1	
104550	**Low discrimination**						**104550**
					OXI	YEP	
Staphylococcus kloosii		71.89	0.79	VPT 10	1	99	
Staphylococcus cohnii subsp. cohnii		27.84	0.58		1	12	
Macrococcus bovicus		0.27	0.10		99	99	
104551	**Very good identification**						**104551**
					NOV	YEP	
Staphylococcus cohnii subsp. cohnii		97.30	0.59	PHS 9	99	12	
Staphylococcus kloosii		2.34	0.38	GAL 10	99	99	
Staphylococcus caprae		0.36	-.07		1	1	
104552	**Very good identification**						**104552**
					NOV	YEP	
Staphylococcus kloosii		99.28	0.80	VPT 10	99	99	
Staphylococcus caprae		0.37	0.02		1	1	
Staphylococcus cohnii subsp. cohnii		0.36	0.17		99	12	
104562	**Good identification**						**104562**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.27	VPT 10				
104570	**Very good identification**							**104570**
					YEP			
Staphylococcus kloosii		98.46	0.67	VPT 10	99			
Staphylococcus cohnii subsp. cohnii		1.54	0.17		12			
104572	**Very good identification**							**104572**
Staphylococcus kloosii		100.0	0.68	VPT 10				
104573	**Good identification**							**104573**
Staphylococcus kloosii		100.0	0.27	VPT 10				
104601	**Low discrimination**							**104601**
					OXI	NOV		
Kocuria kristinae		86.64	0.39	PHS 1	99	1		
Staphylococcus schleiferi subsp. schleiferi		7.00	0.19	MLT 90	1	1		
Staphylococcus cohnii subsp. cohnii		5.73	0.25		1	99		
104603	**Good identification**							**104603**
Kocuria kristinae		100.0	0.27	PHS 1				
104610	**Low discrimination**							**104610**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		75.50	0.27	PHS 9	1	12		
Staphylococcus kloosii		18.21	0.26	SUC 1	1	99		
Macrococcus bovicus		6.29	-.02		99	99		
104612	**Good identification**							**104612**
Staphylococcus kloosii		100.0	0.27	VPT 10				
104640	**Very good identification**							**104640**
					PYR	COA	NOV	
Staphylococcus petrasii subsp. pragensis		98.04	0.59	PHS 1	99	1	1	
Staphylococcus cohnii subsp. cohnii		0.65	0.35		1	1	99	
Staphylococcus aureus subsp. anaerobius		0.57	0.16		1	99	1	
104641	**Good identification**							**104641**
					OXI	NOV	PYR	
Kocuria kristinae		94.72	0.59	PHS 1	99	1	93	
Staphylococcus petrasii subsp. pragensis		3.08	0.18		1	1	99	
Staphylococcus cohnii subsp. cohnii		2.20	0.35		1	99	1	
104643	**Good identification**							**104643**
Kocuria kristinae		100.0	0.47	PHS 1				
104650	**Low discrimination**							**104650**
					NOV	YEP	PYR	
Staphylococcus kloosii		39.11	0.53	VPT 10	99	99	52	
Staphylococcus petrasii subsp. pragensis		32.20	0.18	SUC 9	1	1	99	
Staphylococcus cohnii subsp. cohnii		27.03	0.37		99	12	1	
104651	**Low discrimination**							**104651**
					OXI	NOV		
Kocuria kristinae		58.47	0.31	PHS 1	99	1		
Staphylococcus cohnii subsp. cohnii		41.53	0.38	MAN 4	1	99		
104652	**Good identification**							**104652**
Staphylococcus kloosii		92.54	0.53	VPT 10				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. saprophyticus		7.46	0.02	SUC 9			
104670	**Good identification**						**104670**
Staphylococcus kloosii		100.0	0.40	VPT 10			
104672	**Low discrimination**						**104672**
Staphylococcus kloosii		76.22	0.41	VPT 10			
Staphylococcus arlettae		23.78	0.19	SUC 9			
104701	**Low discrimination**						**104701**
					OXI	NOV	
Kocuria kristinae		77.87	0.30	PHS 1	99	1	
Staphylococcus schleiferi subsp. schleiferi		18.86	0.19	MLT 90	1	1	
Staphylococcus cohnii subsp. cohnii		1.72	0.05		1	99	
104710	**Low discrimination**						**104710**
					YEP		
Staphylococcus kloosii		80.82	0.32	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		19.18	0.07	SUC 9	12		
104712	**Good identification**						**104712**
Staphylococcus kloosii		100.0	0.33	VPT 10			
104741	**Good identification**						**104741**
					OXI	NOV	
Kocuria kristinae		98.73	0.49	PHS 1	99	1	
Staphylococcus cohnii subsp. cohnii		0.77	0.16		1	99	
Rothia mucilaginosa		0.51	-.02		1	1	
104743	**Good identification**						**104743**
					OXI	PYR	
Kocuria kristinae		98.59	0.37	PHS 1	99	93	
Staphylococcus epidermidis		1.42	0.05		1	1	
104750	**Very good identification**						**104750**
					OXI	YEP	
Staphylococcus kloosii		95.34	0.59	VPT 10	1	99	
Staphylococcus cohnii subsp. cohnii		3.77	0.17	SUC 9	1	12	
Macrococcus bovicus		0.89	-.02		99	99	
104752	**Very good identification**						**104752**
					NOV	YEP	
Staphylococcus kloosii		98.64	0.59	VPT 10	99	99	
Staphylococcus caprae		1.36	-.07	SUC 9	1	1	
104770	**Good identification**						**104770**
Staphylococcus kloosii		100.0	0.46	VPT 10			
104772	**Low discrimination**						**104772**
Staphylococcus kloosii		52.34	0.47	VPT 10			
Staphylococcus arlettae		47.66	0.35	SUC 9			
104773	**Good identification**						**104773**
Staphylococcus arlettae		100.0	0.37	VPT 1			
105050	**Low discrimination**						**105050**
					OXI		
Staphylococcus kloosii		81.35	0.32	VPT 10	1		
Macrococcus equipersicus		18.65	0.02	TRE 99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
105052	**Good identification**						**105052**
Staphylococcus kloosii		100.0	0.33	VPT 10			
105150	**Good identification**						**105150**
Staphylococcus kloosii		100.0	0.38	VPT 10			
105152	**Good identification**						**105152**
Staphylococcus kloosii		100.0	0.39	VPT 10			
105172	**Good identification**						**105172**
Staphylococcus kloosii		100.0	0.27	VPT 10			
105410	**Low discrimination**						**105410**
					OXI	YEP	
Staphylococcus kloosii		63.89	0.47	VPT 10	1	99	
Staphylococcus cohnii subsp. cohnii		26.20	0.27	MLT 95	1	12	
Micrococcus carouselicus		9.91	0.08		99	67	
105411	**Good identification**						**105411**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	ESL 1			
105412	**Good identification**						**105412**
Staphylococcus kloosii		100.0	0.48	VPT 10			
105430	**Good identification**						**105430**
Staphylococcus kloosii		100.0	0.35	VPT 10			
105432	**Good identification**						**105432**
Staphylococcus kloosii		100.0	0.35	VPT 10			
105440	**Low discrimination**						**105440**
					YEP		
Staphylococcus cohnii subsp. cohnii		84.17	0.35	ESL 1	12		
Staphylococcus kloosii		15.83	0.32	PHS 9	99		
105441	**Acceptable identification**						**105441**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		90.21	0.35	ESL 1	1	99	
Kocuria kristinae		9.80	0.05	PHS 9	99	1	
105442	**Good identification**						**105442**
Staphylococcus kloosii		100.0	0.33	VPT 10			
105450	**Good identification**						**105450**
					OXI	YEP	
Staphylococcus kloosii		93.24	0.73	VPT 10	1	99	
Staphylococcus cohnii subsp. cohnii		6.37	0.37		1	12	
Micrococcus equipersicus		0.22	0.02		99	99	
105451	**Low discrimination**						**105451**
					YEP		
Staphylococcus cohnii subsp. cohnii		88.00	0.38	ESL 1	12		
Staphylococcus kloosii		12.00	0.32	PHS 9	99		
105452	**Very good identification**						**105452**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.74	VPT 10			
105453	**Low discrimination**						**105453**
					NOV	YEP	
Staphylococcus kloosii		83.48	0.33	VPT 10	99	99	
Aerococcus viridans		16.52	0.18	MNS 1	1	1	
105470	**Very good identification**						**105470**
Staphylococcus kloosii		100.0	0.61	VPT 10			
105472	**Very good identification**						**105472**
Staphylococcus kloosii		100.0	0.62	VPT 10			
105510	**Good identification**						**105510**
					OXI	YEP	
Staphylococcus kloosii		94.54	0.53	VPT 10	1	99	
Macrococcus carouselicus		3.24	0.00	MLT 95	99	67	
Staphylococcus cohnii subsp. cohnii		2.22	0.07		1	12	
105512	**Very good identification**						**105512**
Staphylococcus kloosii		100.0	0.54	VPT 10			
105530	**Good identification**						**105530**
Staphylococcus kloosii		100.0	0.41	VPT 10			
105532	**Good identification**						**105532**
Staphylococcus kloosii		100.0	0.41	VPT 10			
105540	**Low discrimination**						**105540**
					YEP		
Staphylococcus kloosii		76.67	0.38	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		23.33	0.15	MAN 99	12		
105542	**Good identification**						**105542**
Staphylococcus kloosii		100.0	0.39	VPT 10			
105550	**Very good identification**						**105550**
					YEP		
Staphylococcus kloosii		99.61	0.79	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		0.39	0.17		12		
105551	**Low discrimination**						**105551**
					YEP		
Staphylococcus kloosii		70.44	0.38	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		29.56	0.18	MNS 1	12		
105552	**Very good identification**						**105552**
Staphylococcus kloosii		100.0	0.80	VPT 10			
105553	**Acceptable identification**						**105553**
					NOV	YEP	
Staphylococcus kloosii		90.75	0.39	VPT 10	99	99	
Aerococcus viridans		9.25	0.18	MNS 1	1	1	
105562	**Good identification**						**105562**
Staphylococcus kloosii		100.0	0.27	VPT 10			
105570	**Very good identification**						**105570**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.67	VPT 10			
105572	**Very good identification**						**105572**
Staphylococcus kloosii		100.0	0.68	VPT 10			
105573	**Low discrimination**						**105573**
					NOV	YEP	
Staphylococcus kloosii		71.03	0.27	VPT 10	99	99	
Aerococcus viridans		28.97	0.18	MNS 1	1	1	
105601	**Good identification**						**105601**
Kocuria kristinae		100.0	0.27	PHS 1			
105612	**Good identification**						**105612**
Staphylococcus kloosii		100.0	0.27	VPT 10			
105641	**Good identification**						**105641**
Kocuria kristinae		100.0	0.47	PHS 1			
105643	**Good identification**						**105643**
Kocuria kristinae		100.0	0.34	PHS 1			
105650	**Good identification**						**105650**
					OXI		
Staphylococcus kloosii		98.35	0.53	VPT 10	1		
Staphylococcus sciuri		1.65	-.01	SUC 9	99		
105652	**Very good identification**						**105652**
Staphylococcus kloosii		100.0	0.53	VPT 10			
105670	**Good identification**						**105670**
Staphylococcus kloosii		100.0	0.40	VPT 10			
105672	**Low discrimination**						**105672**
Staphylococcus kloosii		76.22	0.41	VPT 10			
Staphylococcus arlettae		23.78	0.19	SUC 9			
105710	**Low discrimination**						**105710**
					OXI		
Staphylococcus kloosii		58.73	0.32	VPT 10	1		
Macrococcus carouselicus		41.27	0.06	SUC 9	99		
105712	**Good identification**						**105712**
Staphylococcus kloosii		100.0	0.33	VPT 10			
105741	**Low discrimination**						**105741**
					OXI		
Kocuria kristinae		84.32	0.37	PHS 1	99		
Rothia mucilaginosa		15.68	0.18		1		
105750	**Very good identification**						**105750**
					OXI		
Staphylococcus kloosii		98.35	0.59	VPT 10	1		
Staphylococcus sciuri		1.65	0.05	SUC 9	99		
105752	**Very good identification**						**105752**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.59	VPT 10			
105770	**Good identification**						**105770**
					OXI		
Staphylococcus kloosii		95.18	0.46	VPT 10	1		
Staphylococcus sciuri		4.82	0.03	SUC 9	99		
105772	**Low discrimination**						**105772**
Staphylococcus kloosii		52.34	0.47	VPT 10			
Staphylococcus arlettae		47.66	0.35	SUC 9			
105773	**Low discrimination**						**105773**
					OXI	YEP	
Staphylococcus arlettae		71.42	0.37	VPT 1	1	99	
Staphylococcus lentus		22.47	0.23	GLR 92	99	1	
Staphylococcus sciuri		3.64	0.05		99	57	
106000	**Low discrimination**						**106000**
					OXI		
Kocuria rosea		58.90	0.47	PHS 1	35		
Dermacoccus nishinomiyaensis		31.58	0.42		93		
Staphylococcus schleiferi subsp. schleiferi		5.60	0.29		1		
106001	**Good identification**						**106001**
Staphylococcus schleiferi subsp. schleiferi		93.10	0.70	ARG 86			
Staphylococcus capitis subsp. capitis		6.11	0.45				
Kocuria rosea		0.52	0.20				
106003	**Low discrimination**						**106003**
Staphylococcus schleiferi subsp. schleiferi		52.19	0.29	ARG 86			
Staphylococcus carnosus subsp. carnosus		40.90	0.27	LAC 1			
Staphylococcus capitis subsp. capitis		6.91	0.10				
106010	**Low discrimination**						**106010**
					OXI		
Kocuria rosea		55.74	0.27	PHS 1	35		
Staphylococcus capitis subsp. capitis		41.54	0.27	MAN 10	1		
Dermacoccus nishinomiyaensis		2.72	0.01		93		
106011	**Very good identification**						**106011**
Staphylococcus capitis subsp. capitis		97.88	0.68	PHS 11			
Staphylococcus schleiferi subsp. schleiferi		1.14	0.29				
Staphylococcus carnosus subsp. carnosus		0.89	0.27				
106013	**Low discrimination**						**106013**
Staphylococcus carnosus subsp. carnosus		64.04	0.39	ARG 99			
Staphylococcus capitis subsp. capitis		35.96	0.33	BGA 90			
106020	**Good identification**						**106020**
					OXI		
Kocuria rosea		95.67	0.39	PHS 1	35		
Kocuria varians		3.12	0.17		13		
Dermacoccus nishinomiyaensis		1.21	0.01		93		
106021	**Low discrimination**						**106021**
Staphylococcus schleiferi subsp. schleiferi		76.75	0.29	ARG 86			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		18.21	0.13	XYL 1			
Staphylococcus capitis subsp. capitis		5.03	0.04				
106031	**Good identification**						**106031**
Staphylococcus capitis subsp. capitis		97.06	0.27	PHS 11			
Kocuria rosea		2.94	-.07	XYL 1			
106041	**Low discrimination**						**106041**
Staphylococcus schleiferi subsp. schleiferi		82.60	0.29	ARG 86			
Staphylococcus epidermidis		6.62	0.10	MLT 1			
Staphylococcus capitis subsp. capitis		5.42	0.04				
106051	**Acceptable identification**						**106051**
					NOV		
Staphylococcus capitis subsp. capitis		91.32	0.27	PHS 11	1		
Staphylococcus cohnii subsp. cohnii		8.68	0.14	MLT 1	99		
106100	**Low discrimination**						**106100**
					OXI		
Dermacoccus nishinomiyaensis		38.25	0.27	VPT 5	93		
Staphylococcus schleiferi subsp. schleiferi		35.60	0.29	PHS 5	1		
Kocuria varians		22.37	0.30		13		
106101	**Very good identification**						**106101**
Staphylococcus schleiferi subsp. schleiferi		99.83	0.70	ARG 86			
Staphylococcus capitis subsp. capitis		0.07	0.04				
Kocuria varians		0.06	0.09				
106103	**Low discrimination**						**106103**
					COA		
Staphylococcus schleiferi subsp. schleiferi		77.33	0.29	ARG 86	1		
Staphylococcus carnosus subsp. carnosus		15.15	0.15	LAC 1	1		
Staphylococcus schleiferi subsp. coagulans		7.52	0.12		99		
106111	**Low discrimination**						**106111**
Staphylococcus schleiferi subsp. schleiferi		48.38	0.29	ARG 86			
Staphylococcus capitis subsp. capitis		42.14	0.27	MAN 1			
Staphylococcus carnosus subsp. carnosus		9.48	0.15				
106113	**Low discrimination**						**106113**
					COA		
Staphylococcus carnosus subsp. carnosus		89.72	0.27	ARG 99	1		
Staphylococcus schleiferi subsp. coagulans		10.28	0.12	BGA 90	99		
106121	**Good identification**						**106121**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.29	ARG 86			
106141	**Low discrimination**						**106141**
Staphylococcus schleiferi subsp. schleiferi		68.78	0.29	ARG 86			
Staphylococcus epidermidis		31.22	0.25	MLT 1			
106142	**Low discrimination**						**106142**
					NOV	OXI	
Staphylococcus epidermidis		56.37	0.30	URE 92	1	1	
Macrococcus caseolyticus		40.80	0.18	SUC 99	99	99	
Kocuria varians		2.83	0.05		1	13	
106143	**Good identification**						**106143**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		97.00	0.36	URE 92			
Staphylococcus caprae		3.00	-.07	SUC 99			
106200	**Low discrimination**						**106200**
					OXI		
Kocuria rosea		65.10	0.27	PHS 1	35		
Dermacoccus nishinomiyaensis		20.05	0.18	SUC 10	93		
Staphylococcus capitis subsp. capitis		6.78	0.09		1		
106201	**Acceptable identification**						**106201**
Staphylococcus capitis subsp. capitis		91.35	0.50	PHS 11			
Staphylococcus schleiferi subsp. schleiferi		7.58	0.29	MAN 93			
Staphylococcus epidermidis		0.61	0.10				
106210	**Low discrimination**						**106210**
					NOV	OXI	
Staphylococcus capitis subsp. capitis		89.13	0.32	PHS 11	1	1	
Kocuria rosea		7.16	0.07	MNS 99	1	35	
Staphylococcus vitulinus		3.71	0.00		99	99	
106211	**Very good identification**						**106211**
Staphylococcus capitis subsp. capitis		100.0	0.73	PHS 11			
106213	**Good identification**						**106213**
Staphylococcus capitis subsp. capitis		99.04	0.39	PHS 11			
Staphylococcus carnosus subsp. carnosus		0.96	-.02	LAC 2			
106231	**Good identification**						**106231**
Staphylococcus capitis subsp. capitis		100.0	0.32	PHS 11			
106240	**Low discrimination**						**106240**
					COA	NOV	
Staphylococcus epidermidis		76.92	0.45	URE 92	1	1	
Staphylococcus aureus subsp. anaerobius		12.49	0.16		99	1	
Staphylococcus hominis subsp. novobiosepticus		2.83	0.08		1	99	
106241	**Good identification**						**106241**
					OXI	PYR	
Staphylococcus epidermidis		96.26	0.51	URE 92	1	1	
Kocuria kristinae		1.79	0.10		99	93	
Staphylococcus capitis subsp. capitis		1.48	0.09		1	1	
106242	**Very good identification**						**106242**
					NOV		
Staphylococcus epidermidis		98.15	0.56	URE 92	1		
Staphylococcus hominis subsp. hominis		0.87	0.17		1		
Staphylococcus hominis subsp. novobiosepticus		0.61	0.03		99		
106243	**Very good identification**						**106243**
					OXI	PYR	
Staphylococcus epidermidis		99.85	0.61	URE 92	1	1	
Kocuria kristinae		0.15	-.03		99	93	
106251	**Good identification**						**106251**
Staphylococcus capitis subsp. capitis		95.28	0.32	PHS 11			
Staphylococcus epidermidis		4.72	0.10	MLT 1			
106301	**Low discrimination**						**106301**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus schleiferi subsp. schleiferi		62.38	0.29	ARG 86				
Staphylococcus epidermidis		28.31	0.25	SUC 1				
Staphylococcus capitis subsp. capitis		7.60	0.09					
106302	**Good identification**							**106302**
Staphylococcus epidermidis		100.0	0.30	URE 92				
106303	**Good identification**							**106303**
					COA			
Staphylococcus epidermidis		95.97	0.36	URE 92	1			
Staphylococcus schleiferi subsp. coagulans		2.05	0.02	MLT 99	99			
Staphylococcus hyicus		1.98	-.07		24			
106311	**Good identification**							**106311**
Staphylococcus capitis subsp. capitis		100.0	0.32	PHS 11				
106340	**Very good identification**							**106340**
					OXI	NOV	YEP	
Staphylococcus epidermidis		96.29	0.61	URE 92	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		1.87	0.18		1	99	1	
Macrococcus caseolyticus		0.99	0.11		99	99	99	
106341	**Very good identification**							**106341**
					NOV			
Staphylococcus epidermidis		99.49	0.66	URE 92	1			
Staphylococcus hominis subsp. novobiosepticus		0.26	0.05		99			
Rothia mucilaginosa		0.14	-.02		1			
106342	**Very good identification**							**106342**
					OXI	NOV	YEP	
Staphylococcus epidermidis		98.95	0.71	URE 92	1	1	1	
Macrococcus caseolyticus		0.48	0.14		99	99	99	
Staphylococcus hominis subsp. novobiosepticus		0.33	0.12		1	99	1	
106343	**Very good identification**							**106343**
					NOV			
Staphylococcus epidermidis		99.96	0.77	URE 92	1			
Staphylococcus hominis subsp. novobiosepticus		0.05	0.00		99			
106352	**Acceptable identification**							**106352**
					PYR			
Staphylococcus epidermidis		90.99	0.30	URE 92	1			
Staphylococcus caprae		5.01	-.07	MAN 1	75			
Staphylococcus haemolyticus		4.00	0.04		98			
106353	**Low discrimination**							**106353**
Staphylococcus epidermidis		82.14	0.36	URE 92				
Staphylococcus caprae		17.86	0.11	MAN 1				
106362	**Good identification**							**106362**
Staphylococcus epidermidis		100.0	0.30	URE 92				
106363	**Good identification**							**106363**
Staphylococcus epidermidis		100.0	0.36	URE 92				
106400	**Low discrimination**							**106400**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		53.31	0.39	ARG 86	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		33.35	0.42	MNS 99	99			
Staphylococcus auricularis		9.88	0.28		1			
106401	**Very good identification**							**106401**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		99.23	0.80	ARG 86	1			
Staphylococcus cohnii subsp. cohnii		0.68	0.43		99			
Staphylococcus capitis subsp. capitis		0.04	0.10		1			
106403	**Low discrimination**							**106403**
Staphylococcus schleiferi subsp. schleiferi		89.93	0.39	ARG 86				
Staphylococcus carnosus subsp. carnosus		10.07	0.19	LAC 1				
106410	**Good identification**							**106410**
					NOV			
Staphylococcus cohnii subsp. cohnii		98.75	0.45	NIT 7	99			
Staphylococcus schleiferi subsp. schleiferi		1.25	-.02	PHS 9	1			
106411	**Low discrimination**							**106411**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		39.05	0.39	ARG 86	1			
Staphylococcus cohnii subsp. cohnii		33.68	0.45	MAN 1	99			
Staphylococcus capitis subsp. capitis		22.91	0.33		1			
106413	**Acceptable identification**							**106413**
Staphylococcus carnosus subsp. carnosus		93.57	0.32	ARG 99				
Staphylococcus capitis subsp. capitis		2.50	-.02	BGA 90				
Staphylococcus schleiferi subsp. schleiferi		2.11	-.02					
106421	**Good identification**							**106421**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ARG 86				
106440	**Low discrimination**							**106440**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		76.40	0.53	NIT 7	1	99	12	
Staphylococcus auricularis		16.69	0.36	PHS 9	1	1	1	
Macrococcus caseolyticus		4.80	0.14		99	99	99	
106441	**Low discrimination**							**106441**
					NOV			
Staphylococcus cohnii subsp. cohnii		66.99	0.53	NIT 7	99			
Staphylococcus schleiferi subsp. schleiferi		31.22	0.39	PHS 9	1			
Staphylococcus auricularis		1.50	0.16		1			
106450	**Very good identification**							**106450**
					PYR	YEP		
Staphylococcus cohnii subsp. cohnii		97.03	0.55	NIT 7	1	12		
Staphylococcus kloosii		1.92	0.32	PHS 9	52	99		
Staphylococcus saprophyticus subsp. bovis		0.69	0.06		99	80		
106451	**Very good identification**							**106451**
					NOV			
Staphylococcus cohnii subsp. cohnii		99.63	0.56	NIT 7	99			
Staphylococcus schleiferi subsp. schleiferi		0.37	-.02	PHS 9	1			
106452	**Low discrimination**							**106452**
					YEP			
Staphylococcus kloosii		68.02	0.33	VPT 10	99			
Staphylococcus cohnii subsp. cohnii		31.98	0.14	NIT 1	12			
106500	**Low discrimination**							**106500**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	YEP	
Staphylococcus schleiferi subsp. schleiferi		81.28	0.39	ARG 86	1	1	1	
Micrococcus caseolyticus		10.12	0.14	MNS 99	99	99	99	
Staphylococcus cohnii subsp. cohnii		5.65	0.23		1	99	12	
106501	**Very good identification**							**106501**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		99.92	0.80	ARG 86	1			
Staphylococcus cohnii subsp. cohnii		0.08	0.23		99			
Rothia mucilaginosa		0.01	-.09		1			
106503	**Good identification**							**106503**
Staphylococcus schleiferi subsp. schleiferi		97.28	0.39	ARG 86				
Staphylococcus carnosus subsp. carnosus		2.72	0.07	LAC 1				
106511	**Low discrimination**							**106511**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		88.98	0.39	ARG 86	1			
Staphylococcus cohnii subsp. cohnii		8.53	0.26	MAN 1	99			
Staphylococcus carnosus subsp. carnosus		2.49	0.07		1			
106521	**Good identification**							**106521**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ARG 86				
106540	**Very good identification**							**106540**
					OXI	NOV	YEP	
Micrococcus caseolyticus		97.79	0.55	PHS 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		1.75	0.33		1	99	12	
Staphylococcus auricularis		0.38	0.16		1	1	1	
106541	**Low discrimination**							**106541**
					OXI	NOV	YEP	
Staphylococcus schleiferi subsp. schleiferi		72.24	0.39	ARG 86	1	1	1	
Staphylococcus cohnii subsp. cohnii		17.23	0.34	MLT 1	1	99	12	
Micrococcus caseolyticus		8.99	0.14		99	99	99	
106542	**Very good identification**							**106542**
					OXI	NOV		
Micrococcus caseolyticus		99.90	0.59	PHS 1	99	99		
Staphylococcus caprae		0.10	-.05		1	1		
106550	**Low discrimination**							**106550**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		47.38	0.35	NIT 7	1	12		
Micrococcus caseolyticus		21.05	0.14	PHS 9	99	99		
Staphylococcus kloosii		16.41	0.38	GAL 10	1	99		
106551	**Low discrimination**							**106551**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		65.28	0.36	NIT 7	99	1		
Staphylococcus caprae		28.70	0.13	PHS 9	1	75		
Staphylococcus saprophyticus subsp. bovis		3.84	0.06	GAL 10	99	99		
106553	**Good identification**							**106553**
Staphylococcus caprae		100.0	0.39	URE 90				
106572	**Good identification**							**106572**
Staphylococcus kloosii		100.0	0.27	VPT 10				
106601	**Low discrimination**							**106601**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	PYR		
Staphylococcus schleiferi subsp. schleiferi		82.39	0.39	ARG 86	1	37		
Kocuria kristinae		8.53	0.16	SUC 1	99	93		
Staphylococcus capitis subsp. capitis		6.76	0.15		1	1		
106611	**Good identification**							**106611**
					NOV			
Staphylococcus capitis subsp. capitis		97.53	0.39	PHS 11	1			
Staphylococcus schleiferi subsp. schleiferi		0.90	-.02	TRE 2	1			
Staphylococcus cohnii subsp. cohnii		0.78	0.04		99			
106620	**Very good identification**							**106620**
Staphylococcus muscae		100.0	0.59	VPT 1				
106640	**Low discrimination**							**106640**
Staphylococcus petrasii subsp. pragensis		76.71	0.39	NIT 10				
Staphylococcus haemolyticus		6.99	0.36	PHS 1				
Staphylococcus auricularis		6.91	0.35					
106641	**Low discrimination**							**106641**
					OXI			
Kocuria kristinae		83.71	0.36	NIT 7	99			
Staphylococcus petrasii subsp. pragensis		4.02	-.02	PHS 1	1			
Staphylococcus auricularis		3.98	0.15		1			
106642	**Low discrimination**							**106642**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		34.11	0.31	ARG 92	1	98	2	
Staphylococcus hominis subsp. hominis		18.86	0.27	PHS 1	1	1	1	
Macrococcus caseolyticus		18.67	0.14		99	80	99	
106650	**Low discrimination**							**106650**
					NOV			
Staphylococcus haemolyticus		60.08	0.31	ARG 92	2			
Staphylococcus saprophyticus subsp. bovis		18.09	0.18	PHS 1	99			
Staphylococcus petrasii subsp. pragensis		11.33	-.02		1			
106701	**Low discrimination**							**106701**
					OXI			
Staphylococcus schleiferi subsp. schleiferi		51.30	0.39	ARG 86	1			
Rothia mucilaginosa		46.42	0.32	SUC 1	1			
Kocuria kristinae		1.77	0.06		99			
106740	**Low discrimination**							**106740**
					OXI	NOV		
Macrococcus caseolyticus		85.03	0.52	PHS 1	99	99		
Rothia mucilaginosa		7.28	0.30		1	1		
Staphylococcus haemolyticus		4.72	0.37		1	2		
106741	**Low discrimination**							**106741**
					OXI	PYR		
Rothia mucilaginosa		78.77	0.39	ESL 90	1	75		
Kocuria kristinae		11.59	0.26	PHS 1	99	93		
Staphylococcus epidermidis		5.64	0.25		1	1		
106742	**Good identification**							**106742**
					OXI	PYR	NOV	
Macrococcus caseolyticus		94.99	0.55	PHS 1	99	80	99	
Staphylococcus epidermidis		1.99	0.30		1	1	1	
Staphylococcus haemolyticus		1.98	0.32		1	98	2	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
106743	**Low discrimination**							**106743**
					OXI	NOV	PYR	
Staphylococcus epidermidis		57.35	0.36	URE 92	1	1	1	
Macrococcus caseolyticus		14.90	0.14	TRE 1	99	99	80	
Kocuria kristinae		9.30	0.14		99	1	93	
106750	**Low discrimination**							**106750**
					OXI	NOV		
Staphylococcus saprophyticus subsp. bovis		43.09	0.31	URE 99	1	99		
Staphylococcus haemolyticus		40.35	0.32		1	2		
Macrococcus caseolyticus		12.51	0.11		99	99		
106752	**Low discrimination**							**106752**
					OXI	NOV		
Staphylococcus haemolyticus		29.81	0.27	ARG 92	1	2		
Staphylococcus caprae		25.31	0.13	PHS 1	1	1		
Macrococcus caseolyticus		24.64	0.14		99	99		
106753	**Low discrimination**							**106753**
					OXI	COA	NOV	
Staphylococcus caprae		63.57	0.30	URE 90	1	1	1	
Staphylococcus aureus subsp. aureus		27.25	0.18	ARG 99	1	99	4	
Staphylococcus lentus		6.63	0.23		99	1	99	
107001	**Acceptable identification**							**107001**
Staphylococcus schleiferi subsp. schleiferi		93.85	0.29	ARG 86				
Staphylococcus capitis subsp. capitis		6.15	0.04	ESL 1				
107011	**Acceptable identification**							**107011**
Staphylococcus capitis subsp. capitis		90.91	0.27	ESL 1				
Staphylococcus carnosus subsp. carnosus		9.09	0.07	PHS 11				
107101	**Good identification**							**107101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.29	ARG 86				
107211	**Good identification**							**107211**
					OXI	NOV		
Staphylococcus capitis subsp. capitis		95.27	0.32	ESL 1	1	1		
Staphylococcus sciuri		4.73	0.10	PHS 11	99	99		
107342	**Good identification**							**107342**
Staphylococcus epidermidis		100.0	0.30	URE 92				
107343	**Good identification**							**107343**
Staphylococcus epidermidis		100.0	0.36	URE 92				
107351	**Low discrimination**							**107351**
Staphylococcus sciuri		87.64	0.28	VPT 1				
Staphylococcus lentus		12.36	0.12	TRE 97				
107401	**Good identification**							**107401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ARG 86				
107450	**Low discrimination**							**107450**
					YEP			
Staphylococcus kloosii		66.26	0.32	VPT 10	99			
Staphylococcus cohnii subsp. cohnii		33.75	0.14	NIT 1	12			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
107452	**Good identification**						**107452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
107501	**Acceptable identification**						**107501**
Staphylococcus schleiferi subsp. schleiferi		92.40	0.39	ARG 86			
Rothia mucilaginosa		7.60	0.10	ESL 1			
107550	**Good identification**						**107550**
Staphylococcus kloosii		100.0	0.38	VPT 10			
107552	**Good identification**						**107552**
Staphylococcus kloosii		100.0	0.39	VPT 10			
107572	**Good identification**						**107572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
107611	**Good identification**						**107611**
					OXI	NOV	
Staphylococcus sciuri		97.24	0.41	VPT 1	99	99	
Staphylococcus lentus		1.53	0.06		99	99	
Staphylococcus capitis subsp. capitis		1.24	-.03		1	1	
107631	**Good identification**						**107631**
Staphylococcus sciuri		97.96	0.39	VPT 1			
Staphylococcus lentus		2.04	0.06				
107650	**Good identification**						**107650**
Staphylococcus sciuri		100.0	0.34	VPT 1			
107651	**Very good identification**						**107651**
					NOV	PYR	
Staphylococcus sciuri		98.87	0.53	VPT 1	99	1	
Staphylococcus lentus		0.84	0.12		99	6	
Kocuria kristinae		0.30	-.05		1	93	
107653	**Low discrimination**						**107653**
Staphylococcus sciuri		84.43	0.37	VPT 1			
Staphylococcus lentus		15.57	0.23	LAC 14			
107670	**Good identification**						**107670**
Staphylococcus sciuri		100.0	0.32	VPT 1			
107671	**Good identification**						**107671**
Staphylococcus sciuri		98.89	0.51	VPT 1			
Staphylococcus lentus		1.11	0.12				
107673	**Low discrimination**						**107673**
Staphylococcus sciuri		80.36	0.34	VPT 1			
Staphylococcus lentus		19.64	0.23	LAC 14			
107700	**Good identification**						**107700**
Rothia mucilaginosa		100.0	0.42	PHS 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
107701	**Good identification**							**107701**
					OXI	NOV		
Rothia mucilaginosa		98.98	0.51	PHS 1	1	1		
Staphylococcus sciuri		0.63	0.16		99	99		
Staphylococcus lentus		0.17	0.06		99	99		
107710	**Low discrimination**							**107710**
					OXI	NOV		
Staphylococcus sciuri		83.52	0.28	VPT 1	99	99		
Rothia mucilaginosa		11.02	0.00	MNS 89	1	1		
Staphylococcus lentus		5.46	0.06		99	99		
107711	**Low discrimination**							**107711**
					OXI	NOV		
Staphylococcus sciuri		54.09	0.47	VPT 1	99	99		
Staphylococcus lentus		43.27	0.47		99	99		
Rothia mucilaginosa		2.65	0.10		1	1		
107713	**Good identification**							**107713**
Staphylococcus lentus		94.57	0.58	VPT 7				
Staphylococcus sciuri		5.43	0.31					
107731	**Low discrimination**							**107731**
Staphylococcus lentus		51.47	0.47	VPT 7				
Staphylococcus sciuri		48.54	0.45					
107733	**Very good identification**							**107733**
Staphylococcus lentus		95.85	0.58	VPT 7				
Staphylococcus sciuri		4.15	0.28					
107740	**Good identification**							**107740**
					OXI	NOV		
Rothia mucilaginosa		98.34	0.49	PHS 1	1	1		
Micrococcus caseolyticus		1.29	0.11		99	99		
Staphylococcus sciuri		0.38	0.09		99	99		
107741	**Very good identification**							**107741**
					OXI	NOV	PYR	
Rothia mucilaginosa		98.44	0.59	PHS 1	1	1	75	
Staphylococcus sciuri		1.01	0.28		99	99	1	
Kocuria kristinae		0.40	0.14		99	1	93	
107750	**Low discrimination**							**107750**
					OXI	NOV		
Staphylococcus sciuri		88.54	0.40	VPT 1	99	99		
Rothia mucilaginosa		7.25	0.08	MNS 89	1	1		
Staphylococcus lentus		3.12	0.12		99	99		
107751	**Low discrimination**							**107751**
					OXI	NOV		
Staphylococcus sciuri		68.40	0.59	VPT 1	99	99		
Staphylococcus lentus		29.53	0.53		99	99		
Rothia mucilaginosa		2.08	0.18		1	1		
107753	**Good identification**							**107753**
					OXI	NOV		
Staphylococcus lentus		90.31	0.65	VPT 7	99	99		
Staphylococcus sciuri		9.61	0.43		99	99		
Staphylococcus caprae		0.09	-.11		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
107770	**Good identification**						**107770**
Staphylococcus sciuri		95.53	0.38	VPT 1			
Staphylococcus lentus		4.47	0.12	MNS 89			
107771	**Low discrimination**						**107771**
					OXI		
Staphylococcus sciuri		61.87	0.56	VPT 1	99		
Staphylococcus lentus		35.41	0.53		99		
Staphylococcus gallinarum		2.73	0.23		1		
107773	**Low discrimination**						**107773**
					OXI		
Staphylococcus lentus		88.15	0.65	VPT 7	99		
Staphylococcus sciuri		7.07	0.40		99		
Staphylococcus gallinarum		4.69	0.31		1		
110000	**Low discrimination**						**110000**
Micrococcus sp.		49.83	0.28	VPT 3			
Dermacoccus nishinomiyaensis		30.12	0.33	ORN 1			
Kocuria rosea		18.30	0.27				
110011	**Good identification**						**110011**
					NOV		
Staphylococcus capitis subsp. capitis		96.09	0.27	ORN 1	1		
Staphylococcus cohnii subsp. cohnii		3.91	0.07	NIT 89	99		
110211	**Good identification**						**110211**
Staphylococcus capitis subsp. capitis		100.0	0.32	ORN 1			
110241	**Good identification**						**110241**
Kocuria kristinae		100.0	0.33	ORN 1			
110400	**Good identification**						**110400**
					NOV	OXI	
Staphylococcus cohnii subsp. cohnii		96.44	0.45	ORN 1	99	1	
Staphylococcus auricularis		2.34	0.08		1	1	
Micrococcus sp.		1.22	-.13		1	89	
110401	**Good identification**						**110401**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		98.51	0.46	ORN 1	1	99	
Kocuria kristinae		1.49	-.02		99	1	
110410	**Good identification**						**110410**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		96.81	0.47	ORN 1	1	12	
Macrococcus bovicus		3.19	0.10		99	99	
110411	**Good identification**						**110411**
Staphylococcus cohnii subsp. cohnii		100.0	0.48	ORN 1			
110440	**Low discrimination**						**110440**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		85.63	0.55	ORN 1	99	1	
Staphylococcus petrasii subsp. pragensis		12.84	0.18		1	99	
Staphylococcus auricularis		1.53	0.16		1	40	
110441	**Very good identification**						**110441**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. cohnii		95.88	0.56	ORN 1	1	99	
Kocuria kristinae		4.12	0.18		99	1	
110450	**Very good identification**						**110450**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		97.63	0.58	ORN 1	1	12	
Macrococcus bovicus		2.37	0.18		99	99	
110451	**Very good identification**						**110451**
Staphylococcus cohnii subsp. cohnii		100.0	0.58	ORN 1			
110510	**Low discrimination**						**110510**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		88.72	0.28	ORN 1	1	12	
Macrococcus bovicus		11.28	0.03	GAL 10	99	99	
110511	**Good identification**						**110511**
Staphylococcus cohnii subsp. cohnii		100.0	0.28	ORN 1			
110540	**Low discrimination**						**110540**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		70.59	0.36	ORN 1	1	12	
Macrococcus caseolyticus		29.41	0.14	GAL 10	99	99	
110541	**Low discrimination**						**110541**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		88.58	0.36	ORN 1	1	99	
Kocuria kristinae		11.42	0.08	GAL 10	99	1	
110550	**Acceptable identification**						**110550**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		91.43	0.38	ORN 1	1	12	
Macrococcus bovicus		8.57	0.10	GAL 10	99	99	
110551	**Good identification**						**110551**
Staphylococcus cohnii subsp. cohnii		100.0	0.39	ORN 1			
110601	**Good identification**						**110601**
					OXI	NOV	
Kocuria kristinae		99.33	0.39	ORN 1	99	1	
Staphylococcus cohnii subsp. cohnii		0.67	0.05	MLT 90	1	99	
110603	**Low discrimination**						**110603**
					OXI		
Kocuria kristinae		82.62	0.27	ORN 1	99		
Staphylococcus lugdunensis		17.38	0.14	MLT 90	1		
110640	**Very good identification**						**110640**
					OXI		
Staphylococcus petrasii subsp. pragensis		99.18	0.59	ORN 1	1		
Staphylococcus petrasii subsp. jettensis		0.33	0.10		1		
Kocuria kristinae		0.31	0.18		99		
110641	**Good identification**						**110641**
					OXI		
Kocuria kristinae		90.98	0.59	ORN 1	99		
Staphylococcus lugdunensis		5.85	0.36		1		
Staphylococcus petrasii subsp. pragensis		2.96	0.18		1		
110643	**Low discrimination**						**110643**
					OXI		
Staphylococcus lugdunensis		69.83	0.55	NIT 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria kristinae		30.17	0.47		99		
110651	**Acceptable identification**						**110651**
					OXI	NOV	
Kocuria kristinae		93.24	0.31	ORN 1	99	1	
Staphylococcus cohnii subsp. cohnii		6.76	0.17	MAN 4	1	99	
110701	**Good identification**						**110701**
					OXI		
Kocuria kristinae		98.05	0.30	ORN 1	99		
Rothia mucilaginosa		1.95	-.09	MLT 90	1		
110741	**Low discrimination**						**110741**
					OXI		
Kocuria kristinae		83.47	0.49	ORN 1	99		
Staphylococcus lugdunensis		16.10	0.36		1		
Rothia mucilaginosa		0.43	-.02		1		
110743	**Low discrimination**						**110743**
					OXI		
Staphylococcus lugdunensis		87.41	0.55	NIT 99	1		
Kocuria kristinae		12.59	0.37		99		
111601	**Good identification**						**111601**
Kocuria kristinae		100.0	0.27	ORN 1			
111641	**Good identification**						**111641**
Kocuria kristinae		100.0	0.47	ORN 1			
111643	**Acceptable identification**						**111643**
					OXI		
Kocuria kristinae		91.45	0.34	ORN 1	99		
Staphylococcus lugdunensis		8.55	0.14		1		
111741	**Low discrimination**						**111741**
					OXI		
Kocuria kristinae		84.32	0.37	ORN 1	99		
Rothia mucilaginosa		15.68	0.18		1		
112000	**Low discrimination**						**112000**
					OXI		
Kocuria rosea		85.97	0.47	ORN 1	35		
Dermacoccus nishinomiyaensis		8.85	0.28		93		
Kocuria varians		3.42	0.27		13		
112010	**Acceptable identification**						**112010**
Kocuria rosea		94.26	0.27	ORN 1			
Staphylococcus capitis subsp. capitis		5.74	0.04	MAN 10			
112011	**Good identification**						**112011**
Staphylococcus capitis subsp. capitis		99.14	0.45	ORN 1			
Kocuria rosea		0.87	0.01				
112020	**Good identification**						**112020**
Kocuria rosea		96.84	0.39	ORN 1			
Kocuria varians		3.16	0.17				
112100	**Low discrimination**						**112100**
					OXI		
Kocuria varians		66.79	0.30	VPT 14	13		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		21.92	0.13	ORN 1	93			
Kocuria rosea		11.30	0.05		35			
112200	**Low discrimination**							**112200**
					OXI			
Kocuria rosea		87.25	0.27	ORN 1	35			
Dermacoccus nishinomiyaensis		5.16	0.03	SUC 10	93			
Kocuria varians		4.26	0.09		13			
112201	**Acceptable identification**							**112201**
Staphylococcus capitis subsp. capitis		94.12	0.28	ORN 1				
Kocuria rosea		5.88	0.01	MAN 93				
112211	**Good identification**							**112211**
Staphylococcus capitis subsp. capitis		100.0	0.51	ORN 1				
112241	**Low discrimination**							**112241**
					OXI	PYR		
Staphylococcus lugdunensis		86.36	0.36	TRE 99	1	96		
Staphylococcus epidermidis		6.90	0.18	LAC 90	1	1		
Kocuria kristinae		5.32	0.10		99	93		
112243	**Very good identification**							**112243**
					OXI	PYR		
Staphylococcus lugdunensis		97.10	0.55	TRE 99	1	96		
Staphylococcus epidermidis		2.73	0.28		1	1		
Kocuria kristinae		0.17	-.03		99	93		
112340	**Low discrimination**							**112340**
					OXI	NOV	YEP	
Staphylococcus epidermidis		40.41	0.28	URE 92	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		32.67	0.18	ORN 12	1	99	1	
Macrococcus caseolyticus		17.28	0.11		99	99	99	
112341	**Low discrimination**							**112341**
					NOV	PYR		
Staphylococcus lugdunensis		64.55	0.36	TRE 99	1	96		
Staphylococcus epidermidis		29.22	0.33	LAC 90	1	1		
Staphylococcus hominis subsp. novobiosepticus		3.18	0.05		99	1		
112342	**Low discrimination**							**112342**
					OXI	NOV	YEP	
Staphylococcus epidermidis		66.14	0.38	URE 92	1	1	1	
Macrococcus caseolyticus		13.39	0.14	ORN 12	99	99	99	
Staphylococcus hominis subsp. novobiosepticus		9.09	0.12		1	99	1	
112343	**Low discrimination**							**112343**
					NOV	PYR		
Staphylococcus lugdunensis		86.04	0.55	TRE 99	1	96		
Staphylococcus epidermidis		13.71	0.43		1	1		
Staphylococcus hominis subsp. novobiosepticus		0.25	0.00		99	1		
112400	**Low discrimination**							**112400**
					NOV			
Staphylococcus auricularis		62.80	0.28	VPT 12	1			
Staphylococcus cohnii subsp. cohnii		21.64	0.22	ORN 1	99			
Kocuria rosea		12.00	0.05		1			
112440	**Low discrimination**							**112440**
					OXI	NOV	YEP	
Staphylococcus auricularis		53.80	0.36	VPT 12	1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. cohnii		25.16	0.32	ORN 1	1	99	12		
Macrococcus caseolyticus		15.48	0.14		99	99	99		
112441	**Low discrimination**								**112441**
					NOV	PYR			
Staphylococcus lugdunensis		74.21	0.36	SUC 99	1	96			
Staphylococcus cohnii subsp. cohnii		20.43	0.33	LAC 90	99	1			
Staphylococcus auricularis		4.48	0.16		1	40			
112443	**Very good identification**								**112443**
Staphylococcus lugdunensis		100.0	0.55	SUC 99					
112450	**Good identification**								**112450**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		96.54	0.34	ORN 1	1	12			
Macrococcus bovicus		3.46	-.02	NIT 7	99	99			
112451	**Good identification**								**112451**
Staphylococcus cohnii subsp. cohnii		100.0	0.35	ORN 1					
112540	**Very good identification**								**112540**
					OXI	NOV	YEP		
Macrococcus caseolyticus		99.43	0.55	ORN 1	99	99	99		
Staphylococcus auricularis		0.39	0.16		1	1	1		
Staphylococcus cohnii subsp. cohnii		0.18	0.13		1	99	12		
112541	**Low discrimination**								**112541**
					OXI	NOV	PYR	YEP	
Staphylococcus lugdunensis		82.39	0.36	SUC 99	1	1	96	50	
Macrococcus caseolyticus		12.88	0.14	LAC 90	99	99	80	99	
Staphylococcus cohnii subsp. cohnii		2.52	0.13		1	99	1	12	
112542	**Very good identification**								**112542**
					OXI	NOV			
Macrococcus caseolyticus		99.61	0.59	ORN 1	99	99			
Staphylococcus lugdunensis		0.39	0.14		1	1			
112543	**Very good identification**								**112543**
					OXI	NOV			
Staphylococcus lugdunensis		97.46	0.55	SUC 99	1	1			
Macrococcus caseolyticus		2.54	0.18		99	99			
112600	**Low discrimination**								**112600**
Staphylococcus auricularis		79.26	0.27	VPT 12					
Staphylococcus petrasii subsp. pragensis		20.74	-.02	ORN 1					
112601	**Low discrimination**								**112601**
					OXI				
Staphylococcus lugdunensis		85.80	0.36	MLT 99	1				
Kocuria kristinae		11.16	0.16	LAC 90	99				
Staphylococcus auricularis		2.05	0.08		1				
112603	**Very good identification**								**112603**
					OXI				
Staphylococcus lugdunensis		99.64	0.55	MLT 99	1				
Kocuria kristinae		0.36	0.04		99				
112640	**Low discrimination**								**112640**
Staphylococcus petrasii subsp. pragensis		68.54	0.39	ORN 1					
Staphylococcus lugdunensis		12.31	0.36	NIT 10					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		6.25	0.36				
112641	**Very good identification**						**112641**
					OXI		
Staphylococcus lugdunensis		98.68	0.77	LAC 90	1		
Kocuria kristinae		1.17	0.36		99		
Staphylococcus auricularis		0.06	0.15		1		
112642	**Good identification**						**112642**
					OXI	NOV	
Staphylococcus lugdunensis		93.03	0.55	MNS 99	1	1	
Staphylococcus haemolyticus		2.95	0.31		1	2	
Macrococcus caseolyticus		1.62	0.14		99	99	
112643	**Excellent identification**						**112643**
					OXI		
Staphylococcus lugdunensis		99.97	0.96		1		
Kocuria kristinae		0.03	0.23		99		
112650	**Low discrimination**						**112650**
Staphylococcus haemolyticus		84.13	0.31	ARG 92			
Staphylococcus petrasii subsp. pragensis		15.87	-.02	ORN 1			
112651	**Good identification**						**112651**
					OXI		
Staphylococcus lugdunensis		95.35	0.36	MAN 1	1		
Kocuria kristinae		4.65	0.07	LAC 90	99		
112653	**Very good identification**						**112653**
					OXI		
Staphylococcus lugdunensis		99.87	0.55	MAN 1	1		
Kocuria kristinae		0.14	-.05		99		
112661	**Good identification**						**112661**
					OXI		
Staphylococcus lugdunensis		98.83	0.36	XYL 1	1		
Kocuria kristinae		1.17	-.05	LAC 90	99		
112663	**Very good identification**						**112663**
Staphylococcus lugdunensis		100.0	0.55	XYL 1			
112701	**Low discrimination**						**112701**
					OXI		
Rothia mucilaginosa		52.14	0.32	ORN 1	1		
Staphylococcus lugdunensis		45.88	0.36	ESL 90	1		
Kocuria kristinae		1.99	0.06		99		
112703	**Very good identification**						**112703**
					OXI		
Staphylococcus lugdunensis		99.75	0.55	MLT 99	1		
Rothia mucilaginosa		0.13	-.09		1		
Kocuria kristinae		0.12	-.06		99		
112740	**Low discrimination**						**112740**
					OXI	NOV	
Macrococcus caseolyticus		79.77	0.52	ORN 1	99	99	
Staphylococcus lugdunensis		7.73	0.36		1	1	
Rothia mucilaginosa		6.83	0.30		1	1	
112741	**Very good identification**						**112741**
					OXI		
Staphylococcus lugdunensis		96.90	0.77	LAC 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Rothia mucilaginosa		2.60	0.39		1		
Kocuria kristinae		0.38	0.26		99		
112742	**Low discrimination**						**112742**
					OXI	NOV	
Micrococcus caseolyticus		62.27	0.55	ORN 1	99	99	
Staphylococcus lugdunensis		36.21	0.55		1	1	
Staphylococcus haemolyticus		1.30	0.32		1	2	
112743	**Excellent identification**						**112743**
					OXI	NOV	
Staphylococcus lugdunensis		99.97	0.96		1	1	
Micrococcus caseolyticus		0.02	0.14		99	99	
Kocuria kristinae		0.01	0.14		99	1	
112750	**Low discrimination**						**112750**
					OXI	NOV	
Staphylococcus haemolyticus		73.91	0.32	ARG 92	1	2	
Micrococcus caseolyticus		22.91	0.11	ORN 1	99	99	
Staphylococcus saprophyticus subsp. bovis		3.19	0.02		1	99	
112751	**Acceptable identification**						**112751**
					OXI		
Staphylococcus lugdunensis		94.93	0.36	MAN 1	1		
Rothia mucilaginosa		2.54	-.02	LAC 90	1		
Kocuria kristinae		1.54	-.02		99		
112752	**Low discrimination**						**112752**
					OXI	NOV	
Staphylococcus haemolyticus		43.35	0.27	ARG 92	1	2	
Micrococcus caseolyticus		35.82	0.14	ORN 1	99	99	
Staphylococcus lugdunensis		20.83	0.14		1	1	
112753	**Very good identification**						**112753**
Staphylococcus lugdunensis		100.0	0.55	MAN 1			
112761	**Good identification**						**112761**
Staphylococcus lugdunensis		97.39	0.36	XYL 1			
Rothia mucilaginosa		2.61	-.02	LAC 90			
112763	**Very good identification**						**112763**
Staphylococcus lugdunensis		100.0	0.55	XYL 1			
113641	**Low discrimination**						**113641**
					OXI		
Staphylococcus lugdunensis		65.20	0.36	ESL 1	1		
Kocuria kristinae		19.09	0.23	LAC 90	99		
Rothia mucilaginosa		15.72	0.18		1		
113643	**Very good identification**						**113643**
					OXI		
Staphylococcus lugdunensis		99.19	0.55	ESL 1	1		
Kocuria kristinae		0.81	0.11		99		
113700	**Good identification**						**113700**
Rothia mucilaginosa		100.0	0.42	ORN 1			
113701	**Good identification**						**113701**
					OXI		
Rothia mucilaginosa		99.89	0.51	ORN 1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Kocuria kristinae		0.11	-.06		99		
113713	**Good identification**						**113713**
Staphylococcus lentus		100.0	0.30	VPT 7			
113733	**Good identification**						**113733**
Staphylococcus lentus		100.0	0.30	VPT 7			
113740	**Good identification**						**113740**
					OXI	NOV	
Rothia mucilaginosa		98.71	0.49	ORN 1	1	1	
Micrococcus caseolyticus		1.29	0.11		99	99	
113741	**Very good identification**						**113741**
					OXI		
Rothia mucilaginosa		95.60	0.59	ORN 1	1		
Staphylococcus lugdunensis		4.01	0.36		1		
Kocuria kristinae		0.39	0.14		99		
113743	**Very good identification**						**113743**
					OXI		
Staphylococcus lugdunensis		97.14	0.55	ESL 1	1		
Rothia mucilaginosa		2.60	0.18		1		
Kocuria kristinae		0.26	0.01		99		
113753	**Low discrimination**						**113753**
					OXI	NOV	
Staphylococcus lentus		84.38	0.36	VPT 7	99	99	
Staphylococcus lugdunensis		15.62	0.14	ORN 1	1	1	
113773	**Good identification**						**113773**
Staphylococcus lentus		100.0	0.36	VPT 7			
114343	**Good identification**						**114343**
Staphylococcus epidermidis		100.0	0.28	URE 92			
114410	**Good identification**						**114410**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	ORN 1			
114411	**Good identification**						**114411**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	ORN 1			
114440	**Good identification**						**114440**
Staphylococcus cohnii subsp. cohnii		100.0	0.35	ORN 1			
114441	**Good identification**						**114441**
Staphylococcus cohnii subsp. cohnii		100.0	0.35	ORN 1			
114450	**Low discrimination**						**114450**
					YEP		
Staphylococcus cohnii subsp. cohnii		87.12	0.37	ORN 1	12		
Staphylococcus kloosii		12.88	0.32	PHS 9	99		
114451	**Good identification**						**114451**
Staphylococcus cohnii subsp. cohnii		100.0	0.38	ORN 1			
114452	**Good identification**						**114452**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.33	VPT 10		
114550	**Low discrimination**					**114550**
					YEP	
Staphylococcus kloosii		72.08	0.38	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		27.92	0.17	ORN 1	12	
114552	**Good identification**					**114552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
114572	**Good identification**					**114572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
115450	**Good identification**					**115450**
Staphylococcus kloosii		100.0	0.32	VPT 10		
115452	**Good identification**					**115452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
115550	**Good identification**					**115550**
Staphylococcus kloosii		100.0	0.38	VPT 10		
115552	**Good identification**					**115552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
115572	**Good identification**					**115572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
116001	**Acceptable identification**					**116001**
Staphylococcus schleiferi subsp. schleiferi		93.85	0.29	ARG 86		
Staphylococcus capitis subsp. capitis		6.15	0.04	ORN 1		
116011	**Good identification**					**116011**
Staphylococcus capitis subsp. capitis		100.0	0.27	ORN 1		
116101	**Good identification**					**116101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.29	ARG 86		
116211	**Good identification**					**116211**
Staphylococcus capitis subsp. capitis		100.0	0.32	ORN 1		
116240	**Good identification**					**116240**
Staphylococcus epidermidis		100.0	0.28	URE 92		
116241	**Good identification**					**116241**
Staphylococcus epidermidis		100.0	0.33	URE 92		
116242	**Good identification**					**116242**
Staphylococcus epidermidis		100.0	0.38	URE 92		
116243	**Acceptable identification**					**116243**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		94.04	0.43	URE 92	1	
Staphylococcus lugdunensis		5.96	0.14	ORN 12	96	
116340	**Good identification**					**116340**
Staphylococcus epidermidis		100.0	0.43	URE 92		
116341	**Good identification**					**116341**
Staphylococcus epidermidis		100.0	0.49	URE 92		
116342	**Very good identification**					**116342**
Staphylococcus epidermidis		100.0	0.53	URE 92		
116343	**Very good identification**					**116343**
					PYR	
Staphylococcus epidermidis		98.89	0.59	URE 92	1	
Staphylococcus lugdunensis		1.11	0.14	ORN 12	96	
116401	**Good identification**					**116401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ARG 86		
116501	**Good identification**					**116501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ARG 86		
116641	**Good identification**					**116641**
					OXI	
Staphylococcus lugdunensis		98.83	0.36	PHS 1	1	
Kocuria kristinae		1.17	-.05	LAC 90	99	
116643	**Very good identification**					**116643**
					PYR	
Staphylococcus lugdunensis		99.84	0.55	PHS 1	96	
Staphylococcus epidermidis		0.16	0.02		1	
116741	**Acceptable identification**					**116741**
					PYR	
Staphylococcus lugdunensis		94.99	0.36	PHS 1	96	
Rothia mucilaginosa		2.54	-.02	LAC 90	75	
Staphylococcus epidermidis		2.46	0.08		1	
116743	**Very good identification**					**116743**
					PYR	
Staphylococcus lugdunensis		99.10	0.55	PHS 1	96	
Staphylococcus epidermidis		0.90	0.18		1	
120000	**Low discrimination**					**120000**
					OXI	
Micrococcus sp.		34.53	0.28	VPT 3	89	
Kocuria varians		30.71	0.44	BGA 1	13	
Dermacoccus nishinomiyaensis		20.87	0.33		93	
120002	**Good identification**					**120002**
					OXI	
Kocuria varians		96.13	0.32	VPT 14	13	
Micrococcus sp.		3.87	-.13		89	
120011	**Good identification**					**120011**
					NOV	
Staphylococcus capitis subsp. capitis		96.09	0.27	BGA 1	1	
Staphylococcus cohnii subsp. cohnii		3.91	0.07	NIT 89	99	
120020	**Low discrimination**					**120020**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI		
Kocuria varians		65.11	0.34	VPT 14	13		
Kocuria rosea		32.79	0.19		35		
Micrococcus sp.		2.11	-.13		89		
120040	**Low discrimination**						**120040**
					OXI		
Kocuria varians		54.30	0.30	VPT 14	13		
Micrococcus sp.		20.93	0.05		89		
Staphylococcus auricularis		17.66	0.19		1		
120100	**Acceptable identification**						**120100**
					OXI		
Kocuria varians		91.42	0.47	VPT 14	13		
Dermacoccus nishinomiyaensis		7.89	0.18		93		
Micrococcus sp.		0.69	-.13		89		
120102	**Good identification**						**120102**
Kocuria varians		100.0	0.36	VPT 14			
120120	**Good identification**						**120120**
Kocuria varians		100.0	0.38	VPT 14			
120122	**Good identification**						**120122**
Kocuria varians		100.0	0.27	VPT 14			
120140	**Good identification**						**120140**
Kocuria varians		100.0	0.34	VPT 14			
120201	**Good identification**						**120201**
					OXI	PYR	
Kocuria kristinae		97.88	0.44	TRE 95	99	93	
Staphylococcus capitis subsp. capitis		1.66	0.09	MLT 90	1	1	
Kocuria varians		0.46	0.04		13	10	
120203	**Good identification**						**120203**
Kocuria kristinae		100.0	0.32	TRE 95			
120211	**Low discrimination**						**120211**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		84.41	0.32	BGA 1	1	1	
Kocuria kristinae		15.59	0.16	NIT 89	99	93	
120241	**Very good identification**						**120241**
Kocuria kristinae		100.0	0.64	TRE 95			
120243	**Good identification**						**120243**
					OXI	PYR	
Kocuria kristinae		99.79	0.52	TRE 95	99	93	
Staphylococcus epidermidis		0.21	0.02		1	1	
120251	**Good identification**						**120251**
Kocuria kristinae		100.0	0.35	TRE 95			
120300	**Good identification**						**120300**
Kocuria varians		100.0	0.30	VPT 14			
120301	**Good identification**						**120301**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	PYR		
Kocuria kristinae		97.94	0.34	TRE 95	99	93		
Kocuria varians		2.06	0.08	MLT 90	13	10		
120341	**Very good identification**							**120341**
					OXI	PYR		
Kocuria kristinae		99.72	0.54	TRE 95	99	93		
Staphylococcus epidermidis		0.28	0.08		1	1		
120343	**Good identification**							**120343**
					OXI	PYR		
Kocuria kristinae		96.52	0.42	TRE 95	99	93		
Staphylococcus epidermidis		3.48	0.18		1	1		
120400	**Low discrimination**							**120400**
					NOV			
Staphylococcus cohnii subsp. cohnii		67.23	0.45	BGA 1	99			
Staphylococcus auricularis		26.30	0.33		1			
Kocuria varians		5.63	0.21		1			
120401	**Low discrimination**							**120401**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		58.13	0.46	BGA 1	1	99		
Kocuria kristinae		29.00	0.30		99	1		
Staphylococcus schleiferi subsp. schleiferi		10.54	0.23		1	1		
120410	**Good identification**							**120410**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		96.81	0.47	BGA 1	1	12		
Macrococcus bovicus		3.19	0.10		99	99		
120411	**Good identification**							**120411**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		98.39	0.48	BGA 1	1	99		
Kocuria kristinae		1.61	0.01		99	1		
120440	**Low discrimination**							**120440**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		68.05	0.55	BGA 1	99	1		
Staphylococcus auricularis		19.61	0.41		1	40		
Staphylococcus petrasii subsp. pragensis		10.20	0.18		1	99		
120441	**Low discrimination**							**120441**
					OXI	NOV		
Kocuria kristinae		57.93	0.49	SUC 99	99	1		
Staphylococcus cohnii subsp. cohnii		40.86	0.56		1	99		
Staphylococcus auricularis		1.21	0.21		1	1		
120443	**Acceptable identification**							**120443**
					OXI	NOV	PYR	
Kocuria kristinae		93.55	0.37	SUC 99	99	1	93	
Aerococcus viridans		2.92	0.28		1	1	95	
Staphylococcus cohnii subsp. cohnii		2.67	0.15		1	99	1	
120450	**Low discrimination**							**120450**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		86.11	0.58	BGA 1	1	12		
Staphylococcus saprophyticus subsp. saprophyticus		10.54	0.23		1	16		
Macrococcus bovicus		2.09	0.18		99	99		
120451	**Very good identification**							**120451**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		95.30	0.58	BGA 1	1	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		4.42	0.21		99	1	93	
Aerococcus viridans		0.28	0.18		1	1	95	
120452	**Acceptable identification**							**120452**
					NOV	YEP	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		94.80	0.38	URE 99	99	16	7	
Staphylococcus kloosii		2.17	0.33	SUC 91	99	99	52	
Aerococcus viridans		1.65	0.30		1	1	95	
120453	**Low discrimination**							**120453**
					OXI	NOV	PYR	
Aerococcus viridans		44.78	0.38	VPT 1	1	1	95	
Kocuria kristinae		19.96	0.08	ESL 90	99	1	93	
Staphylococcus cohnii subsp. cohnii		17.37	0.17		1	99	1	
120463	**Low discrimination**							**120463**
					OXI			
Aerococcus viridans		75.52	0.28	VPT 1	1			
Kocuria kristinae		24.48	-.04	ESL 90	99			
120472	**Low discrimination**							**120472**
					NOV	YEP	PYR	
Aerococcus viridans		52.44	0.30	VPT 1	1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		30.38	-.03	ESL 90	99	16	7	
Staphylococcus kloosii		17.17	0.21		99	99	52	
120473	**Good identification**							**120473**
Aerococcus viridans		100.0	0.38	VPT 1				
120510	**Low discrimination**							**120510**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		88.72	0.28	BGA 1	1	12		
Macrococcus bovicus		11.28	0.03	GAL 10	99	99		
120511	**Good identification**							**120511**
Staphylococcus cohnii subsp. cohnii		100.0	0.28	BGA 1				
120540	**Low discrimination**							**120540**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		54.65	0.36	BGA 1	1	99	12	
Macrococcus caseolyticus		22.77	0.14	GAL 10	99	99	99	
Staphylococcus auricularis		15.75	0.21		1	1	1	
120541	**Low discrimination**							**120541**
					OXI	NOV		
Kocuria kristinae		80.51	0.39	SUC 99	99	1		
Staphylococcus cohnii subsp. cohnii		18.93	0.36		1	99		
Staphylococcus auricularis		0.56	0.02		1	1		
120543	**Acceptable identification**							**120543**
					OXI			
Kocuria kristinae		91.45	0.27	SUC 99	99			
Aerococcus viridans		8.55	0.28		1			
120550	**Low discrimination**							**120550**
					YEP			
Staphylococcus cohnii subsp. cohnii		69.21	0.38	BGA 1	12			
Staphylococcus kloosii		17.67	0.38	GAL 10	99			
Staphylococcus saprophyticus subsp. saprophyticus		6.63	0.01		16			
120551	**Low discrimination**							**120551**
					OXI	NOV	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		85.82	0.39	BGA 1	1	99	1	
Kocuria kristinae		11.95	0.11	GAL 10	99	1	93	
Aerococcus viridans		2.23	0.18		1	1	95	
120553	**Low discrimination**							**120553**
					OXI			
Aerococcus viridans		87.07	0.38	VPT 1	1			
Kocuria kristinae		12.93	-.02	ESL 90	99			
120563	**Good identification**							**120563**
Aerococcus viridans		100.0	0.28	VPT 1				
120572	**Low discrimination**							**120572**
					NOV	YEP		
Aerococcus viridans		61.14	0.30	VPT 1	1	1		
Staphylococcus kloosii		38.86	0.27	ESL 90	99	99		
120573	**Good identification**							**120573**
Aerococcus viridans		100.0	0.38	VPT 1				
120600	**Low discrimination**							**120600**
					OXI			
Kocuria kristinae		38.35	0.30	MLT 90	99			
Staphylococcus petrasii subsp. pragensis		33.70	0.18	MNS 99	1			
Staphylococcus auricularis		25.62	0.33		1			
120601	**Very good identification**							**120601**
					OXI	NOV		
Kocuria kristinae		99.91	0.71	MLT 90	99	1		
Staphylococcus auricularis		0.08	0.13		1	1		
Staphylococcus cohnii subsp. cohnii		0.02	0.05		1	99		
120603	**Very good identification**							**120603**
Kocuria kristinae		100.0	0.58	MLT 90				
120611	**Good identification**							**120611**
					OXI	NOV	PYR	
Kocuria kristinae		98.81	0.42	MAN 4	99	1	93	
Staphylococcus cohnii subsp. cohnii		0.62	0.07	MLT 90	1	99	1	
Staphylococcus capitis subsp. capitis		0.58	-.03		1	1	1	
120613	**Good identification**							**120613**
					OXI			
Kocuria kristinae		95.15	0.30	MAN 4	99			
Aerococcus viridans		4.85	0.25	MLT 90	1			
120621	**Good identification**							**120621**
Kocuria kristinae		100.0	0.30	XYL 1				
120640	**Low discrimination**							**120640**
					OXI			
Staphylococcus petrasii subsp. pragensis		87.96	0.59	BGA 1	1			
Kocuria kristinae		9.10	0.49		99			
Staphylococcus auricularis		1.58	0.40		1			
120641	**Very good identification**							**120641**
					OXI			
Kocuria kristinae		99.88	0.90		99			
Staphylococcus petrasii subsp. pragensis		0.10	0.18		1			
Staphylococcus auricularis		0.02	0.20		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
120642	**Low discrimination**							**120642**
					OXI	NOV	PYR	
Kocuria kristinae		38.68	0.37	MNS 99	99	1	93	
Staphylococcus saprophyticus subsp. saprophyticus		37.45	0.31		1	99	7	
Staphylococcus petrasii subsp. pragensis		15.11	0.18		1	1	99	
120643	**Very good identification**							**120643**
					OXI			
Kocuria kristinae		99.93	0.78		99			
Aerococcus viridans		0.03	0.28		1			
Staphylococcus lugdunensis		0.02	0.02		1			
120650	**Low discrimination**							**120650**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		83.75	0.43	URE 99	1	99	7	
Staphylococcus petrasii subsp. pragensis		7.98	0.18		1	1	99	
Kocuria kristinae		3.40	0.21		99	1	93	
120651	**Very good identification**							**120651**
					OXI	NOV		
Kocuria kristinae		99.47	0.62	MAN 4	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		0.25	0.02		1	99		
Staphylococcus cohnii subsp. cohnii		0.22	0.17		1	99		
120652	**Very good identification**							**120652**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		99.33	0.59	URE 99	1	99	7	
Staphylococcus warneri		0.18	0.23		1	1	8	
Kocuria kristinae		0.18	0.08		99	1	93	
120653	**Acceptable identification**							**120653**
					OXI	NOV	PYR	
Kocuria kristinae		92.63	0.49	MAN 4	99	1	93	
Staphylococcus saprophyticus subsp. saprophyticus		5.27	0.18		1	99	7	
Aerococcus viridans		2.10	0.38		1	1	95	
120661	**Good identification**							**120661**
Kocuria kristinae		100.0	0.49	XYL 1				
120663	**Good identification**							**120663**
					OXI			
Kocuria kristinae		96.98	0.37	XYL 1	99			
Aerococcus viridans		3.02	0.28		1			
120673	**Low discrimination**							**120673**
					OXI	NOV	YEP	
Aerococcus viridans		54.81	0.38	VPT 1	1	1	1	
Kocuria kristinae		24.43	0.08	ESL 90	99	1	15	
Staphylococcus arlettae		20.77	0.18		1	99	99	
120701	**Very good identification**							**120701**
					OXI			
Kocuria kristinae		98.53	0.61	MLT 90	99			
Rothia mucilaginosa		1.47	0.19		1			
120703	**Good identification**							**120703**
Kocuria kristinae		100.0	0.48	MLT 90				
120711	**Good identification**							**120711**
Kocuria kristinae		100.0	0.32	MAN 4				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
120740	**Low discrimination**							**120740**
					OXI			
Kocuria kristinae		59.72	0.39	MNS 99	99			
Staphylococcus petrasii subsp. pragensis		17.50	0.18		1			
Rothia mucilaginosa		7.64	0.17		1			
120741	**Very good identification**							**120741**
					OXI			
Kocuria kristinae		99.61	0.80		99			
Rothia mucilaginosa		0.39	0.27		1			
Staphylococcus auricularis		0.01	0.01		1			
120742	**Low discrimination**							**120742**
					OXI	NOV		
Kocuria kristinae		52.03	0.27	MNS 99	99	1		
Macrococcus caseolyticus		18.83	0.14		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		13.14	0.09		1	99		
120743	**Very good identification**							**120743**
					OXI			
Kocuria kristinae		99.85	0.68		99			
Aerococcus viridans		0.09	0.28		1			
Staphylococcus lugdunensis		0.05	0.02		1			
120751	**Good identification**							**120751**
					OXI			
Kocuria kristinae		99.81	0.52	MAN 4	99			
Aerococcus viridans		0.19	0.18		1			
120752	**Acceptable identification**							**120752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		94.66	0.37	URE 99	99	7		
Aerococcus viridans		1.88	0.30	GAL 8	1	95		
Staphylococcus haemolyticus		1.71	0.15		2	98		
120753	**Acceptable identification**							**120753**
					OXI	NOV	PYR	
Kocuria kristinae		92.35	0.40	MAN 4	99	1	93	
Aerococcus viridans		6.28	0.38		1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		1.37	-.04		1	99	7	
120761	**Good identification**							**120761**
Kocuria kristinae		100.0	0.39	XYL 1				
120763	**Acceptable identification**							**120763**
					OXI			
Kocuria kristinae		91.45	0.27	XYL 1	99			
Aerococcus viridans		8.55	0.28		1			
120772	**Low discrimination**							**120772**
					NOV	PYR		
Staphylococcus arlettae		71.46	0.31	VPT 1	99	1		
Aerococcus viridans		18.91	0.30	GLR 92	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		9.63	-.04		99	7		
120773	**Low discrimination**							**120773**
					OXI	NOV	YEP	
Staphylococcus arlettae		62.99	0.34	VPT 1	1	99	99	
Aerococcus viridans		29.34	0.38	GLR 92	1	1	1	
Kocuria kristinae		4.36	-.02		99	1	15	
121053	**Good identification**							**121053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Aerococcus viridans		100.0	0.31	VPT 1		
121073	**Good identification**					**121073**
Aerococcus viridans		100.0	0.31	VPT 1		
121153	**Good identification**					**121153**
Aerococcus viridans		100.0	0.31	VPT 1		
121173	**Good identification**					**121173**
Aerococcus viridans		100.0	0.31	VPT 1		
121201	**Good identification**					**121201**
Kocuria kristinae		100.0	0.32	TRE 95		
121241	**Good identification**					**121241**
Kocuria kristinae		100.0	0.52	TRE 95		
121243	**Good identification**					**121243**
					OXI	
Kocuria kristinae		98.88	0.39	TRE 95	99	
Aerococcus viridans		1.12	0.21		1	
121273	**Good identification**					**121273**
Aerococcus viridans		100.0	0.31	VPT 1		
121341	**Good identification**					**121341**
					OXI	
Kocuria kristinae		97.39	0.42	TRE 95	99	
Rothia mucilaginosa		2.61	0.05		1	
121343	**Good identification**					**121343**
					OXI	
Kocuria kristinae		96.71	0.29	TRE 95	99	
Aerococcus viridans		3.29	0.21		1	
121353	**Low discrimination**					**121353**
					OXI	
Aerococcus viridans		71.00	0.31	VPT 1	1	
Kocuria kristinae		29.00	0.01	TRE 95	99	
121373	**Good identification**					**121373**
Aerococcus viridans		100.0	0.31	VPT 1		
121402	**Good identification**					**121402**
Aerococcus viridans		100.0	0.27	VPT 1		
121403	**Low discrimination**					**121403**
					OXI	
Aerococcus viridans		71.63	0.35	VPT 1	1	
Kocuria kristinae		28.37	0.05		99	
121412	**Good identification**					**121412**
Aerococcus viridans		100.0	0.37	VPT 1		
121413	**Good identification**					**121413**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.45	VPT 1				
121422	**Good identification**							**121422**
Aerococcus viridans		100.0	0.27	VPT 1				
121423	**Good identification**							**121423**
Aerococcus viridans		100.0	0.35	VPT 1				
121432	**Good identification**							**121432**
Aerococcus viridans		100.0	0.37	VPT 1				
121433	**Good identification**							**121433**
Aerococcus viridans		100.0	0.45	VPT 1				
121441	**Acceptable identification**							**121441**
					OXI	NOV	PYR	
Kocuria kristinae		94.37	0.37	SUC 99	99	1	93	
Aerococcus viridans		2.94	0.28		1	1	95	
Staphylococcus cohnii subsp. cohnii		2.69	0.15		1	99	1	
121442	**Good identification**							**121442**
Aerococcus viridans		100.0	0.40	VPT 1				
121443	**Low discrimination**							**121443**
					OXI			
Aerococcus viridans		52.87	0.47	VPT 1	1			
Kocuria kristinae		47.13	0.24		99			
121450	**Low discrimination**							**121450**
					NOV	YEP	PYR	
Staphylococcus kloosii		34.11	0.32	VPT 10	99	99	52	
Aerococcus viridans		28.21	0.30	PHS 99	1	1	95	
Staphylococcus cohnii subsp. cohnii		23.57	0.16		99	12	1	
121451	**Low discrimination**							**121451**
					OXI	NOV	PYR	
Aerococcus viridans		54.54	0.38	VPT 1	1	1	95	
Kocuria kristinae		24.30	0.08	LAC 90	99	1	93	
Staphylococcus cohnii subsp. cohnii		21.16	0.17		1	99	1	
121452	**Low discrimination**							**121452**
					NOV	YEP	PYR	
Aerococcus viridans		82.65	0.50	VPT 1	1	1	95	
Staphylococcus kloosii		12.03	0.33		99	99	52	
Staphylococcus saprophyticus subsp. saprophyticus		5.32	-.03		99	16	7	
121453	**Very good identification**							**121453**
					OXI			
Aerococcus viridans		98.78	0.57	VPT 1	1			
Kocuria kristinae		1.22	-.04		99			
121461	**Low discrimination**							**121461**
					OXI			
Aerococcus viridans		75.52	0.28	VPT 1	1			
Kocuria kristinae		24.48	-.04	LAC 90	99			
121462	**Good identification**							**121462**
Aerococcus viridans		100.0	0.40	VPT 1				
121463	**Good identification**							**121463**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.47	VPT 1			
121470	**Low discrimination**						**121470**
					NOV	YEP	
Aerococcus viridans		76.79	0.30	VPT 1	1	1	
Staphylococcus kloosii		23.21	0.20	LAC 90	99	99	
121471	**Good identification**						**121471**
Aerococcus viridans		100.0	0.38	VPT 1			
121472	**Good identification**						**121472**
					NOV	YEP	
Aerococcus viridans		96.49	0.50	VPT 1	1	1	
Staphylococcus kloosii		3.51	0.21		99	99	
121473	**Very good identification**						**121473**
Aerococcus viridans		100.0	0.57	VPT 1			
121502	**Good identification**						**121502**
Aerococcus viridans		100.0	0.27	VPT 1			
121503	**Low discrimination**						**121503**
					OXI		
Aerococcus viridans		88.34	0.35	VPT 1	1		
Kocuria kristinae		11.66	-.05		99		
121512	**Good identification**						**121512**
Aerococcus viridans		100.0	0.37	VPT 1			
121513	**Good identification**						**121513**
Aerococcus viridans		100.0	0.45	VPT 1			
121522	**Good identification**						**121522**
Aerococcus viridans		100.0	0.27	VPT 1			
121523	**Good identification**						**121523**
Aerococcus viridans		100.0	0.35	VPT 1			
121532	**Good identification**						**121532**
Aerococcus viridans		100.0	0.37	VPT 1			
121533	**Good identification**						**121533**
Aerococcus viridans		100.0	0.45	VPT 1			
121541	**Low discrimination**						**121541**
					OXI		
Kocuria kristinae		81.11	0.27	SUC 99	99		
Rothia mucilaginosa		11.31	0.05		1		
Aerococcus viridans		7.58	0.28		1		
121542	**Good identification**						**121542**
Aerococcus viridans		100.0	0.40	VPT 1			
121543	**Low discrimination**						**121543**
					OXI		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		77.10	0.47	VPT 1	1		
Kocuria kristinae		22.91	0.15		99		
121550	**Low discrimination**						**121550**
					NOV	YEP	
Staphylococcus kloosii		70.12	0.38	VPT 10	99	99	
Aerococcus viridans		29.88	0.30	PHS 99	1	1	
121551	**Low discrimination**						**121551**
					OXI		
Aerococcus viridans		87.07	0.38	VPT 1	1		
Kocuria kristinae		12.93	-.02	LAC 90	99		
121552	**Low discrimination**						**121552**
					NOV	YEP	
Aerococcus viridans		77.97	0.50	VPT 1	1	1	
Staphylococcus kloosii		22.03	0.39		99	99	
121553	**Very good identification**						**121553**
Aerococcus viridans		100.0	0.57	VPT 1			
121561	**Good identification**						**121561**
Aerococcus viridans		100.0	0.28	VPT 1			
121562	**Good identification**						**121562**
Aerococcus viridans		100.0	0.40	VPT 1			
121563	**Good identification**						**121563**
Aerococcus viridans		100.0	0.47	VPT 1			
121570	**Low discrimination**						**121570**
					NOV	YEP	
Aerococcus viridans		63.02	0.30	VPT 1	1	1	
Staphylococcus kloosii		36.98	0.26	LAC 90	99	99	
121571	**Good identification**						**121571**
Aerococcus viridans		100.0	0.38	VPT 1			
121572	**Acceptable identification**						**121572**
					NOV	YEP	
Aerococcus viridans		93.40	0.50	VPT 1	1	1	
Staphylococcus kloosii		6.60	0.27		99	99	
121573	**Very good identification**						**121573**
Aerococcus viridans		100.0	0.57	VPT 1			
121601	**Very good identification**						**121601**
					OXI		
Kocuria kristinae		99.82	0.58	MLT 90	99		
Rothia mucilaginosa		0.18	-.02		1		
121602	**Low discrimination**						**121602**
					OXI		
Aerococcus viridans		51.97	0.27	VPT 1	1		
Kocuria kristinae		48.03	0.05		99		
121603	**Good identification**						**121603**
					OXI		
Kocuria kristinae		97.51	0.46	MLT 90	99		
Aerococcus viridans		2.49	0.35		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
121611	**Good identification**							**121611**
					OXI			
Kocuria kristinae		95.15	0.30	MAN 4	99			
Aerococcus viridans		4.85	0.25	MLT 90	1			
121612	**Good identification**							**121612**
Aerococcus viridans		100.0	0.37	VPT 1				
121613	**Low discrimination**							**121613**
					OXI			
Aerococcus viridans		64.74	0.45	VPT 1	1			
Kocuria kristinae		35.26	0.17		99			
121622	**Good identification**							**121622**
Aerococcus viridans		100.0	0.27	VPT 1				
121623	**Low discrimination**							**121623**
					OXI			
Aerococcus viridans		71.63	0.35	VPT 1	1			
Kocuria kristinae		28.37	0.05		99			
121632	**Good identification**							**121632**
Aerococcus viridans		100.0	0.37	VPT 1				
121633	**Good identification**							**121633**
Aerococcus viridans		100.0	0.45	VPT 1				
121640	**Low discrimination**							**121640**
					OXI			
Kocuria kristinae		70.45	0.37	MNS 99	99			
Staphylococcus petrasii subsp. pragensis		27.52	0.18		1			
Rothia mucilaginosa		1.09	-.04		1			
121641	**Very good identification**							**121641**
					OXI			
Kocuria kristinae		99.92	0.78		99			
Rothia mucilaginosa		0.05	0.05		1			
Aerococcus viridans		0.03	0.28		1			
121643	**Very good identification**							**121643**
					OXI			
Kocuria kristinae		98.88	0.65		99			
Aerococcus viridans		1.12	0.47		1			
121651	**Good identification**							**121651**
					OXI			
Kocuria kristinae		97.78	0.49	MAN 4	99			
Aerococcus viridans		2.22	0.38		1			
121652	**Low discrimination**							**121652**
					OXI	NOV	PYR	
Aerococcus viridans		59.53	0.50	VPT 1	1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		38.75	0.18		1	99	7	
Kocuria kristinae		1.72	-.04		99	1	93	
121653	**Low discrimination**							**121653**
					OXI			
Kocuria kristinae		55.07	0.37	MAN 4	99			
Aerococcus viridans		44.93	0.57		1			
121661	**Good identification**							**121661**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Kocuria kristinae		96.98	0.37	XYL 1	99			
Aerococcus viridans		3.02	0.28		1			
121662	**Good identification**							**121662**
Aerococcus viridans		100.0	0.40	VPT 1				
121663	**Low discrimination**							**121663**
					OXI			
Aerococcus viridans		52.87	0.47	VPT 1	1			
Kocuria kristinae		47.13	0.24		99			
121670	**Good identification**							**121670**
Aerococcus viridans		100.0	0.30	VPT 1				
121671	**Low discrimination**							**121671**
					OXI			
Aerococcus viridans		69.17	0.38	VPT 1	1			
Kocuria kristinae		30.83	0.08	LAC 90	99			
121672	**Acceptable identification**							**121672**
					NOV	YEP		
Aerococcus viridans		93.10	0.50	VPT 1	1	1		
Staphylococcus arlettae		6.90	0.15		99	99		
121673	**Good identification**							**121673**
					OXI	NOV	YEP	
Aerococcus viridans		94.83	0.57	VPT 1	1	1	1	
Staphylococcus arlettae		3.99	0.18		1	99	99	
Kocuria kristinae		1.17	-.04		99	1	15	
121700	**Low discrimination**							**121700**
					OXI	NOV	PYR	
Rothia mucilaginosa		87.92	0.29	NIT 99	1	1	75	
Macrococcus carouselicus		7.12	0.10		99	99	1	
Kocuria kristinae		4.95	0.07		99	1	93	
121701	**Low discrimination**							**121701**
					OXI			
Kocuria kristinae		65.02	0.48	MLT 90	99			
Rothia mucilaginosa		34.98	0.39		1			
121702	**Low discrimination**							**121702**
					OXI			
Aerococcus viridans		76.45	0.27	VPT 1	1			
Kocuria kristinae		23.55	-.05		99			
121703	**Acceptable identification**							**121703**
					OXI			
Kocuria kristinae		91.06	0.36	MLT 90	99			
Aerococcus viridans		6.97	0.35		1			
Rothia mucilaginosa		1.98	-.02		1			
121712	**Good identification**							**121712**
Aerococcus viridans		100.0	0.37	VPT 1				
121713	**Low discrimination**							**121713**
					OXI	NOV	PYR	
Aerococcus viridans		48.34	0.45	VPT 1	1	1	95	
Staphylococcus lentus		42.88	0.30		99	99	6	
Kocuria kristinae		8.78	0.08		99	1	93	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
121722	**Good identification**							**121722**
Aerococcus viridans		100.0	0.27	VPT 1				
121723	**Low discrimination**							**121723**
					OXI			
Aerococcus viridans		88.34	0.35	VPT 1	1			
Kocuria kristinae		11.66	-.05		99			
121732	**Good identification**							**121732**
Aerococcus viridans		100.0	0.37	VPT 1				
121733	**Low discrimination**							**121733**
					OXI	NOV		
Aerococcus viridans		52.99	0.45	VPT 1	1	1		
Staphylococcus lentus		47.01	0.30		99	99		
121740	**Low discrimination**							**121740**
					OXI			
Rothia mucilaginosa		81.57	0.37	NIT 99	1			
Kocuria kristinae		17.72	0.27		99			
Aerococcus viridans		0.71	0.20		1			
121741	**Low discrimination**							**121741**
					OXI			
Kocuria kristinae		87.69	0.68		99			
Rothia mucilaginosa		12.23	0.47		1			
Aerococcus viridans		0.08	0.28		1			
121742	**Low discrimination**							**121742**
					OXI			
Aerococcus viridans		54.88	0.40	VPT 1	1			
Kocuria kristinae		38.05	0.15		99			
Rothia mucilaginosa		7.08	-.04		1			
121743	**Very good identification**							**121743**
					OXI			
Kocuria kristinae		96.19	0.56		99			
Aerococcus viridans		3.27	0.47		1			
Rothia mucilaginosa		0.54	0.05		1			
121750	**Low discrimination**							**121750**
					OXI			
Aerococcus viridans		50.88	0.30	VPT 1	1			
Rothia mucilaginosa		19.68	-.04	LAC 90	1			
Kocuria kristinae		17.64	-.02		99			
121751	**Low discrimination**							**121751**
					OXI	NOV	PYR	
Kocuria kristinae		84.78	0.40	MAN 4	99	1	93	
Staphylococcus lentus		6.59	0.25		99	99	6	
Aerococcus viridans		5.77	0.38		1	1	95	
121752	**Acceptable identification**							**121752**
					NOV	YEP	PYR	
Aerococcus viridans		92.21	0.50	VPT 1	1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		5.22	-.04		99	16	7	
Staphylococcus kloosii		2.58	0.18		99	99	52	
121753	**Low discrimination**							**121753**
					OXI	NOV	PYR	
Aerococcus viridans		53.80	0.57	VPT 1	1	1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		24.22	0.36		99	99	6	
Kocuria kristinae		21.98	0.27		99	1	93	
121761	**Low discrimination**							**121761**
					OXI			
Kocuria kristinae		81.11	0.27	XYL 1	99			
Rothia mucilaginosa		11.31	0.05		1			
Aerococcus viridans		7.58	0.28		1			
121762	**Good identification**							**121762**
Aerococcus viridans		100.0	0.40	VPT 1				
121763	**Low discrimination**							**121763**
					OXI			
Aerococcus viridans		77.10	0.47	VPT 1	1			
Kocuria kristinae		22.91	0.15		99			
121770	**Good identification**							**121770**
Aerococcus viridans		100.0	0.30	VPT 1				
121772	**Low discrimination**							**121772**
					NOV	YEP		
Aerococcus viridans		70.43	0.50	VPT 1	1	1		
Staphylococcus arlettae		29.57	0.31		99	99		
121773	**Low discrimination**							**121773**
					OXI	NOV	YEP	
Aerococcus viridans		59.21	0.57	VPT 1	1	1	1	
Staphylococcus lentus		26.66	0.36		99	99	1	
Staphylococcus arlettae		14.13	0.34		1	99	99	
122000	**Low discrimination**							**122000**
					OXI			
Kocuria varians		52.41	0.58	VPT 14	13			
Kocuria rosea		39.89	0.47		35			
Dermacoccus nishinomiyaensis		4.10	0.28		93			
122001	**Low discrimination**							**122001**
Staphylococcus schleiferi subsp. schleiferi		38.73	0.33	ARG 86				
Kocuria varians		31.04	0.36	PHS 99				
Kocuria rosea		14.30	0.20					
122002	**Good identification**							**122002**
Kocuria varians		95.29	0.47	VPT 14				
Kocuria rosea		2.60	0.05					
Staphylococcus auricularis		2.12	0.11					
122010	**Low discrimination**							**122010**
Kocuria rosea		84.72	0.27	BGA 1				
Kocuria varians		10.12	0.17	MAN 10				
Staphylococcus capitis subsp. capitis		5.16	0.04					
122011	**Good identification**							**122011**
Staphylococcus capitis subsp. capitis		98.15	0.45	BGA 1				
Staphylococcus carnosus subsp. carnosus		0.99	0.05					
Kocuria rosea		0.86	0.01					
122020	**Low discrimination**							**122020**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		51.86	0.49	VPT 14			
Kocuria rosea		48.14	0.39				
122021	**Low discrimination**						**122021**
Kocuria varians		64.03	0.27	VPT 14			
Kocuria rosea		35.98	0.13	MNS 8			
122022	**Good identification**						**122022**
Kocuria varians		96.78	0.37	VPT 14			
Kocuria rosea		3.22	-.02				
122040	**Low discrimination**						**122040**
Kocuria varians		61.18	0.44	VPT 14			
Staphylococcus auricularis		36.67	0.39				
Kocuria rosea		2.14	0.05				
122042	**Low discrimination**						**122042**
Kocuria varians		80.90	0.33	VPT 14			
Staphylococcus auricularis		19.10	0.19				
122060	**Good identification**						**122060**
Kocuria varians		95.90	0.35	VPT 14			
Kocuria rosea		4.10	-.02				
122100	**Very good identification**						**122100**
					OXI		
Kocuria varians		98.11	0.62	VPT 14	13		
Dermacoccus nishinomiyaensis		0.98	0.13		93		
Kocuria rosea		0.50	0.05		35		
122101	**Low discrimination**						**122101**
Kocuria varians		54.59	0.40	VPT 14			
Staphylococcus schleiferi subsp. schleiferi		45.41	0.33	MNS 8			
122102	**Good identification**						**122102**
Kocuria varians		100.0	0.50	VPT 14			
122103	**Good identification**						**122103**
Kocuria varians		100.0	0.28	VPT 14			
122120	**Good identification**						**122120**
Kocuria varians		99.38	0.52	VPT 14			
Kocuria rosea		0.62	-.02				
122121	**Good identification**						**122121**
Kocuria varians		100.0	0.30	VPT 14			
122122	**Good identification**						**122122**
Kocuria varians		100.0	0.41	VPT 14			
122140	**Low discrimination**						**122140**
					NOV	OXI	
Kocuria varians		88.33	0.48	VPT 14	1	13	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Micrococcus caseolyticus		7.25	0.14		99	99	
Staphylococcus auricularis		3.92	0.19		1	1	
122142	**Low discrimination**						**122142**
					NOV	OXI	
Kocuria varians		68.36	0.37	VPT 14	1	13	
Micrococcus caseolyticus		29.82	0.18		99	99	
Staphylococcus epidermidis		1.82	0.02		1	1	
122160	**Good identification**						**122160**
Kocuria varians		100.0	0.39	VPT 14			
122162	**Good identification**						**122162**
Kocuria varians		100.0	0.27	VPT 14			
122200	**Low discrimination**						**122200**
Kocuria varians		49.06	0.40	VPT 14			
Kocuria rosea		30.42	0.27	SUC 12			
Staphylococcus auricularis		18.73	0.30				
122201	**Low discrimination**						**122201**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		53.83	0.28	BGA 1	1	1	
Kocuria kristinae		29.49	0.21	MAN 93	99	93	
Kocuria varians		8.96	0.18		13	10	
122202	**Low discrimination**						**122202**
Kocuria varians		86.93	0.29	VPT 14			
Staphylococcus auricularis		13.07	0.11	SUC 12			
122211	**Good identification**						**122211**
Staphylococcus capitis subsp. capitis		100.0	0.51	BGA 1			
122220	**Low discrimination**						**122220**
Kocuria varians		56.93	0.31	VPT 14			
Kocuria rosea		43.07	0.19	SUC 12			
122240	**Low discrimination**						**122240**
					PYR		
Staphylococcus haemolyticus		45.98	0.42	ARG 92	98		
Staphylococcus auricularis		34.17	0.38	TRE 93	40		
Kocuria varians		8.42	0.27		10		
122241	**Low discrimination**						**122241**
					OXI	PYR	
Kocuria kristinae		87.70	0.41	NIT 7	99	93	
Staphylococcus epidermidis		6.32	0.23	TRE 95	1	1	
Staphylococcus auricularis		3.37	0.18		1	40	
122242	**Low discrimination**						**122242**
					PYR		
Staphylococcus haemolyticus		55.40	0.36	ARG 92	98		
Staphylococcus epidermidis		26.00	0.28	TRE 93	1		
Staphylococcus auricularis		8.13	0.18		40		
122243	**Low discrimination**						**122243**
					OXI	PYR	
Kocuria kristinae		50.27	0.28	NIT 7	99	93	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		45.86	0.33	TRE 95	1	1		
Staphylococcus lugdunensis		2.25	0.02		1	96		
122250	**Good identification**							**122250**
Staphylococcus haemolyticus		100.0	0.37	ARG 92				
122252	**Good identification**							**122252**
Staphylococcus haemolyticus		100.0	0.32	ARG 92				
122300	**Low discrimination**							**122300**
Kocuria varians		89.50	0.44	VPT 14				
Rothia mucilaginosa		7.16	0.10	SUC 12				
Staphylococcus auricularis		2.53	0.11					
122302	**Good identification**							**122302**
Kocuria varians		95.90	0.32	VPT 14				
Staphylococcus epidermidis		4.11	0.02	SUC 12				
122320	**Good identification**							**122320**
Kocuria varians		100.0	0.34	VPT 14				
122340	**Low discrimination**							**122340**
					PYR			
Staphylococcus haemolyticus		44.65	0.43	ARG 92	98			
Staphylococcus epidermidis		18.71	0.33	TRE 93	1			
Kocuria varians		10.88	0.30		10			
122341	**Low discrimination**							**122341**
					OXI	PYR		
Staphylococcus epidermidis		36.47	0.39	URE 92	1	1		
Kocuria kristinae		29.79	0.31		99	93		
Rothia mucilaginosa		29.13	0.27		1	75		
122342	**Low discrimination**							**122342**
					OXI	PYR	NOV	
Staphylococcus epidermidis		59.75	0.43	URE 92	1	1	1	
Staphylococcus haemolyticus		25.33	0.38		1	98	2	
Macrococcus caseolyticus		6.60	0.14		99	80	99	
122343	**Acceptable identification**							**122343**
					OXI	NOV	PYR	
Staphylococcus epidermidis		91.72	0.49	URE 92	1	1	1	
Kocuria kristinae		5.91	0.19		99	1	93	
Staphylococcus hominis subsp. novobiosepticus		0.93	0.00		1	99	1	
122350	**Good identification**							**122350**
					NOV			
Staphylococcus haemolyticus		96.64	0.38	ARG 92	2			
Staphylococcus saprophyticus subsp. bovis		3.36	0.05	TRE 93	99			
122352	**Good identification**							**122352**
					PYR			
Staphylococcus haemolyticus		96.10	0.33	ARG 92	98			
Staphylococcus epidermidis		3.90	0.02	TRE 93	1			
122400	**Low discrimination**							**122400**
					NOV			
Staphylococcus auricularis		86.63	0.53	VPT 12	1			
Kocuria varians		10.06	0.35	BGA 14	1			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		1.85	0.22		99			
122401	**Low discrimination**							**122401**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		76.30	0.43	ARG 86	1			
Staphylococcus auricularis		16.89	0.33	PHS 99	1			
Staphylococcus cohnii subsp. cohnii		3.52	0.23		99			
122402	**Low discrimination**							**122402**
Staphylococcus auricularis		77.24	0.33	VPT 12				
Kocuria varians		22.76	0.24	BGA 14				
122440	**Good identification**							**122440**
					OXI	NOV	YEP	
Staphylococcus auricularis		92.70	0.60	VPT 12	1	1	1	
Staphylococcus cohnii subsp. cohnii		2.69	0.32	BGA 14	1	99	12	
Macrococcus caseolyticus		1.66	0.14		99	99	99	
122441	**Low discrimination**							**122441**
					OXI	NOV		
Staphylococcus auricularis		58.70	0.41	VPT 12	1	1		
Kocuria kristinae		23.54	0.26	BGA 14	99	1		
Staphylococcus cohnii subsp. cohnii		16.61	0.33	MNS 10	1	99		
122442	**Low discrimination**							**122442**
					OXI	NOV		
Staphylococcus auricularis		74.51	0.41	VPT 12	1	1		
Macrococcus caseolyticus		17.96	0.18	BGA 14	99	99		
Staphylococcus haemolyticus		5.47	0.19	LAC 10	1	2		
122450	**Low discrimination**							**122450**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		51.12	0.34	BGA 1	99	1		
Staphylococcus saprophyticus subsp. bovis		21.28	0.22	NIT 7	99	99		
Staphylococcus auricularis		13.98	0.19		1	40		
122451	**Low discrimination**							**122451**
					OXI	PYR	NOV	
Staphylococcus cohnii subsp. cohnii		87.53	0.35	BGA 1	1	1	99	
Staphylococcus saprophyticus subsp. bovis		8.41	0.09	NIT 7	1	99	99	
Kocuria kristinae		4.06	-.02		99	93	1	
122500	**Low discrimination**							**122500**
					NOV	OXI		
Kocuria varians		48.75	0.38	VPT 14	1	13		
Staphylococcus auricularis		31.11	0.33	TRE 7	1	1		
Macrococcus caseolyticus		11.66	0.14		99	99		
122501	**Low discrimination**							**122501**
Staphylococcus schleiferi subsp. schleiferi		82.43	0.43	ARG 86				
Rothia mucilaginosa		12.43	0.19	PHS 99				
Kocuria varians		2.49	0.17					
122520	**Good identification**							**122520**
Kocuria varians		100.0	0.29	VPT 14				
122540	**Good identification**							**122540**
					OXI	NOV		
Macrococcus caseolyticus		91.12	0.55	BGA 1	99	99		
Staphylococcus auricularis		5.73	0.41		1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Rothia mucilaginosa		1.30	0.17		1	1	
122541	**Low discrimination**						**122541**
					OXI	NOV	
Rothia mucilaginosa		59.21	0.27	ESL 90	1	1	
Macrococcus caseolyticus		13.96	0.14	SUC 99	99	99	
Kocuria kristinae		11.62	0.16		99	1	
122542	**Very good identification**						**122542**
					OXI	NOV	
Macrococcus caseolyticus		99.02	0.59	BGA 1	99	99	
Staphylococcus auricularis		0.46	0.21		1	1	
Staphylococcus haemolyticus		0.34	0.20		1	2	
122550	**Low discrimination**						**122550**
					OXI	NOV	
Staphylococcus saprophyticus subsp. bovis		66.09	0.34	URE 99	1	99	
Macrococcus caseolyticus		19.18	0.14		99	99	
Staphylococcus haemolyticus		10.32	0.20		1	2	
122600	**Acceptable identification**						**122600**
Staphylococcus auricularis		90.10	0.52	VPT 12			
Staphylococcus haemolyticus		3.30	0.24	BGA 14			
Rothia mucilaginosa		2.46	0.10				
122601	**Low discrimination**						**122601**
					OXI		
Kocuria kristinae		86.19	0.47	NIT 7	99		
Staphylococcus auricularis		7.73	0.33	MLT 90	1		
Rothia mucilaginosa		5.70	0.19		1		
122602	**Low discrimination**						**122602**
					PYR		
Staphylococcus auricularis		79.54	0.33	VPT 12	40		
Staphylococcus haemolyticus		14.75	0.19	BGA 14	98		
Kocuria varians		3.46	0.06	LAC 10	10		
122603	**Acceptable identification**						**122603**
					OXI		
Kocuria kristinae		93.38	0.35	NIT 7	99		
Staphylococcus auricularis		3.72	0.13	MLT 90	1		
Staphylococcus lugdunensis		1.98	0.02		1		
122640	**Low discrimination**						**122640**
Staphylococcus haemolyticus		45.74	0.65	ARG 92			
Staphylococcus auricularis		29.42	0.60				
Staphylococcus petrasii subsp. pragensis		20.27	0.39				
122641	**Good identification**						**122641**
					OXI		
Kocuria kristinae		94.39	0.67	NIT 7	99		
Staphylococcus auricularis		2.19	0.40		1		
Rothia mucilaginosa		1.62	0.27		1		
122642	**Low discrimination**						**122642**
					OXI	NOV	
Staphylococcus haemolyticus		83.51	0.60	ARG 92	1	2	
Staphylococcus auricularis		10.61	0.40		1	1	
Macrococcus caseolyticus		1.85	0.14		99	99	
122643	**Low discrimination**						**122643**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI				
Kocuria kristinae		78.86	0.55	NIT 7	99				
Staphylococcus lugdunensis		18.37	0.43		1				
Staphylococcus haemolyticus		1.79	0.29		1				
122650	**Good identification**								**122650**
					NOV				
Staphylococcus haemolyticus		90.03	0.60	ARG 92	2				
Staphylococcus saprophyticus subsp. bovis		6.57	0.34		99				
Staphylococcus auricularis		1.00	0.19		1				
122651	**Low discrimination**								**122651**
					OXI	NOV			
Kocuria kristinae		81.60	0.39	NIT 7	99	1			
Staphylococcus haemolyticus		11.57	0.29	MAN 4	1	2			
Staphylococcus saprophyticus subsp. bovis		6.83	0.22		1	99			
122652	**Low discrimination**								**122652**
					NOV	PYR			
Staphylococcus haemolyticus		89.98	0.55	ARG 92	2	98			
Staphylococcus saprophyticus subsp. saprophyticus		6.59	0.18		99	7			
Staphylococcus saprophyticus subsp. bovis		2.92	0.22		99	99			
122700	**Acceptable identification**								**122700**
Rothia mucilaginosa		92.72	0.51	ESL 90					
Staphylococcus auricularis		3.81	0.33						
Staphylococcus haemolyticus		1.42	0.25						
122701	**Good identification**								**122701**
					OXI				
Rothia mucilaginosa		94.96	0.60	ESL 90	1				
Kocuria kristinae		4.83	0.38		99				
Staphylococcus auricularis		0.14	0.13		1				
122713	**Low discrimination**								**122713**
					OXI	NOV	COA	PYR	
Staphylococcus lentus		86.13	0.30	VPT 7	99	99	1	6	
Staphylococcus intermedius		8.56	0.05	ESL 99	1	1	99	90	
Kocuria kristinae		5.31	-.03		99	1	1	93	
122733	**Good identification**								**122733**
Staphylococcus lentus		100.0	0.30	VPT 7					
122740	**Low discrimination**								**122740**
					OXI	NOV			
Rothia mucilaginosa		45.99	0.58	ESL 90	1	1			
Staphylococcus haemolyticus		29.79	0.66		1	2			
Macrococcus caseolyticus		21.69	0.52		99	99			
122741	**Low discrimination**								**122741**
					OXI				
Rothia mucilaginosa		82.63	0.68	ESL 90	1				
Kocuria kristinae		16.22	0.57		99				
Staphylococcus haemolyticus		0.55	0.35		1				
122742	**Low discrimination**								**122742**
					OXI	NOV			
Macrococcus caseolyticus		64.77	0.55	BGA 1	99	99			
Staphylococcus haemolyticus		33.36	0.61		1	2			
Rothia mucilaginosa		0.93	0.17		1	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
122743	**Low discrimination**							**122743**
					OXI			
Kocuria kristinae		48.78	0.45	NIT 7	99			
Staphylococcus lugdunensis		34.09	0.43		1			
Rothia mucilaginosa		10.04	0.27		1			
122750	**Low discrimination**							**122750**
					NOV			
Staphylococcus haemolyticus		76.89	0.61	ARG 92	2			
Staphylococcus saprophyticus subsp. bovis		19.90	0.47		99			
Rothia mucilaginosa		2.04	0.17		1			
122751	**Low discrimination**							**122751**
					OXI	NOV		
Rothia mucilaginosa		31.06	0.27	ESL 90	1	1		
Staphylococcus saprophyticus subsp. bovis		25.24	0.34	MAN 1	1	99		
Kocuria kristinae		25.15	0.29		99	1		
122752	**Low discrimination**							**122752**
					OXI	NOV		
Staphylococcus haemolyticus		86.65	0.56	ARG 92	1	2		
Staphylococcus saprophyticus subsp. bovis		9.97	0.34		1	99		
Macrococcus caseolyticus		2.89	0.14		99	99		
122753	**Low discrimination**							**122753**
					OXI	NOV	PYR	
Staphylococcus lentus		50.97	0.36	VPT 7	99	99	6	
Staphylococcus haemolyticus		15.03	0.25	ESL 99	1	2	98	
Staphylococcus saprophyticus subsp. bovis		13.98	0.22		1	99	99	
122761	**Low discrimination**							**122761**
					OXI			
Rothia mucilaginosa		83.60	0.27	ESL 90	1			
Kocuria kristinae		16.41	0.16	XYL 1	99			
122773	**Low discrimination**							**122773**
					OXI			
Staphylococcus lentus		81.62	0.36	VPT 7	99			
Staphylococcus xylosus		18.39	0.16	ESL 99	1			
123241	**Low discrimination**							**123241**
					OXI			
Kocuria kristinae		89.40	0.28	NIT 7	99			
Rothia mucilaginosa		10.60	0.05	TRE 95	1			
123300	**Good identification**							**123300**
Rothia mucilaginosa		98.62	0.29	TRE 99				
Kocuria varians		1.38	0.03					
123301	**Good identification**							**123301**
					OXI			
Rothia mucilaginosa		99.27	0.39	TRE 99	1			
Kocuria kristinae		0.73	-.01		99			
123313	**Good identification**							**123313**
Staphylococcus lentus		100.0	0.30	VPT 7				
123333	**Good identification**							**123333**
Staphylococcus lentus		100.0	0.30	VPT 7				
123340	**Good identification**							**123340**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Rothia mucilaginosa		98.62	0.37	TRE 99	1	75	
Staphylococcus hominis subsp. novobiosepticus		0.85	-.04		99	1	
Staphylococcus haemolyticus		0.53	0.02		2	98	
123341	**Good identification**						**123341**
					OXI		
Rothia mucilaginosa		97.24	0.47	TRE 99	1		
Kocuria kristinae		2.76	0.19		99		
123353	**Good identification**						**123353**
Staphylococcus lentus		100.0	0.36	VPT 7			
123373	**Good identification**						**123373**
Staphylococcus lentus		100.0	0.36	VPT 7			
123500	**Good identification**						**123500**
Rothia mucilaginosa		100.0	0.29	SUC 99			
123501	**Good identification**						**123501**
Rothia mucilaginosa		99.26	0.39	SUC 99			
Staphylococcus schleiferi subsp. schleiferi		0.74	0.02				
123513	**Good identification**						**123513**
Staphylococcus lentus		100.0	0.30	VPT 7			
123533	**Good identification**						**123533**
Staphylococcus lentus		100.0	0.30	VPT 7			
123540	**Acceptable identification**						**123540**
					OXI	NOV	
Rothia mucilaginosa		92.71	0.37	SUC 99	1	1	
Macrococcus caseolyticus		7.29	0.14		99	99	
123541	**Good identification**						**123541**
					OXI		
Rothia mucilaginosa		99.46	0.47	SUC 99	1		
Kocuria kristinae		0.54	0.04		99		
123553	**Good identification**						**123553**
					OXI	NOV	
Staphylococcus lentus		97.81	0.36	VPT 7	99	99	
Aerococcus viridans		2.19	0.16	SUC 99	1	1	
123573	**Good identification**						**123573**
					OXI	NOV	
Staphylococcus lentus		97.81	0.36	VPT 7	99	99	
Aerococcus viridans		2.19	0.16	SUC 99	1	1	
123600	**Acceptable identification**						**123600**
					OXI		
Rothia mucilaginosa		94.90	0.29	GAL 99	1		
Staphylococcus auricularis		3.90	0.11		1		
Kocuria kristinae		1.21	-.06		99		
123601	**Low discrimination**						**123601**
					OXI		
Rothia mucilaginosa		70.44	0.39	GAL 99	1		
Kocuria kristinae		29.57	0.35		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
123613	**Good identification**							**123613**
					NOV	PYR		
Staphylococcus lentus		95.58	0.30	VPT 7	99	6		
Kocuria kristinae		4.42	-.06	GAL 99	1	93		
123633	**Good identification**							**123633**
Staphylococcus lentus		100.0	0.30	VPT 7				
123640	**Low discrimination**							**123640**
					OXI			
Rothia mucilaginosa		84.59	0.37	GAL 99	1			
Staphylococcus haemolyticus		5.40	0.24		1			
Kocuria kristinae		4.15	0.14		99			
123641	**Low discrimination**							**123641**
					OXI			
Kocuria kristinae		61.82	0.55	NIT 7	99			
Rothia mucilaginosa		38.18	0.47		1			
123643	**Good identification**							**123643**
					OXI			
Kocuria kristinae		96.68	0.42	NIT 7	99			
Rothia mucilaginosa		2.41	0.05		1			
Staphylococcus lugdunensis		0.91	0.02		1			
123653	**Low discrimination**							**123653**
					OXI	NOV	PYR	
Staphylococcus lentus		81.48	0.36	VPT 7	99	99	6	
Kocuria kristinae		16.69	0.14	GAL 99	99	1	93	
Aerococcus viridans		1.83	0.16		1	1	95	
123673	**Good identification**							**123673**
					OXI	NOV		
Staphylococcus lentus		97.81	0.36	VPT 7	99	99		
Aerococcus viridans		2.19	0.16	GAL 99	1	1		
123700	**Very good identification**							**123700**
Rothia mucilaginosa		100.0	0.70					
123701	**Very good identification**							**123701**
					OXI	NOV	PYR	
Rothia mucilaginosa		99.83	0.80		1	1	75	
Kocuria kristinae		0.14	0.25		99	1	93	
Staphylococcus lentus		0.03	0.18		99	99	6	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
123702	**Good identification**							**123702**
Rothia mucilaginosa		100.0	0.29	LAC 1				
123703	**Low discrimination**							**123703**
					OXI	NOV	PYR	
Rothia mucilaginosa		88.53	0.39	LAC 1	1	1	75	
Staphylococcus lentus		8.38	0.30		99	99	6	
Kocuria kristinae		3.10	0.13		99	1	93	
123710	**Acceptable identification**							**123710**
					OXI	NOV		
Rothia mucilaginosa		92.59	0.29	MAN 1	1	1		
Staphylococcus lentus		7.41	0.18		99	99		
123711	**Low discrimination**							**123711**
					OXI	NOV	PYR	
Staphylococcus lentus		72.33	0.59	VPT 7	99	99	6	
Rothia mucilaginosa		27.38	0.39		1	1	75	
Kocuria kristinae		0.16	-.03		99	1	93	
123712	**Good identification**							**123712**
Staphylococcus lentus		100.0	0.30	VPT 7				
123713	**Very good identification**							**123713**
					OXI	NOV		
Staphylococcus lentus		99.89	0.71	VPT 7	99	99		
Rothia mucilaginosa		0.11	-.02		1	1		
123720	**Good identification**							**123720**
Rothia mucilaginosa		100.0	0.29	XYL 1				
123721	**Good identification**							**123721**
					OXI	NOV		
Rothia mucilaginosa		97.40	0.39	XYL 1	1	1		
Staphylococcus lentus		2.60	0.18		99	99		
123723	**Acceptable identification**							**123723**
					OXI	NOV		
Staphylococcus lentus		90.35	0.30	VPT 7	99	99		
Rothia mucilaginosa		9.65	-.02	MAN 99	1	1		
123731	**Very good identification**							**123731**
					OXI	NOV		
Staphylococcus lentus		99.62	0.59	VPT 7	99	99		
Rothia mucilaginosa		0.38	-.02		1	1		
123732	**Good identification**							**123732**
Staphylococcus lentus		100.0	0.30	VPT 7				
123733	**Very good identification**							**123733**
Staphylococcus lentus		100.0	0.71	VPT 7				
123740	**Very good identification**							**123740**
					OXI	NOV		
Rothia mucilaginosa		99.86	0.78		1	1		
Staphylococcus haemolyticus		0.07	0.25		1	2		
Micrococcus caseolyticus		0.05	0.11		99	99		
123741	**Very good identification**							**123741**
					OXI	NOV	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Rothia mucilaginosa		99.44	0.88		1	1	75	
Kocuria kristinae		0.54	0.45		99	1	93	
Staphylococcus lentus		0.02	0.25		99	99	6	
123742	**Low discrimination**							**123742**
					OXI	NOV		
Rothia mucilaginosa		89.36	0.37	LAC 1	1	1		
Macrococcus caseolyticus		7.02	0.14		99	99		
Staphylococcus haemolyticus		3.62	0.20		1	2		
123743	**Low discrimination**							**123743**
					OXI	NOV	PYR	
Rothia mucilaginosa		81.90	0.47	LAC 1	1	1	75	
Kocuria kristinae		11.05	0.32		99	1	93	
Staphylococcus lentus		6.74	0.36		99	99	6	
123750	**Low discrimination**							**123750**
					OXI	NOV		
Rothia mucilaginosa		89.06	0.37	MAN 1	1	1		
Staphylococcus lentus		6.20	0.25		99	99		
Staphylococcus haemolyticus		3.77	0.20		1	2		
123751	**Low discrimination**							**123751**
					OXI	NOV	PYR	
Staphylococcus lentus		69.05	0.66	VPT 7	99	99	6	
Rothia mucilaginosa		30.04	0.47		1	1	75	
Kocuria kristinae		0.68	0.16		99	1	93	
123752	**Low discrimination**							**123752**
					OXI	NOV		
Staphylococcus lentus		87.94	0.36	VPT 7	99	99		
Staphylococcus haemolyticus		8.47	0.15	MNS 99	1	2		
Rothia mucilaginosa		3.60	-.04		1	1		
123753	**Very good identification**							**123753**
					OXI	NOV	PYR	
Staphylococcus lentus		99.79	0.77	VPT 7	99	99	6	
Rothia mucilaginosa		0.12	0.05		1	1	75	
Kocuria kristinae		0.07	0.04		99	1	93	
123760	**Good identification**							**123760**
Rothia mucilaginosa		100.0	0.37	XYL 1				
123761	**Good identification**							**123761**
					OXI	NOV	PYR	
Rothia mucilaginosa		97.21	0.47	XYL 1	1	1	75	
Staphylococcus lentus		2.26	0.25		99	99	6	
Kocuria kristinae		0.53	0.04		99	1	93	
123763	**Low discrimination**							**123763**
					OXI	NOV		
Staphylococcus lentus		89.07	0.36	VPT 7	99	99		
Rothia mucilaginosa		10.93	0.05	MAN 99	1	1		
123771	**Very good identification**							**123771**
					OXI	NOV		
Staphylococcus lentus		99.31	0.66	VPT 7	99	99		
Rothia mucilaginosa		0.44	0.05		1	1		
Staphylococcus sciuri		0.26	0.11		99	99		
123772	**Good identification**							**123772**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lentus		100.0	0.36	VPT 7			
123773	**Very good identification**						**123773**
					OXI	NOV	
Staphylococcus lentus		99.92	0.77	VPT 7	99	99	
Staphylococcus xylosus		0.06	0.04		1	99	
Aerococcus viridans		0.02	0.16		1	1	
124001	**Very good identification**						**124001**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.54	ARG 86			
124050	**Good identification**						**124050**
Staphylococcus kloosii		100.0	0.32	VPT 10			
124052	**Good identification**						**124052**
Staphylococcus kloosii		100.0	0.33	VPT 10			
124101	**Very good identification**						**124101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.54	ARG 86			
124150	**Good identification**						**124150**
Staphylococcus kloosii		100.0	0.38	VPT 10			
124152	**Good identification**						**124152**
Staphylococcus kloosii		100.0	0.39	VPT 10			
124172	**Good identification**						**124172**
Staphylococcus kloosii		100.0	0.27	VPT 10			
124342	**Good identification**						**124342**
Staphylococcus epidermidis		100.0	0.28	URE 92			
124343	**Good identification**						**124343**
					OXI	PYR	
Staphylococcus epidermidis		95.30	0.33	URE 92	1	1	
Kocuria kristinae		4.71	0.01	NIT 97	99	93	
124401	**Very good identification**						**124401**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		99.45	0.64	ARG 86	1		
Staphylococcus cohnii subsp. cohnii		0.55	0.25	NIT 90	99		
124410	**Low discrimination**						**124410**
					YEP		
Staphylococcus kloosii		70.92	0.47	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		29.08	0.27	MLT 95	12		
124412	**Good identification**						**124412**
Staphylococcus kloosii		100.0	0.48	VPT 10			
124430	**Good identification**						**124430**
Staphylococcus kloosii		100.0	0.35	VPT 10			
124432	**Good identification**						**124432**
Staphylococcus kloosii		100.0	0.35	VPT 10			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
124440	**Low discrimination**						**124440**
					YEP		
Staphylococcus cohnii subsp. cohnii		84.17	0.35	BGA 1	12		
Staphylococcus kloosii		15.83	0.32	PHS 9	99		
124441	**Low discrimination**						**124441**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		58.02	0.35	BGA 1	1	99	
Staphylococcus schleiferi subsp. schleiferi		33.58	0.23	PHS 9	1	1	
Kocuria kristinae		8.40	0.08		99	1	
124442	**Good identification**						**124442**
Staphylococcus kloosii		100.0	0.33	VPT 10			
124450	**Good identification**						**124450**
					YEP		
Staphylococcus kloosii		93.60	0.73	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		6.40	0.37		12		
124451	**Low discrimination**						**124451**
					YEP		
Staphylococcus cohnii subsp. cohnii		88.00	0.38	BGA 1	12		
Staphylococcus kloosii		12.00	0.32	PHS 9	99		
124452	**Very good identification**						**124452**
Staphylococcus kloosii		99.56	0.74	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		0.45	-.03				
124453	**Low discrimination**						**124453**
Staphylococcus kloosii		71.22	0.33	VPT 10			
Staphylococcus cohnii subsp. urealyticum		28.79	0.13	MNS 1			
124470	**Very good identification**						**124470**
Staphylococcus kloosii		100.0	0.61	VPT 10			
124472	**Very good identification**						**124472**
Staphylococcus kloosii		100.0	0.62	VPT 10			
124501	**Very good identification**						**124501**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		99.94	0.64	ARG 86	1		
Staphylococcus cohnii subsp. cohnii		0.06	0.05	NIT 90	99		
124510	**Very good identification**						**124510**
					YEP		
Staphylococcus kloosii		97.71	0.53	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		2.29	0.07	MLT 95	12		
124512	**Very good identification**						**124512**
Staphylococcus kloosii		100.0	0.54	VPT 10			
124530	**Good identification**						**124530**
Staphylococcus kloosii		100.0	0.41	VPT 10			
124532	**Good identification**						**124532**
Staphylococcus kloosii		100.0	0.41	VPT 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
124540	**Low discrimination**					**124540**
					YEP	
Staphylococcus kloosii		76.67	0.38	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		23.33	0.15	MAN 99	12	
124542	**Good identification**					**124542**
Staphylococcus kloosii		100.0	0.39	VPT 10		
124550	**Very good identification**					**124550**
					YEP	
Staphylococcus kloosii		99.61	0.79	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		0.39	0.17		12	
124551	**Low discrimination**					**124551**
					YEP	
Staphylococcus kloosii		70.44	0.38	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		29.56	0.18	MNS 1	12	
124552	**Very good identification**					**124552**
Staphylococcus kloosii		100.0	0.80	VPT 10		
124553	**Acceptable identification**					**124553**
Staphylococcus kloosii		91.81	0.39	VPT 10		
Staphylococcus cohnii subsp. urealyticum		8.19	0.06	MNS 1		
124562	**Good identification**					**124562**
Staphylococcus kloosii		100.0	0.27	VPT 10		
124570	**Very good identification**					**124570**
Staphylococcus kloosii		100.0	0.67	VPT 10		
124572	**Very good identification**					**124572**
Staphylococcus kloosii		100.0	0.68	VPT 10		
124573	**Good identification**					**124573**
Staphylococcus kloosii		100.0	0.27	VPT 10		
124601	**Low discrimination**					**124601**
					OXI	
Kocuria kristinae		73.35	0.30	PHS 1	99	
Staphylococcus schleiferi subsp. schleiferi		26.65	0.23	MLT 90	1	
124612	**Good identification**					**124612**
Staphylococcus kloosii		100.0	0.27	VPT 10		
124641	**Good identification**					**124641**
Kocuria kristinae		100.0	0.49	PHS 1		
124643	**Good identification**					**124643**
Kocuria kristinae		100.0	0.37	PHS 1		
124650	**Acceptable identification**					**124650**
Staphylococcus kloosii		91.97	0.53	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		8.03	0.02	SUC 9		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
124652	**Low discrimination**						**124652**
Staphylococcus kloosii		68.65	0.53	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		31.35	0.18	SUC 9			
124670	**Good identification**						**124670**
Staphylococcus kloosii		100.0	0.40	VPT 10			
124672	**Low discrimination**						**124672**
Staphylococcus kloosii		76.22	0.41	VPT 10			
Staphylococcus arlettae		23.78	0.19	SUC 9			
124710	**Good identification**						**124710**
Staphylococcus kloosii		100.0	0.32	VPT 10			
124712	**Good identification**						**124712**
Staphylococcus kloosii		100.0	0.33	VPT 10			
124741	**Good identification**						**124741**
Kocuria kristinae		100.0	0.39	PHS 1			
124743	**Acceptable identification**						**124743**
					OXI	COA	
Kocuria kristinae		93.01	0.27	PHS 1	99	1	
Staphylococcus intermedius		6.99	0.07		1	99	
124750	**Very good identification**						**124750**
Staphylococcus kloosii		100.0	0.59	VPT 10			
124752	**Very good identification**						**124752**
Staphylococcus kloosii		98.00	0.59	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		2.01	-.04	SUC 9			
124770	**Good identification**						**124770**
Staphylococcus kloosii		100.0	0.46	VPT 10			
124772	**Low discrimination**						**124772**
Staphylococcus kloosii		52.34	0.47	VPT 10			
Staphylococcus arlettae		47.66	0.35	SUC 9			
124773	**Good identification**						**124773**
Staphylococcus arlettae		100.0	0.37	VPT 1			
125050	**Good identification**						**125050**
Staphylococcus kloosii		100.0	0.32	VPT 10			
125052	**Good identification**						**125052**
Staphylococcus kloosii		100.0	0.33	VPT 10			
125150	**Good identification**						**125150**
Staphylococcus kloosii		100.0	0.38	VPT 10			
125152	**Good identification**						**125152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.39	VPT 10			
125172	**Good identification**						**125172**
Staphylococcus kloosii		100.0	0.27	VPT 10			
125410	**Good identification**						**125410**
Staphylococcus kloosii		100.0	0.47	VPT 10			
125412	**Good identification**						**125412**
Staphylococcus kloosii		100.0	0.48	VPT 10			
125430	**Good identification**						**125430**
Staphylococcus kloosii		100.0	0.35	VPT 10			
125432	**Good identification**						**125432**
Staphylococcus kloosii		100.0	0.35	VPT 10			
125440	**Good identification**						**125440**
Staphylococcus kloosii		100.0	0.32	VPT 10			
125442	**Good identification**						**125442**
Staphylococcus kloosii		100.0	0.33	VPT 10			
125450	**Very good identification**						**125450**
Staphylococcus kloosii		100.0	0.73	VPT 10			
125451	**Good identification**						**125451**
Staphylococcus kloosii		100.0	0.32	VPT 10			
125452	**Very good identification**						**125452**
Staphylococcus kloosii		100.0	0.74	VPT 10			
125453	**Low discrimination**						**125453**
					NOV	YEP	
Staphylococcus kloosii		75.81	0.33	VPT 10	99	99	
Aerococcus viridans		12.28	0.16	MNS 1	1	1	
Staphylococcus cohnii subsp. urealyticum		11.92	0.05		99	64	
125470	**Very good identification**						**125470**
Staphylococcus kloosii		100.0	0.61	VPT 10			
125472	**Very good identification**						**125472**
Staphylococcus kloosii		100.0	0.62	VPT 10			
125510	**Very good identification**						**125510**
Staphylococcus kloosii		100.0	0.53	VPT 10			
125512	**Very good identification**						**125512**
Staphylococcus kloosii		100.0	0.54	VPT 10			
125530	**Good identification**						**125530**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.41	VPT 10			
125532	**Good identification**						**125532**
Staphylococcus kloosii		100.0	0.41	VPT 10			
125540	**Good identification**						**125540**
Staphylococcus kloosii		100.0	0.38	VPT 10			
125542	**Good identification**						**125542**
Staphylococcus kloosii		100.0	0.39	VPT 10			
125550	**Very good identification**						**125550**
Staphylococcus kloosii		100.0	0.79	VPT 10			
125551	**Good identification**						**125551**
Staphylococcus kloosii		100.0	0.38	VPT 10			
125552	**Very good identification**						**125552**
Staphylococcus kloosii		100.0	0.80	VPT 10			
125553	**Acceptable identification**						**125553**
					NOV	YEP	
Staphylococcus kloosii		92.30	0.39	VPT 10	99	99	
Aerococcus viridans		7.70	0.16	MNS 1	1	1	
125562	**Good identification**						**125562**
Staphylococcus kloosii		100.0	0.27	VPT 10			
125570	**Very good identification**						**125570**
Staphylococcus kloosii		100.0	0.67	VPT 10			
125572	**Very good identification**						**125572**
Staphylococcus kloosii		100.0	0.68	VPT 10			
125573	**Low discrimination**						**125573**
					NOV	YEP	
Staphylococcus kloosii		74.98	0.27	VPT 10	99	99	
Aerococcus viridans		25.02	0.16	MNS 1	1	1	
125612	**Good identification**						**125612**
Staphylococcus kloosii		100.0	0.27	VPT 10			
125641	**Good identification**						**125641**
Kocuria kristinae		100.0	0.37	PHS 1			
125650	**Good identification**						**125650**
Staphylococcus kloosii		100.0	0.53	VPT 10			
125652	**Very good identification**						**125652**
Staphylococcus kloosii		100.0	0.53	VPT 10			
125670	**Good identification**						**125670**
Staphylococcus kloosii		100.0	0.40	VPT 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
125672	**Low discrimination**							**125672**
Staphylococcus kloosii		76.22	0.41	VPT 10				
Staphylococcus arlettae		23.78	0.19	SUC 9				
125710	**Good identification**							**125710**
Staphylococcus kloosii		100.0	0.32	VPT 10				
125712	**Good identification**							**125712**
Staphylococcus kloosii		100.0	0.33	VPT 10				
125741	**Low discrimination**							**125741**
					OXI			
Kocuria kristinae		87.76	0.27	PHS 1	99			
Rothia mucilaginosa		12.24	0.05		1			
125750	**Very good identification**							**125750**
Staphylococcus kloosii		100.0	0.59	VPT 10				
125752	**Very good identification**							**125752**
Staphylococcus kloosii		100.0	0.59	VPT 10				
125770	**Good identification**							**125770**
Staphylococcus kloosii		100.0	0.46	VPT 10				
125772	**Low discrimination**							**125772**
Staphylococcus kloosii		52.34	0.47	VPT 10				
Staphylococcus arlettae		47.66	0.35	SUC 9				
125773	**Low discrimination**							**125773**
					OXI	NOV	YEP	
Staphylococcus arlettae		74.47	0.37	VPT 1	1	99	99	
Staphylococcus lentus		23.43	0.23	GLR 92	99	99	1	
Aerococcus viridans		2.10	0.16		1	1	1	
126000	**Low discrimination**							**126000**
Staphylococcus schleiferi subsp. schleiferi		83.20	0.33	ARG 86				
Kocuria varians		7.75	0.17	MNS 99				
Kocuria rosea		5.89	0.05					
126001	**Very good identification**							**126001**
Staphylococcus schleiferi subsp. schleiferi		98.78	0.74	ARG 86				
Staphylococcus carnosus subsp. carnosus		1.17	0.34					
Staphylococcus capitis subsp. capitis		0.04	0.04					
126003	**Low discrimination**							**126003**
Staphylococcus carnosus subsp. carnosus		82.46	0.47	ARG 99				
Staphylococcus schleiferi subsp. schleiferi		17.54	0.33					
126011	**Low discrimination**							**126011**
Staphylococcus carnosus subsp. carnosus		74.84	0.47	ARG 99				
Staphylococcus schleiferi subsp. schleiferi		15.92	0.33					
Staphylococcus capitis subsp. capitis		9.24	0.27					
126013	**Very good identification**							**126013**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		100.0	0.59	ARG 99			
126021	**Good identification**						**126021**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86			
126041	**Good identification**						**126041**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86			
126100	**Low discrimination**						**126100**
Staphylococcus schleiferi subsp. schleiferi		87.75	0.33	ARG 86			
Kocuria varians		12.25	0.21	MNS 99			
126101	**Very good identification**						**126101**
					COA		
Staphylococcus schleiferi subsp. schleiferi		99.68	0.74	ARG 86	1		
Staphylococcus carnosus subsp. carnosus		0.30	0.22		1		
Staphylococcus schleiferi subsp. coagulans		0.02	0.02		99		
126103	**Low discrimination**						**126103**
					COA		
Staphylococcus carnosus subsp. carnosus		51.20	0.34	ARG 99	1		
Staphylococcus schleiferi subsp. schleiferi		43.56	0.33		1		
Staphylococcus schleiferi subsp. coagulans		5.25	0.18		99		
126111	**Low discrimination**						**126111**
					COA		
Staphylococcus carnosus subsp. carnosus		53.59	0.34	ARG 99	1		
Staphylococcus schleiferi subsp. schleiferi		45.59	0.33		1		
Staphylococcus schleiferi subsp. coagulans		0.83	0.01		99		
126113	**Good identification**						**126113**
					COA		
Staphylococcus carnosus subsp. carnosus		97.69	0.47	ARG 99	1		
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99		
126121	**Good identification**						**126121**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86			
126141	**Acceptable identification**						**126141**
Staphylococcus schleiferi subsp. schleiferi		92.97	0.33	ARG 86			
Staphylococcus epidermidis		7.03	0.13	MLT 1			
126201	**Acceptable identification**						**126201**
Staphylococcus schleiferi subsp. schleiferi		92.49	0.33	ARG 86			
Staphylococcus capitis subsp. capitis		7.51	0.09	SUC 1			
126211	**Good identification**						**126211**
Staphylococcus capitis subsp. capitis		95.78	0.32	BGA 1			
Staphylococcus carnosus subsp. carnosus		4.22	0.05	PHS 11			
126240	**Good identification**						**126240**
Staphylococcus epidermidis		100.0	0.33	URE 92			
126241	**Good identification**						**126241**
					OXI	PYR	
Staphylococcus epidermidis		97.59	0.39	URE 92	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria kristinae		2.42	0.00		99	93	
126242	**Good identification**						**126242**
Staphylococcus epidermidis		100.0	0.43	URE 92			
126243	**Good identification**						**126243**
Staphylococcus epidermidis		100.0	0.49	URE 92			
126301	**Acceptable identification**						**126301**
Staphylococcus schleiferi subsp. schleiferi		92.97	0.33	ARG 86			
Staphylococcus epidermidis		7.03	0.13	SUC 1			
126340	**Good identification**						**126340**
					PYR		
Staphylococcus epidermidis		99.58	0.49	URE 92	1		
Staphylococcus haemolyticus		0.42	0.02		98		
126341	**Very good identification**						**126341**
Staphylococcus epidermidis		100.0	0.54	URE 92			
126342	**Very good identification**						**126342**
Staphylococcus epidermidis		100.0	0.59	URE 92			
126343	**Very good identification**						**126343**
					COA	PYR	
Staphylococcus epidermidis		99.78	0.64	URE 92	1	1	
Staphylococcus intermedius		0.22	0.07		99	90	
126400	**Good identification**						**126400**
Staphylococcus schleiferi subsp. schleiferi		98.03	0.43	ARG 86			
Staphylococcus auricularis		1.97	0.12	MNS 99			
126401	**Very good identification**						**126401**
Staphylococcus schleiferi subsp. schleiferi		99.83	0.84	ARG 86			
Staphylococcus carnosus subsp. carnosus		0.17	0.27				
126403	**Low discrimination**						**126403**
Staphylococcus schleiferi subsp. schleiferi		59.82	0.43	ARG 86			
Staphylococcus carnosus subsp. carnosus		40.18	0.39	LAC 1			
126411	**Low discrimination**						**126411**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		59.61	0.43	ARG 86	1		
Staphylococcus carnosus subsp. carnosus		40.04	0.39	MAN 1	1		
Staphylococcus cohnii subsp. cohnii		0.35	0.04		99		
126413	**Good identification**						**126413**
Staphylococcus carnosus subsp. carnosus		99.63	0.51	ARG 99			
Staphylococcus schleiferi subsp. schleiferi		0.38	0.02				
126421	**Good identification**						**126421**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86			
126441	**Good identification**						**126441**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		98.58	0.43	ARG 86	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. cohnii		1.43	0.12	MLT 1	99		
126450	**Low discrimination**						**126450**
					PYR	YEP	
Staphylococcus kloosii		48.89	0.32	VPT 10	52	99	
Staphylococcus saprophyticus subsp. bovis		26.21	0.09	NIT 1	99	80	
Staphylococcus cohnii subsp. cohnii		24.90	0.14		1	12	
126452	**Good identification**						**126452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
126500	**Good identification**						**126500**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86			
126501	**Very good identification**						**126501**
Staphylococcus schleiferi subsp. schleiferi		99.96	0.84	ARG 86			
Staphylococcus carnosus subsp. carnosus		0.04	0.14				
126503	**Low discrimination**						**126503**
Staphylococcus schleiferi subsp. schleiferi		85.62	0.43	ARG 86			
Staphylococcus carnosus subsp. carnosus		14.38	0.27	LAC 1			
126511	**Low discrimination**						**126511**
Staphylococcus schleiferi subsp. schleiferi		85.62	0.43	ARG 86			
Staphylococcus carnosus subsp. carnosus		14.38	0.27	MAN 1			
126513	**Good identification**						**126513**
Staphylococcus carnosus subsp. carnosus		98.52	0.39	ARG 99			
Staphylococcus schleiferi subsp. schleiferi		1.48	0.02				
126521	**Good identification**						**126521**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86			
126541	**Good identification**						**126541**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86			
126552	**Low discrimination**						**126552**
Staphylococcus kloosii		79.69	0.39	VPT 10			
Staphylococcus saprophyticus subsp. bovis		20.31	0.09	NIT 1			
126572	**Good identification**						**126572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
126601	**Good identification**						**126601**
					OXI		
Staphylococcus schleiferi subsp. schleiferi		97.75	0.43	ARG 86	1		
Kocuria kristinae		2.25	0.06	SUC 1	99		
126701	**Low discrimination**						**126701**
					COA		
Staphylococcus schleiferi subsp. schleiferi		84.51	0.43	ARG 86	1		
Rothia mucilaginosa		12.74	0.19	SUC 1	1		
Staphylococcus intermedius		2.10	0.11		99		
126703	**Low discrimination**						**126703**
					COA	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		89.17	0.27	VPT 1	99	90		
Staphylococcus schleiferi subsp. schleiferi		5.90	0.02	URE 90	1	37		
Staphylococcus hyicus		4.93	-.05		24	1		
126713	**Low discrimination**							**126713**
					OXI	COA	NOV	
Staphylococcus intermedius		89.24	0.36	VPT 1	1	99	1	
Staphylococcus lentus		6.94	0.17	URE 90	99	1	99	
Staphylococcus felis		2.35	-.04		1	1	1	
126741	**Low discrimination**							**126741**
					OXI	COA		
Rothia mucilaginosa		71.30	0.27	ESL 90	1	1		
Kocuria kristinae		13.99	0.16	PHS 1	99	1		
Staphylococcus intermedius		7.55	0.14		1	99		
126743	**Low discrimination**							**126743**
					OXI	COA	PYR	
Staphylococcus intermedius		67.72	0.30	VPT 1	1	99	90	
Staphylococcus epidermidis		23.61	0.23	URE 90	1	1	1	
Kocuria kristinae		5.11	0.04		99	1	93	
126750	**Low discrimination**							**126750**
					NOV			
Staphylococcus saprophyticus subsp. bovis		84.86	0.34	URE 99	99			
Staphylococcus haemolyticus		13.24	0.20		2			
Staphylococcus kloosii		1.90	0.18		99			
126753	**Low discrimination**							**126753**
					OXI	PYR	COA	
Staphylococcus intermedius		83.86	0.39	VPT 1	1	90	99	
Staphylococcus lentus		8.83	0.23	URE 90	99	6	1	
Staphylococcus aureus subsp. aureus		4.03	-.02		1	10	99	
127001	**Low discrimination**							**127001**
Staphylococcus schleiferi subsp. schleiferi		88.45	0.33	ARG 86				
Staphylococcus carnosus subsp. carnosus		11.55	0.15	ESL 1				
127003	**Good identification**							**127003**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ARG 99				
127011	**Good identification**							**127011**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ARG 99				
127013	**Good identification**							**127013**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ARG 99				
127101	**Good identification**							**127101**
					COA			
Staphylococcus schleiferi subsp. schleiferi		95.10	0.33	ARG 86	1			
Staphylococcus carnosus subsp. carnosus		3.11	0.02	ESL 1	1			
Staphylococcus schleiferi subsp. coagulans		1.79	0.01		99			
127113	**Low discrimination**							**127113**
					COA			
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ARG 99	1			
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	ESL 10	99			
127401	**Good identification**							**127401**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. schleiferi		98.17	0.43	ARG 86			
Staphylococcus carnosus subsp. carnosus		1.83	0.07	ESL 1			
127413	**Good identification**						**127413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ARG 99			
127450	**Good identification**						**127450**
Staphylococcus kloosii		100.0	0.32	VPT 10			
127452	**Good identification**						**127452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
127501	**Good identification**						**127501**
Staphylococcus schleiferi subsp. schleiferi		98.65	0.43	ARG 86			
Rothia mucilaginosa		1.35	-.02	ESL 1			
127550	**Good identification**						**127550**
Staphylococcus kloosii		100.0	0.38	VPT 10			
127552	**Good identification**						**127552**
Staphylococcus kloosii		100.0	0.39	VPT 10			
127572	**Good identification**						**127572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
127700	**Good identification**						**127700**
Rothia mucilaginosa		100.0	0.29	PHS 1			
127701	**Good identification**						**127701**
					OXI	NOV	
Rothia mucilaginosa		98.61	0.39	PHS 1	1	1	
Staphylococcus schleiferi subsp. schleiferi		0.73	0.02		1	1	
Staphylococcus lentus		0.66	0.06		99	99	
127711	**Good identification**						**127711**
					OXI	NOV	
Staphylococcus lentus		97.28	0.47	VPT 7	99	99	
Rothia mucilaginosa		1.49	-.02		1	1	
Staphylococcus sciuri		1.23	0.06		99	99	
127713	**Very good identification**						**127713**
Staphylococcus lentus		100.0	0.58	VPT 7			
127731	**Good identification**						**127731**
Staphylococcus lentus		99.06	0.47	VPT 7			
Staphylococcus sciuri		0.94	0.04				
127733	**Very good identification**						**127733**
Staphylococcus lentus		100.0	0.58	VPT 7			
127740	**Good identification**						**127740**
Rothia mucilaginosa		100.0	0.37	PHS 1			
127741	**Good identification**						**127741**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	PYR	
Rothia mucilaginosa		98.89	0.47	PHS 1	1	1	75	
Staphylococcus lentus		0.57	0.12		99	99	6	
Kocuria kristinae		0.54	0.04		99	1	93	
127751	**Very good identification**							**127751**
					OXI	NOV		
Staphylococcus lentus		96.06	0.53	VPT 7	99	99		
Staphylococcus sciuri		2.25	0.18		99	99		
Rothia mucilaginosa		1.69	0.05		1	1		
127753	**Very good identification**							**127753**
					OXI	COA		
Staphylococcus lentus		99.80	0.65	VPT 7	99	1		
Staphylococcus sciuri		0.11	0.02		99	1		
Staphylococcus intermedius		0.10	-.02		1	99		
127771	**Good identification**							**127771**
					OXI			
Staphylococcus lentus		92.96	0.53	VPT 7	99			
Staphylococcus gallinarum		5.40	0.20		1			
Staphylococcus sciuri		1.64	0.15		99			
127773	**Very good identification**							**127773**
					OXI			
Staphylococcus lentus		96.15	0.65	VPT 7	99			
Staphylococcus gallinarum		3.86	0.28		1			
130601	**Good identification**							**130601**
Kocuria kristinae		100.0	0.30	ORN 1				
130641	**Good identification**							**130641**
					OXI			
Kocuria kristinae		95.40	0.49	ORN 1	99			
Staphylococcus lugdunensis		4.60	0.23		1			
130643	**Low discrimination**							**130643**
					OXI			
Staphylococcus lugdunensis		63.45	0.43	NIT 99	1			
Kocuria kristinae		36.55	0.37		99			
130741	**Low discrimination**							**130741**
					OXI			
Kocuria kristinae		87.36	0.39	ORN 1	99			
Staphylococcus lugdunensis		12.64	0.23		1			
130743	**Low discrimination**							**130743**
					OXI			
Staphylococcus lugdunensis		83.89	0.43	NIT 99	1			
Kocuria kristinae		16.11	0.27		99			
131641	**Good identification**							**131641**
Kocuria kristinae		100.0	0.37	ORN 1				
131741	**Low discrimination**							**131741**
					OXI			
Kocuria kristinae		87.76	0.27	ORN 1	99			
Rothia mucilaginosa		12.24	0.05		1			
132243	**Good identification**							**132243**
					PYR			
Staphylococcus lugdunensis		97.27	0.43	TRE 99	96			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		2.73	0.16		1		
132343	**Low discrimination**						**132343**
					PYR		
Staphylococcus lugdunensis		86.26	0.43	TRE 99	96		
Staphylococcus epidermidis		13.74	0.31		1		
132443	**Good identification**						**132443**
Staphylococcus lugdunensis		100.0	0.43	SUC 99			
132543	**Good identification**						**132543**
Staphylococcus lugdunensis		100.0	0.43	SUC 99			
132603	**Good identification**						**132603**
					OXI		
Staphylococcus lugdunensis		99.52	0.43	MLT 99	1		
Kocuria kristinae		0.48	-.06		99		
132641	**Very good identification**						**132641**
					OXI		
Staphylococcus lugdunensis		98.45	0.64	LAC 90	1		
Kocuria kristinae		1.55	0.26		99		
132642	**Good identification**						**132642**
Staphylococcus lugdunensis		96.92	0.43	MNS 99			
Staphylococcus haemolyticus		3.08	0.19				
132643	**Very good identification**						**132643**
					OXI		
Staphylococcus lugdunensis		99.96	0.84		1		
Kocuria kristinae		0.04	0.14		99		
132653	**Good identification**						**132653**
Staphylococcus lugdunensis		100.0	0.43	MAN 1			
132663	**Good identification**						**132663**
Staphylococcus lugdunensis		100.0	0.43	XYL 1			
132703	**Good identification**						**132703**
Staphylococcus lugdunensis		100.0	0.43	MLT 99			
132741	**Very good identification**						**132741**
					OXI		
Staphylococcus lugdunensis		96.90	0.64	LAC 90	1		
Rothia mucilaginosa		2.60	0.27		1		
Kocuria kristinae		0.51	0.16		99		
132742	**Acceptable identification**						**132742**
					OXI	NOV	
Staphylococcus lugdunensis		90.48	0.43	MNS 99	1	1	
Macrococcus caseolyticus		6.29	0.14		99	99	
Staphylococcus haemolyticus		3.24	0.20		1	2	
132743	**Very good identification**						**132743**
					OXI		
Staphylococcus lugdunensis		99.99	0.84		1		
Kocuria kristinae		0.02	0.04		99		
132753	**Good identification**						**132753**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lugdunensis		100.0	0.43	MAN 1			
132763	**Good identification**						**132763**
Staphylococcus lugdunensis		100.0	0.43	XYL 1			
133643	**Good identification**						**133643**
					OXI		
Staphylococcus lugdunensis		98.93	0.43	ESL 1	1		
Kocuria kristinae		1.07	0.01		99		
133700	**Good identification**						**133700**
Rothia mucilaginosa		100.0	0.29	ORN 1			
133701	**Good identification**						**133701**
Rothia mucilaginosa		100.0	0.39	ORN 1			
133713	**Good identification**						**133713**
Staphylococcus lentus		100.0	0.30	VPT 7			
133733	**Good identification**						**133733**
Staphylococcus lentus		100.0	0.30	VPT 7			
133740	**Good identification**						**133740**
Rothia mucilaginosa		100.0	0.37	ORN 1			
133741	**Good identification**						**133741**
					OXI		
Rothia mucilaginosa		95.48	0.47	ORN 1	1		
Staphylococcus lugdunensis		4.00	0.23		1		
Kocuria kristinae		0.52	0.04		99		
133743	**Good identification**						**133743**
Staphylococcus lugdunensis		97.39	0.43	ESL 1			
Rothia mucilaginosa		2.61	0.05				
133753	**Good identification**						**133753**
					OXI	NOV	
Staphylococcus lentus		95.58	0.36	VPT 7	99	99	
Staphylococcus lugdunensis		4.42	0.02	ORN 1	1	1	
133773	**Good identification**						**133773**
Staphylococcus lentus		100.0	0.36	VPT 7			
134450	**Good identification**						**134450**
Staphylococcus kloosii		100.0	0.32	VPT 10			
134452	**Good identification**						**134452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
134550	**Good identification**						**134550**
Staphylococcus kloosii		100.0	0.38	VPT 10			
134552	**Good identification**						**134552**
Staphylococcus kloosii		100.0	0.39	VPT 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
134572	**Good identification**					**134572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
135450	**Good identification**					**135450**
Staphylococcus kloosii		100.0	0.32	VPT 10		
135452	**Good identification**					**135452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
135550	**Good identification**					**135550**
Staphylococcus kloosii		100.0	0.38	VPT 10		
135552	**Good identification**					**135552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
135572	**Good identification**					**135572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
136001	**Good identification**					**136001**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86		
136101	**Good identification**					**136101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86		
136243	**Acceptable identification**					**136243**
					PYR	
Staphylococcus epidermidis		94.04	0.31	URE 92	1	
Staphylococcus lugdunensis		5.96	0.02	ORN 12	96	
136340	**Good identification**					**136340**
Staphylococcus epidermidis		100.0	0.31	URE 92		
136341	**Good identification**					**136341**
Staphylococcus epidermidis		100.0	0.36	URE 92		
136342	**Good identification**					**136342**
Staphylococcus epidermidis		100.0	0.41	URE 92		
136343	**Good identification**					**136343**
					PYR	
Staphylococcus epidermidis		98.89	0.47	URE 92	1	
Staphylococcus lugdunensis		1.11	0.02	ORN 12	96	
136401	**Good identification**					**136401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86		
136501	**Good identification**					**136501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86		
136643	**Good identification**					**136643**
Staphylococcus lugdunensis		100.0	0.43	PHS 1		
136743	**Good identification**					**136743**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR			
Staphylococcus lugdunensis		99.10	0.43	PHS 1	96			
Staphylococcus epidermidis		0.90	0.05		1			
140000	**Good identification**							**140000**
Kocuria rosea		90.94	0.62	NIT 90				
Micrococcus sp.		5.08	0.28					
Dermacoccus nishinomiyaensis		3.07	0.33					
140001	**Low discrimination**							**140001**
					NOV			
Kocuria rosea		82.06	0.35	NIT 90	1			
Staphylococcus cohnii subsp. cohnii		14.37	0.31	MNS 5	99			
Staphylococcus capitis subsp. capitis		1.79	0.04		1			
140010	**Low discrimination**							**140010**
					NOV	YEP	OXI	
Kocuria rosea		89.78	0.42	NIT 90	1	1	35	
Staphylococcus cohnii subsp. cohnii		8.75	0.32	MAN 10	99	12	1	
Macrococcus bovicus		1.01	0.07		99	99	99	
140011	**Low discrimination**							**140011**
					NOV			
Staphylococcus capitis subsp. capitis		46.50	0.27	GLR 1	1			
Staphylococcus cohnii subsp. cohnii		35.70	0.33	NIT 89	99			
Kocuria rosea		17.79	0.16		1			
140020	**Very good identification**							**140020**
					OXI			
Kocuria rosea		99.75	0.54	NIT 90	35			
Micrococcus sp.		0.13	-.13		89			
Kocuria varians		0.12	0.03		13			
140021	**Good identification**							**140021**
Kocuria rosea		100.0	0.28	NIT 90				
140030	**Good identification**							**140030**
Kocuria rosea		100.0	0.34	NIT 90				
140040	**Low discrimination**							**140040**
					NOV	OXI		
Staphylococcus cohnii subsp. cohnii		64.27	0.41	TRE 99	99	1		
Kocuria rosea		24.09	0.21		1	35		
Micrococcus sp.		10.03	0.05		1	89		
140041	**Good identification**							**140041**
					NOV			
Staphylococcus cohnii subsp. cohnii		98.21	0.41	TRE 99	99			
Kocuria rosea		1.79	-.06		1			
140050	**Low discrimination**							**140050**
					NOV	YEP	OXI	
Staphylococcus cohnii subsp. cohnii		89.47	0.43	TRE 99	99	12	1	
Macrococcus bovicus		7.61	0.14		99	99	99	
Kocuria rosea		2.93	0.01		1	1	35	
140051	**Good identification**							**140051**
Staphylococcus cohnii subsp. cohnii		100.0	0.43	TRE 99				
140200	**Acceptable identification**							**140200**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria rosea		93.54	0.42	NIT 90				
Micrococcus sp.		4.65	0.07	SUC 10				
Dermacoccus nishinomiyaensis		1.81	0.08					
140211	**Good identification**							**140211**
Staphylococcus capitis subsp. capitis		97.76	0.32	GLR 1				
Kocuria rosea		2.24	-.04	NIT 89				
140220	**Good identification**							**140220**
Kocuria rosea		100.0	0.34	NIT 90				
140241	**Good identification**							**140241**
Kocuria kristinae		100.0	0.33	GLR 1				
140250	**Low discrimination**							**140250**
					OXI	NOV		
Staphylococcus warneri		76.28	0.32	URE 98	1	1		
Staphylococcus pasteuri		14.16	0.02	TRE 95	1	1		
Macrococcus bovicus		9.56	0.02		99	99		
140400	**Very good identification**							**140400**
					NOV			
Staphylococcus cohnii subsp. cohnii		98.37	0.71		99			
Kocuria rosea		1.18	0.21		1			
Staphylococcus auricularis		0.39	0.18		1			
140401	**Very good identification**							**140401**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		99.86	0.72		99	1		
Kocuria kristinae		0.08	-.02		1	93		
Kocuria rosea		0.06	-.06		1	1		
140402	**Good identification**							**140402**
Staphylococcus cohnii subsp. cohnii		100.0	0.30	LAC 1				
140403	**Acceptable identification**							**140403**
Staphylococcus cohnii subsp. cohnii		91.24	0.31	LAC 1				
Staphylococcus cohnii subsp. urealyticum		8.76	0.11					
140410	**Very good identification**							**140410**
					NOV	YEP	OXI	
Staphylococcus cohnii subsp. cohnii		99.72	0.73		99	12	1	
Macrococcus bovicus		0.17	0.10		99	99	99	
Kocuria rosea		0.10	0.01		1	1	35	
140411	**Very good identification**							**140411**
Staphylococcus cohnii subsp. cohnii		99.92	0.74					
Staphylococcus cohnii subsp. urealyticum		0.08	0.11					
140412	**Good identification**							**140412**
Staphylococcus cohnii subsp. cohnii		100.0	0.32	LAC 1				
140413	**Low discrimination**							**140413**
Staphylococcus cohnii subsp. cohnii		56.73	0.33	LAC 1				
Staphylococcus cohnii subsp. urealyticum		43.27	0.31					

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
140420	**Low discrimination**							**140420**
					NOV			
Staphylococcus cohnii subsp. cohnii		66.28	0.30	XYL 1	99			
Kocuria rosea		33.72	0.13		1			
140421	**Good identification**							**140421**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	XYL 1				
140430	**Good identification**							**140430**
					NOV			
Staphylococcus cohnii subsp. cohnii		95.75	0.32	XYL 1	99			
Kocuria rosea		4.25	-.07		1			
140431	**Good identification**							**140431**
Staphylococcus cohnii subsp. cohnii		100.0	0.33	XYL 1				
140440	**Very good identification**							**140440**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		98.63	0.82		99	1		
Staphylococcus petrasii subsp. pragensis		0.78	0.18		1	99		
Staphylococcus warneri		0.30	0.35		1	8		
140441	**Very good identification**							**140441**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		99.73	0.82		1	99		
Kocuria kristinae		0.23	0.18		99	1		
Staphylococcus auricularis		0.03	0.06		1	1		
140442	**Low discrimination**							**140442**
					NOV			
Staphylococcus cohnii subsp. cohnii		88.40	0.41	LAC 1	99			
Staphylococcus warneri		8.78	0.25		1			
Staphylococcus auricularis		2.82	0.06		1			
140443	**Low discrimination**							**140443**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		81.42	0.41	LAC 1	1	99		
Staphylococcus cohnii subsp. urealyticum		13.99	0.26		1	99		
Kocuria kristinae		4.59	0.05		99	1		
140450	**Very good identification**							**140450**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		99.59	0.84		1	99	12	
Staphylococcus warneri		0.17	0.32		1	1	43	
Macrococcus bovicus		0.13	0.18		99	99	99	
140451	**Very good identification**							**140451**
Staphylococcus cohnii subsp. cohnii		99.86	0.85					
Staphylococcus cohnii subsp. urealyticum		0.14	0.26					
140452	**Low discrimination**							**140452**
					NOV	YEP		
Staphylococcus cohnii subsp. cohnii		83.50	0.43	LAC 1	99	12		
Staphylococcus kloosii		7.22	0.33		99	99		
Staphylococcus warneri		4.72	0.22		1	43		
140453	**Low discrimination**							**140453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		57.00	0.47	URE 99	99	53		
Staphylococcus cohnii subsp. cohnii		41.77	0.43		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		1.23	0.24		1	95		
140460	**Good identification**							**140460**
Staphylococcus cohnii subsp. cohnii		100.0	0.41	XYL 1				
140461	**Good identification**							**140461**
Staphylococcus cohnii subsp. cohnii		100.0	0.41	XYL 1				
140470	**Good identification**							**140470**
					YEP			
Staphylococcus cohnii subsp. cohnii		98.04	0.43	XYL 1	12			
Staphylococcus kloosii		1.96	0.20		99			
140471	**Good identification**							**140471**
Staphylococcus cohnii subsp. cohnii		100.0	0.43	XYL 1				
140500	**Good identification**							**140500**
Staphylococcus cohnii subsp. cohnii		100.0	0.52	GAL 10				
140501	**Good identification**							**140501**
Staphylococcus cohnii subsp. cohnii		100.0	0.52	GAL 10				
140510	**Very good identification**							**140510**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		99.33	0.54	GAL 10	1	12		
Macrococcus bovicus		0.67	0.03		99	99		
140511	**Very good identification**							**140511**
Staphylococcus cohnii subsp. cohnii		100.0	0.55	GAL 10				
140540	**Very good identification**							**140540**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		96.75	0.62	GAL 10	1	99	12	
Macrococcus caseolyticus		2.14	0.14		99	99	99	
Staphylococcus warneri		0.83	0.24		1	1	43	
140541	**Very good identification**							**140541**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		99.32	0.63	GAL 10	1	99		
Kocuria kristinae		0.68	0.08		99	1		
140550	**Very good identification**							**140550**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		97.66	0.64	GAL 10	1	12		
Staphylococcus kloosii		1.38	0.39		1	99		
Macrococcus bovicus		0.49	0.10		99	99		
140551	**Very good identification**							**140551**
Staphylococcus cohnii subsp. cohnii		99.48	0.65	GAL 10				
Staphylococcus cohnii subsp. urealyticum		0.52	0.19					
140552	**Low discrimination**							**140552**
					NOV	YEP		
Staphylococcus kloosii		54.07	0.39	VPT 10	99	99		
Staphylococcus cohnii subsp. cohnii		35.77	0.23	PHS 99	99	12		
Staphylococcus warneri		5.75	0.12		1	43		
140553	**Low discrimination**							**140553**
					NOV	PYR		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		80.62	0.39	URE 99	99	53		
Staphylococcus cohnii subsp. cohnii		15.32	0.24		99	1		
Aerococcus viridans		4.06	0.24		1	95		
140572	**Low discrimination**							**140572**
					NOV	YEP		
Staphylococcus kloosii		50.27	0.27	VPT 10	99	99		
Staphylococcus arlettae		33.35	0.12	PHS 99	99	99		
Aerococcus viridans		16.39	0.16		1	1		
140601	**Low discrimination**							**140601**
					OXI	NOV		
Kocuria kristinae		88.71	0.39	GLR 1	99	1		
Staphylococcus cohnii subsp. cohnii		11.29	0.31	MLT 90	1	99		
140603	**Good identification**							**140603**
Kocuria kristinae		100.0	0.27	GLR 1				
140610	**Low discrimination**							**140610**
					NOV	YEP		
Staphylococcus pasteuri		50.40	0.27	URE 99	1	86		
Staphylococcus cohnii subsp. cohnii		31.39	0.32	MLT 86	99	12		
Staphylococcus warneri		16.85	0.32		1	43		
140611	**Low discrimination**							**140611**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		76.54	0.33	SUC 1	1	99	1	
Kocuria kristinae		19.69	0.11		99	1	93	
Staphylococcus capitis subsp. capitis		3.78	-.03		1	1	1	
140640	**Good identification**							**140640**
					NOV	PYR		
Staphylococcus petrasii subsp. pragensis		90.17	0.59	GLR 1	1	99		
Staphylococcus warneri		6.55	0.61		1	8		
Staphylococcus cohnii subsp. cohnii		1.16	0.41		99	1		
140641	**Good identification**							**140641**
					OXI	NOV	PYR	
Kocuria kristinae		92.31	0.59	GLR 1	99	1	93	
Staphylococcus cohnii subsp. cohnii		4.13	0.41		1	99	1	
Staphylococcus petrasii subsp. pragensis		3.00	0.18		1	1	99	
140642	**Low discrimination**							**140642**
					PYR	YEP		
Staphylococcus warneri		58.99	0.51	URE 98	8	43		
Staphylococcus petrasii subsp. pragensis		24.60	0.18		99	1		
Staphylococcus pasteuri		8.24	0.19		1	86		
140643	**Good identification**							**140643**
					OXI	PYR		
Kocuria kristinae		99.37	0.47	GLR 1	99	93		
Staphylococcus warneri		0.63	0.16		1	8		
140650	**Low discrimination**							**140650**
					NOV	YEP		
Staphylococcus warneri		39.71	0.58	URE 98	1	43		
Staphylococcus pasteuri		38.39	0.43		1	86		
Staphylococcus cohnii subsp. cohnii		12.33	0.43		99	12		
140651	**Low discrimination**							**140651**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		54.92	0.43	SUC 1	1	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		40.15	0.31		99	1	93	
Staphylococcus warneri		3.33	0.23		1	1	8	
140652	**Low discrimination**							**140652**
					NOV			
Staphylococcus pasteuri		49.89	0.35	URE 99	1			
Staphylococcus warneri		42.11	0.48		1			
Staphylococcus saprophyticus subsp. saprophyticus		5.20	0.09		99			
140672	**Acceptable identification**							**140672**
					NOV	PYR	YEP	
Staphylococcus arlettae		94.83	0.37	VPT 1	99	1	99	
Aerococcus viridans		2.67	0.16		1	95	1	
Staphylococcus pasteuri		2.50	-.06		1	1	86	
140673	**Good identification**							**140673**
					NOV	YEP		
Staphylococcus arlettae		95.28	0.40	VPT 1	99	99		
Aerococcus viridans		4.72	0.24		1	1		
140701	**Acceptable identification**							**140701**
					OXI	NOV		
Kocuria kristinae		94.13	0.30	GLR 1	99	1		
Staphylococcus cohnii subsp. cohnii		3.99	0.11	MLT 90	1	99		
Rothia mucilaginosa		1.88	-.09		1	1		
140740	**Low discrimination**							**140740**
					PYR			
Staphylococcus warneri		55.70	0.51	URE 98	8			
Staphylococcus petrasii subsp. pragensis		24.52	0.18		99			
Staphylococcus haemolyticus		7.82	0.26		98			
140741	**Good identification**							**140741**
					OXI	NOV		
Kocuria kristinae		97.60	0.49	GLR 1	99	1		
Staphylococcus cohnii subsp. cohnii		1.46	0.22		1	99		
Rothia mucilaginosa		0.50	-.02		1	1		
140742	**Low discrimination**							**140742**
					OXI	PYR	NOV	
Staphylococcus warneri		59.56	0.41	URE 98	1	8	1	
Macrococcus caseolyticus		24.29	0.14		99	80	99	
Staphylococcus haemolyticus		14.12	0.21		1	98	2	
140743	**Good identification**							**140743**
Kocuria kristinae		100.0	0.37	GLR 1				
140750	**Low discrimination**							**140750**
					NOV	PYR		
Staphylococcus warneri		77.97	0.48	URE 98	1	8		
Staphylococcus haemolyticus		8.88	0.21		2	98		
Staphylococcus cohnii subsp. cohnii		8.52	0.23		99	1		
140752	**Low discrimination**							**140752**
					NOV	PYR		
Staphylococcus warneri		71.35	0.38	URE 98	1	8		
Staphylococcus haemolyticus		13.71	0.16		2	98		
Staphylococcus arlettae		5.86	0.12		99	1		
140772	**Good identification**							**140772**
					NOV	YEP		
Staphylococcus arlettae		99.51	0.53	VPT 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		0.49	0.16		1	1	
140773	**Very good identification**						**140773**
					NOV	YEP	
Staphylococcus arlettae		97.05	0.55	VPT 1	99	99	
Staphylococcus xylosus		2.11	0.12		99	40	
Aerococcus viridans		0.85	0.24		1	1	
141400	**Low discrimination**						**141400**
					OXI		
Staphylococcus cohnii subsp. cohnii		71.90	0.30	ESL 1	1		
Macrococcus carouselicus		28.10	0.12		99		
141401	**Good identification**						**141401**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	ESL 1			
141410	**Low discrimination**						**141410**
					OXI		
Staphylococcus cohnii subsp. cohnii		83.59	0.32	ESL 1	1		
Macrococcus carouselicus		16.41	0.08		99		
141411	**Good identification**						**141411**
Staphylococcus cohnii subsp. cohnii		100.0	0.33	ESL 1			
141433	**Good identification**						**141433**
Aerococcus viridans		100.0	0.31	VPT 1			
141440	**Good identification**						**141440**
Staphylococcus cohnii subsp. cohnii		100.0	0.41	ESL 1			
141441	**Acceptable identification**						**141441**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		94.66	0.41	ESL 1	1	99	
Kocuria kristinae		5.34	0.05		99	1	
141443	**Low discrimination**						**141443**
					NOV		
Aerococcus viridans		62.76	0.34	VPT 1	1		
Staphylococcus cohnii subsp. urealyticum		37.24	0.18		99		
141450	**Low discrimination**						**141450**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		88.86	0.43	ESL 1	1	12	
Staphylococcus kloosii		7.10	0.33		1	99	
Macrococcus equipersicus		1.56	0.02		99	99	
141451	**Acceptable identification**						**141451**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		92.42	0.43	ESL 1	99	1	
Staphylococcus cohnii subsp. urealyticum		4.85	0.18		99	53	
Aerococcus viridans		2.73	0.24		1	95	
141452	**Low discrimination**						**141452**
					NOV	YEP	
Aerococcus viridans		58.74	0.36	VPT 1	1	1	
Staphylococcus kloosii		41.26	0.33		99	99	
141453	**Low discrimination**						**141453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		66.67	0.39	URE 99	99		
Aerococcus viridans		33.33	0.43		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
141463	**Good identification**							**141463**
Aerococcus viridans		100.0	0.34	VPT 1				
141472	**Low discrimination**							**141472**
					NOV	YEP		
Aerococcus viridans		85.06	0.36	VPT 1	1	1		
Staphylococcus kloosii		14.94	0.21		99	99		
141473	**Good identification**							**141473**
Aerococcus viridans		100.0	0.43	VPT 1				
141513	**Low discrimination**							**141513**
					NOV			
Aerococcus viridans		62.30	0.31	VPT 1	1			
Staphylococcus cohnii subsp. urealyticum		37.70	0.15		99			
141533	**Good identification**							**141533**
Aerococcus viridans		100.0	0.31	VPT 1				
141543	**Low discrimination**							**141543**
					NOV			
Aerococcus viridans		79.72	0.34	VPT 1	1			
Staphylococcus cohnii subsp. urealyticum		20.28	0.10		99			
141550	**Low discrimination**							**141550**
					NOV	YEP	PYR	
Staphylococcus kloosii		55.40	0.39	VPT 10	99	99	52	
Staphylococcus cohnii subsp. cohnii		39.71	0.23	PHS 99	99	12	1	
Aerococcus viridans		4.89	0.16		1	1	95	
141552	**Low discrimination**							**141552**
					NOV	YEP		
Staphylococcus kloosii		57.69	0.39	VPT 10	99	99		
Aerococcus viridans		42.31	0.36	PHS 99	1	1		
141553	**Low discrimination**							**141553**
					NOV			
Aerococcus viridans		53.84	0.43	VPT 1	1			
Staphylococcus cohnii subsp. urealyticum		46.16	0.31		99			
141563	**Good identification**							**141563**
Aerococcus viridans		100.0	0.34	VPT 1				
141572	**Low discrimination**							**141572**
					NOV	YEP		
Aerococcus viridans		63.82	0.36	VPT 1	1	1		
Staphylococcus kloosii		21.75	0.27		99	99		
Staphylococcus arlettae		14.43	0.12		99	99		
141573	**Low discrimination**							**141573**
					NOV	YEP		
Aerococcus viridans		88.62	0.43	VPT 1	1	1		
Staphylococcus arlettae		11.38	0.14		99	99		
141601	**Good identification**							**141601**
Kocuria kristinae		100.0	0.27	GLR 1				
141613	**Good identification**							**141613**
Aerococcus viridans		100.0	0.31	VPT 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
141633	**Good identification**							**141633**
Aerococcus viridans		100.0	0.31	VPT 1				
141641	**Good identification**							**141641**
Kocuria kristinae		100.0	0.47	GLR 1				
141643	**Acceptable identification**							**141643**
					OXI			
Kocuria kristinae		92.54	0.34	GLR 1	99			
Aerococcus viridans		7.46	0.34		1			
141652	**Acceptable identification**							**141652**
					PYR	YEP		
Aerococcus viridans		90.55	0.36	VPT 1	95	1		
Staphylococcus pasteuri		9.45	-.06		1	86		
141653	**Low discrimination**							**141653**
					OXI	NOV		
Aerococcus viridans		81.03	0.43	VPT 1	1	1		
Kocuria kristinae		13.95	0.06		99	1		
Staphylococcus cohnii subsp. urealyticum		5.01	0.07		1	99		
141663	**Good identification**							**141663**
Aerococcus viridans		100.0	0.34	VPT 1				
141672	**Low discrimination**							**141672**
					NOV	YEP		
Staphylococcus arlettae		79.80	0.37	VPT 1	99	99		
Aerococcus viridans		20.20	0.36		1	1		
141673	**Low discrimination**							**141673**
					NOV	YEP		
Staphylococcus arlettae		69.18	0.40	VPT 1	99	99		
Aerococcus viridans		30.82	0.43		1	1		
141713	**Good identification**							**141713**
Aerococcus viridans		100.0	0.31	VPT 1				
141733	**Low discrimination**							**141733**
					NOV	YEP		
Aerococcus viridans		66.06	0.31	VPT 1	1	1		
Staphylococcus arlettae		33.94	0.14		99	99		
141741	**Low discrimination**							**141741**
					OXI			
Kocuria kristinae		84.32	0.37	GLR 1	99			
Rothia mucilaginosa		15.68	0.18		1			
141752	**Low discrimination**							**141752**
					NOV	YEP		
Aerococcus viridans		73.48	0.36	VPT 1	1	1		
Staphylococcus arlettae		16.62	0.12		99	99		
Staphylococcus kloosii		9.91	0.19		99	99		
141753	**Low discrimination**							**141753**
					OXI	NOV	YEP	
Aerococcus viridans		84.33	0.43	VPT 1	1	1	1	
Staphylococcus arlettae		10.83	0.14		1	99	99	
Kocuria kristinae		4.84	-.04		99	1	15	
141763	**Low discrimination**							**141763**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Aerococcus viridans		72.18	0.34	VPT 1	1	1	
Staphylococcus arlettae		27.82	0.14		99	99	
141772	**Good identification**						**141772**
					NOV	YEP	
Staphylococcus arlettae		95.72	0.53	VPT 1	99	99	
Aerococcus viridans		4.28	0.36		1	1	
141773	**Good identification**						**141773**
					NOV	YEP	
Staphylococcus arlettae		92.25	0.55	VPT 1	99	99	
Aerococcus viridans		7.25	0.43		1	1	
Staphylococcus xylosus		0.50	-.01		99	40	
142000	**Very good identification**						**142000**
					OXI		
Kocuria rosea		99.64	0.81		35		
Dermacoccus nishinomiyaensis		0.21	0.28		93		
Kocuria varians		0.08	0.27		13		
142001	**Very good identification**						**142001**
					NOV		
Kocuria rosea		97.80	0.55	MNS 5	1		
Staphylococcus capitis subsp. capitis		1.92	0.22		1		
Staphylococcus cohnii subsp. cohnii		0.14	0.08		99		
142002	**Good identification**						**142002**
Kocuria rosea		97.77	0.40	LAC 1			
Kocuria varians		2.23	0.15				
142010	**Very good identification**						**142010**
					NOV		
Kocuria rosea		99.79	0.62	MAN 10	1		
Staphylococcus capitis subsp. capitis		0.13	0.04		1		
Staphylococcus cohnii subsp. cohnii		0.08	0.09		99		
142011	**Low discrimination**						**142011**
					NOV		
Staphylococcus capitis subsp. capitis		69.80	0.45	GLR 1	1		
Kocuria rosea		29.71	0.35		1		
Staphylococcus cohnii subsp. cohnii		0.50	0.10		99		
142020	**Very good identification**						**142020**
Kocuria rosea		99.93	0.74				
Kocuria varians		0.07	0.17				
142021	**Good identification**						**142021**
Kocuria rosea		100.0	0.47	MNS 5			
142022	**Good identification**						**142022**
Kocuria rosea		98.17	0.33	LAC 1			
Kocuria varians		1.84	0.06				
142030	**Very good identification**						**142030**
Kocuria rosea		100.0	0.54	MAN 10			
142031	**Acceptable identification**						**142031**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		94.75	0.28	MAN 10			
Staphylococcus capitis subsp. capitis		5.25	0.04	MNS 5			
142040	**Low discrimination**						**142040**
					NOV		
Kocuria rosea		89.95	0.40	MLT 1	1		
Staphylococcus auricularis		6.00	0.24		1		
Staphylococcus cohnii subsp. cohnii		2.01	0.17		99		
142060	**Good identification**						**142060**
Kocuria rosea		98.57	0.33	MLT 1			
Kocuria varians		1.43	0.04				
142100	**Low discrimination**						**142100**
					OXI		
Kocuria rosea		86.13	0.40	GAL 1	35		
Kocuria varians		10.44	0.30		13		
Dermacoccus nishinomiyaensis		3.43	0.13		93		
142120	**Acceptable identification**						**142120**
Kocuria rosea		90.96	0.33	GAL 1			
Kocuria varians		9.04	0.21				
142200	**Very good identification**						**142200**
					OXI		
Kocuria rosea		99.50	0.62	SUC 10	35		
Staphylococcus auricularis		0.24	0.16		1		
Dermacoccus nishinomiyaensis		0.12	0.03		93		
142201	**Low discrimination**						**142201**
Kocuria rosea		75.28	0.35	SUC 10			
Staphylococcus capitis subsp. capitis		24.72	0.28	MNS 5			
142210	**Good identification**						**142210**
					NOV	OXI	
Kocuria rosea		96.95	0.42	SUC 10	1	35	
Staphylococcus capitis subsp. capitis		2.02	0.10	MAN 10	1	1	
Staphylococcus vitulinus		1.03	0.00		99	99	
142211	**Good identification**						**142211**
Staphylococcus capitis subsp. capitis		97.52	0.51	GLR 1			
Kocuria rosea		2.48	0.16				
142220	**Very good identification**						**142220**
Kocuria rosea		100.0	0.54	SUC 10			
142221	**Good identification**						**142221**
Kocuria rosea		100.0	0.28	SUC 10			
142230	**Acceptable identification**						**142230**
					NOV		
Kocuria rosea		93.71	0.34	SUC 10	1		
Staphylococcus vitulinus		6.29	0.09	MAN 10	99		
142240	**Low discrimination**						**142240**
					PYR		
Staphylococcus haemolyticus		64.30	0.43	ARG 92	98		
Kocuria rosea		14.52	0.21	TRE 93	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus auricularis		8.05	0.23		40				
142242	**Low discrimination**								**142242**
					NOV	PYR			
Staphylococcus haemolyticus		86.48	0.38	ARG 92	2	98			
Staphylococcus hominis subsp. novobiosepticus		5.10	0.03	TRE 93	99	1			
Staphylococcus warneri		3.94	0.17		1	8			
142250	**Low discrimination**								**142250**
					PYR				
Staphylococcus haemolyticus		80.76	0.38	ARG 92	98				
Staphylococcus pasteuri		8.12	0.10	TRE 93	1				
Staphylococcus warneri		7.66	0.24		8				
142252	**Low discrimination**								**142252**
					PYR				
Staphylococcus haemolyticus		88.55	0.33	ARG 92	98				
Staphylococcus pasteuri		6.47	0.02	TRE 93	1				
Staphylococcus warneri		4.98	0.14		8				
142340	**Low discrimination**								**142340**
					OXI	NOV	YEP	PYR	
Staphylococcus haemolyticus		76.24	0.44	ARG 92	1	2	20	98	
Staphylococcus hominis subsp. novobiosepticus		12.48	0.18	TRE 93	1	99	1	1	
Macrococcus caseolyticus		6.60	0.11		99	99	99	80	
142342	**Low discrimination**								**142342**
					OXI	NOV	YEP	PYR	
Staphylococcus haemolyticus		66.75	0.39	ARG 92	1	2	20	98	
Macrococcus caseolyticus		15.41	0.14	TRE 93	99	99	99	80	
Staphylococcus hominis subsp. novobiosepticus		10.46	0.12		1	99	1	1	
142350	**Good identification**								**142350**
					PYR				
Staphylococcus haemolyticus		97.41	0.39	ARG 92	98				
Staphylococcus warneri		2.59	0.14	TRE 93	8				
142352	**Good identification**								**142352**
Staphylococcus haemolyticus		100.0	0.34	ARG 92					
142400	**Low discrimination**								**142400**
					NOV				
Kocuria rosea		49.21	0.40	TRE 1	1				
Staphylococcus cohnii subsp. cohnii		34.32	0.48		99				
Staphylococcus auricularis		16.17	0.38		1				
142401	**Low discrimination**								**142401**
					NOV				
Staphylococcus cohnii subsp. cohnii		88.83	0.49	NIT 7	99				
Kocuria rosea		6.19	0.14		1				
Staphylococcus auricularis		4.29	0.18		1				
142410	**Low discrimination**								**142410**
					NOV				
Staphylococcus cohnii subsp. cohnii		88.88	0.50	NIT 7	99				
Kocuria rosea		11.12	0.21		1				
142411	**Good identification**								**142411**
					NOV				
Staphylococcus cohnii subsp. cohnii		96.62	0.51	NIT 7	99				
Staphylococcus capitis subsp. capitis		2.79	0.11		1				
Kocuria rosea		0.59	-.06		1				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
142420	**Good identification**							**142420**
					NOV			
Kocuria rosea		98.38	0.33	TRE 1	1			
Staphylococcus cohnii subsp. cohnii		1.62	0.07		99			
142440	**Low discrimination**							**142440**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		69.29	0.59	NIT 7	1	99	12	
Staphylococcus auricularis		24.06	0.46		1	1	1	
Macrococcus caseolyticus		2.26	0.14		99	99	99	
142441	**Very good identification**							**142441**
					OXI	NOV		
Staphylococcus cohnii subsp. cohnii		96.35	0.59	NIT 7	1	99		
Staphylococcus auricularis		3.43	0.26		1	1		
Kocuria kristinae		0.22	-.05		99	1		
142450	**Very good identification**							**142450**
					PYR	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		95.95	0.61	NIT 7	1	99	12	
Staphylococcus haemolyticus		1.32	0.20		98	2	20	
Staphylococcus pasteuri		0.99	0.10		1	1	86	
142451	**Very good identification**							**142451**
Staphylococcus cohnii subsp. cohnii		99.68	0.61	NIT 7				
Staphylococcus cohnii subsp. urealyticum		0.32	0.11					
142453	**Low discrimination**							**142453**
Staphylococcus cohnii subsp. urealyticum		76.19	0.31	URE 99				
Staphylococcus cohnii subsp. cohnii		23.81	0.20					
142500	**Low discrimination**							**142500**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		37.75	0.29	NIT 7	1	99	12	
Macrococcus caseolyticus		35.10	0.14	GAL 10	99	99	99	
Staphylococcus auricularis		17.79	0.18		1	1	1	
142501	**Low discrimination**							**142501**
					NOV			
Staphylococcus cohnii subsp. cohnii		88.29	0.29	NIT 7	99			
Staphylococcus schleiferi subsp. schleiferi		6.15	-.02	GAL 10	1			
Rothia mucilaginosa		5.56	-.09		1			
142510	**Good identification**							**142510**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	NIT 7				
142511	**Good identification**							**142511**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	NIT 7				
142540	**Good identification**							**142540**
					OXI	NOV	YEP	
Macrococcus caseolyticus		94.48	0.55	GLR 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		3.25	0.39		1	99	12	
Staphylococcus auricularis		1.13	0.26		1	1	1	
142541	**Low discrimination**							**142541**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		73.91	0.40	NIT 7	1	99	12	
Macrococcus caseolyticus		20.03	0.14	GAL 10	99	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Rothia mucilaginosa		3.43	-.02		1	1	1		
142542	**Very good identification**								**142542**
					OXI	NOV			
Micrococcus caseolyticus		99.52	0.59	GLR 1	99	99			
Staphylococcus haemolyticus		0.39	0.21		1	2			
Staphylococcus auricularis		0.09	0.06		1	1			
142550	**Low discrimination**								**142550**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		65.32	0.41	NIT 7	1	99	1	12	
Micrococcus caseolyticus		15.07	0.14	GAL 10	99	99	80	99	
Staphylococcus haemolyticus		9.14	0.21		1	2	98	20	
142551	**Good identification**								**142551**
					PYR				
Staphylococcus cohnii subsp. cohnii		97.03	0.42	NIT 7	1				
Staphylococcus saprophyticus subsp. bovis		2.97	0.06	GAL 10	99				
142600	**Low discrimination**								**142600**
					YEP				
Staphylococcus auricularis		48.34	0.37	VPT 12	1				
Kocuria rosea		17.71	0.21	GLR 3	1				
Staphylococcus pasteuri		12.15	0.19		86				
142610	**Low discrimination**								**142610**
					PYR				
Staphylococcus pasteuri		83.17	0.35	URE 99	1				
Staphylococcus haemolyticus		6.88	0.20	MLT 86	98				
Staphylococcus warneri		4.87	0.24		8				
142612	**Low discrimination**								**142612**
					PYR				
Staphylococcus pasteuri		86.08	0.27	URE 99	1				
Staphylococcus haemolyticus		9.81	0.15	MLT 86	98				
Staphylococcus warneri		4.11	0.14		8				
142640	**Low discrimination**								**142640**
					PYR				
Staphylococcus haemolyticus		56.78	0.66	ARG 92	98				
Staphylococcus petrasii subsp. pragensis		22.30	0.39		99				
Staphylococcus warneri		6.25	0.53		8				
142641	**Low discrimination**								**142641**
					OXI				
Kocuria kristinae		59.50	0.36	GLR 1	99				
Staphylococcus haemolyticus		22.29	0.35	NIT 7	1				
Staphylococcus auricularis		8.67	0.25		1				
142642	**Low discrimination**								**142642**
					PYR				
Staphylococcus haemolyticus		84.72	0.61	ARG 92	98				
Staphylococcus warneri		5.53	0.43		8				
Staphylococcus pasteuri		4.41	0.27		1				
142650	**Low discrimination**								**142650**
					PYR				
Staphylococcus haemolyticus		52.30	0.61	ARG 92	98				
Staphylococcus pasteuri		39.20	0.51		1				
Staphylococcus warneri		7.10	0.50		8				
142651	**Low discrimination**								**142651**
					PYR	NOV	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus haemolyticus		53.47	0.30	ARG 92	98	2	20	
Staphylococcus cohnii subsp. cohnii		13.86	0.20	MNS 3	1	99	12	
Staphylococcus pasteuri		13.09	0.10		1	1	86	
142652	**Low discrimination**							**142652**
					PYR			
Staphylococcus haemolyticus		61.37	0.56	ARG 92	98			
Staphylococcus pasteuri		33.40	0.43		1			
Staphylococcus warneri		4.94	0.41		8			
142701	**Good identification**							**142701**
					OXI			
Rothia mucilaginosa		96.32	0.32	GLR 1	1			
Kocuria kristinae		3.68	0.06	ESL 90	99			
142740	**Low discrimination**							**142740**
					OXI	NOV		
Staphylococcus haemolyticus		57.19	0.67	ARG 92	1	2		
Macrococcus caseolyticus		36.91	0.52		99	99		
Rothia mucilaginosa		3.16	0.30		1	1		
142741	**Low discrimination**							**142741**
					OXI			
Rothia mucilaginosa		72.06	0.39	GLR 1	1			
Staphylococcus haemolyticus		13.44	0.36	ESL 90	1			
Kocuria kristinae		10.61	0.26		99			
142742	**Low discrimination**							**142742**
					OXI	PYR	NOV	
Macrococcus caseolyticus		62.64	0.55	GLR 1	99	80	99	
Staphylococcus haemolyticus		36.40	0.62		1	98	2	
Staphylococcus warneri		0.67	0.33		1	8	1	
142743	**Low discrimination**							**142743**
					OXI	NOV		
Staphylococcus haemolyticus		37.46	0.31	ARG 92	1	2		
Staphylococcus lugdunensis		24.74	0.21	MNS 3	1	1		
Macrococcus caseolyticus		21.05	0.14		99	99		
142750	**Good identification**							**142750**
					NOV	PYR		
Staphylococcus haemolyticus		91.40	0.62	ARG 92	2	98		
Staphylococcus saprophyticus subsp. bovis		3.50	0.31		99	99		
Staphylococcus warneri		3.47	0.40		1	8		
142751	**Low discrimination**							**142751**
					NOV			
Staphylococcus haemolyticus		66.11	0.31	ARG 92	2			
Staphylococcus saprophyticus subsp. bovis		20.44	0.18	MNS 3	99			
Rothia mucilaginosa		6.10	-.02		1			
142752	**Good identification**							**142752**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		93.11	0.57	ARG 92	1	98	2	
Macrococcus caseolyticus		2.76	0.14		99	80	99	
Staphylococcus warneri		2.10	0.30		1	8	1	
142773	**Acceptable identification**							**142773**
Staphylococcus xylosus		94.56	0.30	VPT 14				
Staphylococcus arlettae		5.44	0.14	URE 99				
143000	**Good identification**							**143000**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria rosea		100.0	0.40	ESL 1				
143020	**Good identification**							**143020**
Kocuria rosea		100.0	0.33	ESL 1				
143651	**Good identification**							**143651**
					NOV	PYR		
Staphylococcus sciuri		96.98	0.32	VPT 1	99	1		
Kocuria kristinae		3.02	-.05	GLR 14	1	93		
143671	**Good identification**							**143671**
Staphylococcus sciuri		100.0	0.30	VPT 1				
143700	**Good identification**							**143700**
Rothia mucilaginosa		100.0	0.42	GLR 1				
143701	**Good identification**							**143701**
					OXI			
Rothia mucilaginosa		99.89	0.51	GLR 1	1			
Kocuria kristinae		0.11	-.06		99			
143713	**Low discrimination**							**143713**
Staphylococcus lentus		88.04	0.30	VPT 7				
Staphylococcus sciuri		11.96	0.10	GLR 1				
143733	**Acceptable identification**							**143733**
Staphylococcus lentus		90.71	0.30	VPT 7				
Staphylococcus sciuri		9.29	0.07	GLR 1				
143740	**Good identification**							**143740**
					OXI	NOV		
Rothia mucilaginosa		96.77	0.49	GLR 1	1	1		
Staphylococcus haemolyticus		1.96	0.26		1	2		
Micrococcus caseolyticus		1.27	0.11		99	99		
143741	**Very good identification**							**143741**
					OXI	NOV	PYR	
Rothia mucilaginosa		99.50	0.59	GLR 1	1	1	75	
Kocuria kristinae		0.41	0.14		99	1	93	
Staphylococcus sciuri		0.10	0.07		99	99	1	
143751	**Low discrimination**							**143751**
					OXI	NOV		
Staphylococcus sciuri		66.67	0.38	VPT 1	99	99		
Rothia mucilaginosa		21.17	0.18	GLR 14	1	1		
Staphylococcus lentus		12.16	0.25		99	99		
143753	**Low discrimination**							**143753**
					OXI			
Staphylococcus lentus		78.44	0.36	VPT 7	99			
Staphylococcus sciuri		19.74	0.22	GLR 1	99			
Staphylococcus xylosus		1.82	-.04		1			
143771	**Low discrimination**							**143771**
					OXI			
Staphylococcus sciuri		72.84	0.35	VPT 1	99			
Staphylococcus lentus		17.62	0.25	GLR 14	99			
Staphylococcus xylosus		5.87	0.09		1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
143773	**Low discrimination**					**143773**
					OXI	
Staphylococcus lentus		58.92	0.36	VPT 7	99	
Staphylococcus xylosus		15.76	0.18	GLR 1	1	
Staphylococcus sciuri		11.19	0.19		99	
144052	**Good identification**					**144052**
Staphylococcus kloosii		100.0	0.33	VPT 10		
144150	**Good identification**					**144150**
Staphylococcus kloosii		100.0	0.39	VPT 10		
144152	**Good identification**					**144152**
Staphylococcus kloosii		100.0	0.39	VPT 10		
144172	**Good identification**					**144172**
Staphylococcus kloosii		100.0	0.27	VPT 10		
144400	**Good identification**					**144400**
Staphylococcus cohnii subsp. cohnii		100.0	0.51	PHS 9		
144401	**Acceptable identification**					**144401**
					NOV	
Staphylococcus cohnii subsp. cohnii		93.92	0.51	PHS 9	99	
Staphylococcus schleiferi subsp. schleiferi		6.09	0.19		1	
144410	**Low discrimination**					**144410**
					YEP	
Staphylococcus cohnii subsp. cohnii		88.14	0.53	PHS 9	12	
Staphylococcus kloosii		11.86	0.47		99	
144411	**Very good identification**					**144411**
Staphylococcus cohnii subsp. cohnii		98.86	0.54	PHS 9		
Staphylococcus cohnii subsp. urealyticum		1.14	0.14			
144412	**Acceptable identification**					**144412**
					YEP	
Staphylococcus kloosii		93.52	0.48	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		6.48	0.12	MLT 95	12	
144413	**Acceptable identification**					**144413**
Staphylococcus cohnii subsp. urealyticum		92.04	0.35	URE 99		
Staphylococcus cohnii subsp. cohnii		7.96	0.12			
144430	**Low discrimination**					**144430**
					YEP	
Staphylococcus kloosii		76.91	0.35	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		23.09	0.12	MLT 95	12	
144432	**Good identification**					**144432**
Staphylococcus kloosii		100.0	0.36	VPT 10		
144440	**Very good identification**					**144440**
					YEP	
Staphylococcus cohnii subsp. cohnii		98.97	0.61	PHS 9	12	
Staphylococcus kloosii		1.03	0.33		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
144441	**Very good identification**					**144441**
	Staphylococcus cohnii subsp. cohnii	99.74	0.62	PHS 9		
	Staphylococcus cohnii subsp. urealyticum	0.26	0.09			
144442	**Low discrimination**					**144442**
					YEP	
	Staphylococcus kloosii	52.68	0.33	VPT 10	99	
	Staphylococcus cohnii subsp. cohnii	47.32	0.20	MAN 99	12	
144443	**Low discrimination**					**144443**
	Staphylococcus cohnii subsp. urealyticum	72.26	0.30	URE 99		
	Staphylococcus cohnii subsp. cohnii	27.74	0.21	MAN 91		
144450	**Low discrimination**					**144450**
					YEP	
	Staphylococcus cohnii subsp. cohnii	55.32	0.63	PHS 9	12	
	Staphylococcus kloosii	44.68	0.74		99	
144451	**Very good identification**					**144451**
					YEP	
	Staphylococcus cohnii subsp. cohnii	97.26	0.64	PHS 9	12	
	Staphylococcus cohnii subsp. urealyticum	2.01	0.30		64	
	Staphylococcus kloosii	0.73	0.33		99	
144452	**Very good identification**					**144452**
					YEP	
	Staphylococcus kloosii	98.60	0.74	VPT 10	99	
	Staphylococcus cohnii subsp. cohnii	1.14	0.22		12	
	Staphylococcus cohnii subsp. urealyticum	0.26	0.10		64	
144453	**Acceptable identification**					**144453**
					YEP	
	Staphylococcus cohnii subsp. urealyticum	91.97	0.51	URE 99	64	
	Staphylococcus cohnii subsp. cohnii	4.44	0.23		12	
	Staphylococcus kloosii	3.59	0.33		99	
144470	**Very good identification**					**144470**
					YEP	
	Staphylococcus kloosii	95.24	0.61	VPT 10	99	
	Staphylococcus cohnii subsp. cohnii	4.77	0.22		12	
144472	**Very good identification**					**144472**
	Staphylococcus kloosii	100.0	0.62	VPT 10		
144500	**Good identification**					**144500**
	Staphylococcus cohnii subsp. cohnii	100.0	0.31	PHS 9		
144501	**Low discrimination**					**144501**
					NOV	
	Staphylococcus cohnii subsp. cohnii	63.17	0.32	PHS 9	99	
	Staphylococcus schleiferi subsp. schleiferi	36.83	0.19	GAL 10	1	
144510	**Low discrimination**					**144510**
					YEP	
	Staphylococcus kloosii	70.16	0.53	VPT 10	99	
	Staphylococcus cohnii subsp. cohnii	29.84	0.33	MLT 95	12	
144511	**Good identification**					**144511**
	Staphylococcus cohnii subsp. cohnii	95.73	0.34	PHS 9		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		4.27	0.07	GAL 10		
144512	**Very good identification**					**144512**
Staphylococcus kloosii		100.0	0.54	VPT 10		
144513	**Good identification**					**144513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	URE 99		
144530	**Good identification**					**144530**
Staphylococcus kloosii		100.0	0.41	VPT 10		
144532	**Good identification**					**144532**
Staphylococcus kloosii		100.0	0.42	VPT 10		
144540	**Low discrimination**					**144540**
					YEP	
Staphylococcus cohnii subsp. cohnii		84.65	0.41	PHS 9	12	
Staphylococcus kloosii		15.35	0.39	GAL 10	99	
144541	**Good identification**					**144541**
Staphylococcus cohnii subsp. cohnii		100.0	0.42	PHS 9		
144542	**Good identification**					**144542**
Staphylococcus kloosii		100.0	0.39	VPT 10		
144550	**Good identification**					**144550**
					YEP	
Staphylococcus kloosii		93.38	0.80	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		6.62	0.43		12	
144551	**Low discrimination**					**144551**
					YEP	
Staphylococcus cohnii subsp. cohnii		82.56	0.44	PHS 9	12	
Staphylococcus kloosii		10.86	0.39	GAL 10	99	
Staphylococcus cohnii subsp. urealyticum		6.58	0.22		64	
144552	**Very good identification**					**144552**
Staphylococcus kloosii		100.0	0.80	VPT 10		
144553	**Low discrimination**					**144553**
Staphylococcus cohnii subsp. urealyticum		84.98	0.43	URE 99		
Staphylococcus kloosii		15.02	0.39			
144562	**Good identification**					**144562**
Staphylococcus kloosii		100.0	0.27	VPT 10		
144570	**Very good identification**					**144570**
Staphylococcus kloosii		100.0	0.67	VPT 10		
144572	**Very good identification**					**144572**
Staphylococcus kloosii		99.01	0.68	VPT 10		
Staphylococcus arlettae		1.00	0.15			
144610	**Low discrimination**					**144610**
					YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		56.86	0.27	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		43.15	0.12	SUC 9	12		
144612	**Good identification**						**144612**
Staphylococcus kloosii		100.0	0.27	VPT 10			
144650	**Low discrimination**						**144650**
					NOV	YEP	
Staphylococcus kloosii		82.76	0.53	VPT 10	99	99	
Staphylococcus cohnii subsp. cohnii		10.47	0.22	SUC 9	99	12	
Staphylococcus warneri		3.44	0.17		1	43	
144652	**Very good identification**						**144652**
					NOV		
Staphylococcus kloosii		98.51	0.54	VPT 10	99		
Staphylococcus pasteuri		1.49	-.06	SUC 9	1		
144670	**Good identification**						**144670**
Staphylococcus kloosii		100.0	0.41	VPT 10			
144672	**Low discrimination**						**144672**
Staphylococcus arlettae		77.51	0.41	VPT 1			
Staphylococcus kloosii		22.49	0.41				
144673	**Good identification**						**144673**
Staphylococcus arlettae		100.0	0.43	VPT 1			
144710	**Good identification**						**144710**
Staphylococcus kloosii		100.0	0.33	VPT 10			
144712	**Good identification**						**144712**
Staphylococcus kloosii		100.0	0.33	VPT 10			
144750	**Very good identification**						**144750**
Staphylococcus kloosii		100.0	0.59	VPT 10			
144752	**Very good identification**						**144752**
Staphylococcus kloosii		97.52	0.60	VPT 10			
Staphylococcus arlettae		2.48	0.15	SUC 9			
144770	**Acceptable identification**						**144770**
Staphylococcus kloosii		90.08	0.47	VPT 10			
Staphylococcus arlettae		9.92	0.15	SUC 9			
144772	**Good identification**						**144772**
Staphylococcus arlettae		90.96	0.56	VPT 1			
Staphylococcus kloosii		9.04	0.47				
144773	**Very good identification**						**144773**
Staphylococcus arlettae		99.59	0.59	VPT 1			
Staphylococcus xylosus		0.41	0.00				
145050	**Good identification**						**145050**
Staphylococcus kloosii		100.0	0.33	VPT 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
145052	**Good identification**					**145052**
Staphylococcus kloosii		100.0	0.33	VPT 10		
145150	**Good identification**					**145150**
Staphylococcus kloosii		100.0	0.39	VPT 10		
145152	**Good identification**					**145152**
Staphylococcus kloosii		100.0	0.39	VPT 10		
145172	**Good identification**					**145172**
Staphylococcus kloosii		100.0	0.27	VPT 10		
145410	**Acceptable identification**					**145410**
					YEP	
Staphylococcus kloosii		93.02	0.47	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		6.98	0.12	MLT 95	12	
145412	**Good identification**					**145412**
Staphylococcus kloosii		100.0	0.48	VPT 10		
145413	**Good identification**					**145413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	URE 99		
145430	**Good identification**					**145430**
Staphylococcus kloosii		100.0	0.35	VPT 10		
145432	**Good identification**					**145432**
Staphylococcus kloosii		100.0	0.36	VPT 10		
145440	**Low discrimination**					**145440**
					YEP	
Staphylococcus kloosii		50.68	0.33	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		49.32	0.20	MAN 99	12	
145442	**Good identification**					**145442**
Staphylococcus kloosii		100.0	0.33	VPT 10		
145450	**Very good identification**					**145450**
					YEP	
Staphylococcus kloosii		98.77	0.74	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		1.24	0.22		12	
145452	**Very good identification**					**145452**
Staphylococcus kloosii		100.0	0.74	VPT 10		
145453	**Acceptable identification**					**145453**
Staphylococcus cohnii subsp. urealyticum		90.88	0.42	URE 99		
Staphylococcus kloosii		9.12	0.33			
145470	**Very good identification**					**145470**
Staphylococcus kloosii		100.0	0.61	VPT 10		
145472	**Very good identification**					**145472**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.62	VPT 10		
145510	**Very good identification**					**145510**
Staphylococcus kloosii		100.0	0.53	VPT 10		
145512	**Very good identification**					**145512**
Staphylococcus kloosii		100.0	0.54	VPT 10		
145530	**Good identification**					**145530**
Staphylococcus kloosii		100.0	0.41	VPT 10		
145532	**Good identification**					**145532**
Staphylococcus kloosii		100.0	0.42	VPT 10		
145540	**Good identification**					**145540**
Staphylococcus kloosii		100.0	0.39	VPT 10		
145542	**Good identification**					**145542**
Staphylococcus kloosii		100.0	0.39	VPT 10		
145550	**Very good identification**					**145550**
Staphylococcus kloosii		100.0	0.80	VPT 10		
145551	**Low discrimination**					**145551**
Staphylococcus kloosii		80.92	0.39	VPT 10		
Staphylococcus cohnii subsp. urealyticum		19.08	0.14	MNS 1		
145552	**Very good identification**					**145552**
Staphylococcus kloosii		100.0	0.80	VPT 10		
145553	**Low discrimination**					**145553**
Staphylococcus cohnii subsp. urealyticum		68.76	0.35	URE 99		
Staphylococcus kloosii		31.24	0.39			
145562	**Good identification**					**145562**
Staphylococcus kloosii		100.0	0.27	VPT 10		
145570	**Very good identification**					**145570**
Staphylococcus kloosii		100.0	0.67	VPT 10		
145572	**Very good identification**					**145572**
Staphylococcus kloosii		99.01	0.68	VPT 10		
Staphylococcus arlettae		1.00	0.15			
145610	**Good identification**					**145610**
Staphylococcus kloosii		100.0	0.27	VPT 10		
145612	**Good identification**					**145612**
Staphylococcus kloosii		100.0	0.27	VPT 10		
145650	**Very good identification**					**145650**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.53	VPT 10		
145652	**Very good identification**					**145652**
Staphylococcus kloosii		100.0	0.54	VPT 10		
145670	**Good identification**					**145670**
Staphylococcus kloosii		100.0	0.41	VPT 10		
145672	**Low discrimination**					**145672**
Staphylococcus arlettae		77.51	0.41	VPT 1		
Staphylococcus kloosii		22.49	0.41			
145673	**Good identification**					**145673**
Staphylococcus arlettae		100.0	0.43	VPT 1		
145710	**Good identification**					**145710**
Staphylococcus kloosii		100.0	0.33	VPT 10		
145712	**Good identification**					**145712**
Staphylococcus kloosii		100.0	0.33	VPT 10		
145750	**Very good identification**					**145750**
Staphylococcus kloosii		100.0	0.59	VPT 10		
145752	**Very good identification**					**145752**
Staphylococcus kloosii		97.52	0.60	VPT 10		
Staphylococcus arlettae		2.48	0.15	SUC 9		
145770	**Acceptable identification**					**145770**
Staphylococcus kloosii		90.08	0.47	VPT 10		
Staphylococcus arlettae		9.92	0.15	SUC 9		
145772	**Good identification**					**145772**
Staphylococcus arlettae		90.96	0.56	VPT 1		
Staphylococcus kloosii		9.04	0.47			
145773	**Very good identification**					**145773**
Staphylococcus arlettae		99.71	0.59	VPT 1		
Staphylococcus gallinarum		0.29	-.04			
146000	**Good identification**					**146000**
Kocuria rosea		98.91	0.40	PHS 1		
Dermacoccus nishinomiyaensis		1.09	0.01			
146001	**Low discrimination**					**146001**
Staphylococcus schleiferi subsp. schleiferi		74.72	0.29	ARG 86		
Kocuria rosea		20.38	0.14	GLR 1		
Staphylococcus capitis subsp. capitis		4.90	0.04			
146011	**Good identification**					**146011**
Staphylococcus capitis subsp. capitis		96.64	0.27	GLR 1		
Kocuria rosea		3.36	-.06	PHS 11		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
146020	**Good identification**					**146020**
Kocuria rosea		100.0	0.33	PHS 1		
146101	**Good identification**					**146101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.29	ARG 86		
146211	**Good identification**					**146211**
Staphylococcus capitis subsp. capitis		100.0	0.32	GLR 1		
146342	**Good identification**					**146342**
Staphylococcus epidermidis		100.0	0.30	URE 92		
146343	**Good identification**					**146343**
Staphylococcus epidermidis		100.0	0.36	URE 92		
146400	**Low discrimination**					**146400**
					NOV	
Staphylococcus cohnii subsp. cohnii		81.22	0.28	NIT 7	99	
Kocuria rosea		11.89	-.01	PHS 9	1	
Staphylococcus schleiferi subsp. schleiferi		6.89	-.02		1	
146401	**Low discrimination**					**146401**
					NOV	
Staphylococcus schleiferi subsp. schleiferi		88.57	0.39	ARG 86	1	
Staphylococcus cohnii subsp. cohnii		11.43	0.28	GLR 1	99	
146410	**Good identification**					**146410**
Staphylococcus cohnii subsp. cohnii		100.0	0.30	NIT 7		
146411	**Acceptable identification**					**146411**
					NOV	
Staphylococcus cohnii subsp. cohnii		94.21	0.30	NIT 7	99	
Staphylococcus schleiferi subsp. schleiferi		5.79	-.02	PHS 9	1	
146440	**Good identification**					**146440**
					NOV	
Staphylococcus cohnii subsp. cohnii		96.58	0.38	NIT 7	99	
Staphylococcus auricularis		3.42	0.05	PHS 9	1	
146441	**Good identification**					**146441**
					NOV	
Staphylococcus cohnii subsp. cohnii		97.59	0.39	NIT 7	99	
Staphylococcus schleiferi subsp. schleiferi		2.41	-.02	PHS 9	1	
146450	**Acceptable identification**					**146450**
					YEP	
Staphylococcus cohnii subsp. cohnii		90.22	0.40	NIT 7	12	
Staphylococcus kloosii		9.78	0.33	PHS 9	99	
146451	**Good identification**					**146451**
Staphylococcus cohnii subsp. cohnii		95.38	0.41	NIT 7		
Staphylococcus cohnii subsp. urealyticum		4.62	0.14	PHS 9		
146452	**Good identification**					**146452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
146453	**Good identification**					**146453**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	URE 99			
146501	**Good identification**						**146501**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		98.59	0.39	ARG 86	1		
Staphylococcus cohnii subsp. cohnii		1.41	0.09	GLR 1	99		
146550	**Low discrimination**						**146550**
					PYR	YEP	
Staphylococcus kloosii		58.65	0.39	VPT 10	52	99	
Staphylococcus cohnii subsp. cohnii		30.98	0.20	NIT 1	1	12	
Staphylococcus saprophyticus subsp. bovis		10.37	0.06		99	80	
146552	**Good identification**						**146552**
Staphylococcus kloosii		100.0	0.39	VPT 10			
146553	**Low discrimination**						**146553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		81.02	0.28	URE 99	99		
Staphylococcus caprae		18.98	-.02		1		
146572	**Good identification**						**146572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
146620	**Good identification**						**146620**
Staphylococcus muscae		100.0	0.39	VPT 1			
146703	**Good identification**						**146703**
Staphylococcus hyicus		100.0	0.29	VPT 1			
147450	**Good identification**						**147450**
Staphylococcus kloosii		100.0	0.33	VPT 10			
147452	**Good identification**						**147452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
147453	**Good identification**						**147453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	URE 99			
147550	**Good identification**						**147550**
Staphylococcus kloosii		100.0	0.39	VPT 10			
147552	**Good identification**						**147552**
Staphylococcus kloosii		100.0	0.39	VPT 10			
147572	**Good identification**						**147572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
147651	**Good identification**						**147651**
Staphylococcus sciuri		100.0	0.37	VPT 1			
147671	**Good identification**						**147671**
Staphylococcus sciuri		100.0	0.34	VPT 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
147711	**Good identification**					**147711**
Staphylococcus sciuri		95.27	0.31	VPT 1		
Staphylococcus lentus		4.73	0.06	GLR 14		
147731	**Acceptable identification**					**147731**
Staphylococcus sciuri		93.83	0.28	VPT 1		
Staphylococcus lentus		6.17	0.06	GLR 14		
147751	**Good identification**					**147751**
Staphylococcus sciuri		97.39	0.43	VPT 1		
Staphylococcus lentus		2.61	0.12	GLR 14		
147753	**Low discrimination**					**147753**
					OXI	
Staphylococcus sciuri		60.74	0.27	VPT 1	99	
Staphylococcus lentus		35.44	0.23	GLR 14	99	
Staphylococcus gallinarum		3.82	-.04	LAC 14	1	
147771	**Low discrimination**					**147771**
					OXI	
Staphylococcus sciuri		60.87	0.40	VPT 1	99	
Staphylococcus gallinarum		33.44	0.29	GLR 14	1	
Staphylococcus equorum		2.71	0.09		1	
147773	**Low discrimination**					**147773**
					OXI	
Staphylococcus gallinarum		73.40	0.37	VPT 1	1	
Staphylococcus sciuri		8.88	0.24	URE 92	99	
Staphylococcus lentus		6.87	0.23		99	
150400	**Good identification**					**150400**
Staphylococcus cohnii subsp. cohnii		100.0	0.30	ORN 1		
150401	**Good identification**					**150401**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	ORN 1		
150410	**Good identification**					**150410**
Staphylococcus cohnii subsp. cohnii		100.0	0.32	ORN 1		
150411	**Good identification**					**150411**
Staphylococcus cohnii subsp. cohnii		100.0	0.33	ORN 1		
150440	**Good identification**					**150440**
Staphylococcus cohnii subsp. cohnii		100.0	0.41	ORN 1		
150441	**Good identification**					**150441**
Staphylococcus cohnii subsp. cohnii		100.0	0.41	ORN 1		
150450	**Good identification**					**150450**
Staphylococcus cohnii subsp. cohnii		100.0	0.43	ORN 1		
150451	**Good identification**					**150451**
Staphylococcus cohnii subsp. cohnii		100.0	0.43	ORN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
152000	**Good identification**					**152000**
Kocuria rosea		100.0	0.40	ORN 1		
152020	**Good identification**					**152020**
Kocuria rosea		100.0	0.33	ORN 1		
152641	**Good identification**					**152641**
					OXI	
Staphylococcus lugdunensis		99.42	0.42	GLR 2	1	
Kocuria kristinae		0.58	-.05	LAC 90	99	
152643	**Very good identification**					**152643**
Staphylococcus lugdunensis		100.0	0.62	GLR 2		
152741	**Good identification**					**152741**
Staphylococcus lugdunensis		98.69	0.42	GLR 2		
Rothia mucilaginosa		1.31	-.02	LAC 90		
152743	**Very good identification**					**152743**
Staphylococcus lugdunensis		100.0	0.62	GLR 2		
154452	**Good identification**					**154452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
154550	**Good identification**					**154550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
154552	**Good identification**					**154552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
154572	**Good identification**					**154572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
155450	**Good identification**					**155450**
Staphylococcus kloosii		100.0	0.33	VPT 10		
155452	**Good identification**					**155452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
155550	**Good identification**					**155550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
155552	**Good identification**					**155552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
155572	**Good identification**					**155572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
160400	**Acceptable identification**					**160400**
					NOV	
Staphylococcus cohnii subsp. cohnii		94.03	0.30	BGA 1	99	
Staphylococcus auricularis		5.97	0.02		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
160401	**Good identification**						**160401**
Staphylococcus cohnii subsp. cohnii		100.0	0.31	BGA 1			
160410	**Good identification**						**160410**
Staphylococcus cohnii subsp. cohnii		100.0	0.32	BGA 1			
160411	**Low discrimination**						**160411**
Staphylococcus cohnii subsp. cohnii		89.84	0.33	BGA 1			
Staphylococcus cohnii subsp. urealyticum		10.16	0.14				
160413	**Good identification**						**160413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	URE 99			
160440	**Low discrimination**						**160440**
					NOV		
Staphylococcus cohnii subsp. cohnii		89.18	0.41	BGA 1	99		
Staphylococcus warneri		6.65	0.22		1		
Staphylococcus auricularis		4.17	0.10		1		
160441	**Acceptable identification**						**160441**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		90.86	0.41	BGA 1	1	99	
Kocuria kristinae		6.83	0.08		99	1	
Staphylococcus cohnii subsp. urealyticum		2.32	0.09		1	99	
160443	**Acceptable identification**						**160443**
					OXI	NOV	
Staphylococcus cohnii subsp. urealyticum		93.20	0.30	URE 99	1	99	
Kocuria kristinae		6.80	-.04	MAN 91	99	1	
160450	**Low discrimination**						**160450**
					NOV	YEP	
Staphylococcus cohnii subsp. cohnii		88.08	0.43	BGA 1	99	12	
Staphylococcus kloosii		7.03	0.33		99	99	
Staphylococcus warneri		3.73	0.19		1	43	
160451	**Low discrimination**						**160451**
Staphylococcus cohnii subsp. cohnii		83.16	0.43	BGA 1			
Staphylococcus cohnii subsp. urealyticum		16.84	0.30				
160452	**Low discrimination**						**160452**
Staphylococcus kloosii		43.85	0.33	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		37.64	0.03	PHS 99			
Staphylococcus cohnii subsp. urealyticum		11.35	0.10				
160453	**Good identification**						**160453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.84	0.51	URE 99	99		
Aerococcus viridans		1.17	0.22		1		
160513	**Good identification**						**160513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	URE 99			
160550	**Low discrimination**						**160550**
					NOV	YEP	
Staphylococcus kloosii		55.46	0.39	VPT 10	99	99	
Staphylococcus cohnii subsp. cohnii		39.75	0.23	PHS 99	99	12	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		4.79	0.09		1	43	
160552	**Good identification**						**160552**
Staphylococcus kloosii		100.0	0.39	VPT 10			
160553	**Good identification**						**160553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.32	0.43	URE 99	99		
Aerococcus viridans		2.68	0.22		1		
160572	**Low discrimination**						**160572**
Staphylococcus kloosii		60.12	0.27	VPT 10			
Staphylococcus arlettae		39.88	0.12	PHS 99			
160601	**Good identification**						**160601**
Kocuria kristinae		100.0	0.30	GLR 1			
160640	**Low discrimination**						**160640**
					OXI	PYR	
Staphylococcus warneri		59.40	0.48	URE 98	1	8	
Staphylococcus petrasii subsp. pragensis		33.03	0.18		1	99	
Kocuria kristinae		3.42	0.08		99	93	
160641	**Good identification**						**160641**
					OXI	PYR	
Kocuria kristinae		99.64	0.49	GLR 1	99	93	
Staphylococcus warneri		0.36	0.14		1	8	
160642	**Low discrimination**						**160642**
					NOV	PYR	
Staphylococcus warneri		83.76	0.39	URE 98	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		7.07	-.04		99	7	
Staphylococcus haemolyticus		5.56	0.07		2	98	
160643	**Good identification**						**160643**
Kocuria kristinae		100.0	0.37	GLR 1			
160650	**Low discrimination**						**160650**
					NOV		
Staphylococcus warneri		80.94	0.46	URE 98	1		
Staphylococcus saprophyticus subsp. saprophyticus		13.32	0.09		99		
Staphylococcus pasteuri		3.16	0.02		1		
160672	**Good identification**						**160672**
Staphylococcus arlettae		100.0	0.37	VPT 1			
160673	**Acceptable identification**						**160673**
					NOV	YEP	
Staphylococcus arlettae		91.81	0.40	VPT 1	99	99	
Staphylococcus xylosus		4.47	0.03		99	40	
Aerococcus viridans		3.72	0.22		1	1	
160740	**Low discrimination**						**160740**
					OXI	PYR	
Staphylococcus warneri		83.25	0.38	URE 98	1	8	
Staphylococcus haemolyticus		11.69	0.13		1	98	
Kocuria kristinae		5.06	-.02		99	93	
160741	**Good identification**						**160741**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria kristinae		100.0	0.39	GLR 1			
160742	**Low discrimination**						**160742**
					PYR		
Staphylococcus warneri		80.84	0.28	URE 98	8		
Staphylococcus haemolyticus		19.16	0.08		98		
160743	**Good identification**						**160743**
Kocuria kristinae		100.0	0.27	GLR 1			
160750	**Low discrimination**						**160750**
					NOV	PYR	
Staphylococcus warneri		85.17	0.35	URE 98	1	8	
Staphylococcus haemolyticus		9.70	0.09		2	98	
Staphylococcus kloosii		5.13	0.18		99	52	
160772	**Good identification**						**160772**
Staphylococcus arlettae		100.0	0.53	VPT 1			
160773	**Good identification**						**160773**
					NOV	YEP	
Staphylococcus arlettae		91.41	0.55	VPT 1	99	99	
Staphylococcus xylosus		7.94	0.24		99	40	
Aerococcus viridans		0.65	0.22		1	1	
161413	**Low discrimination**						**161413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		72.13	0.27	URE 99	99		
Aerococcus viridans		27.87	0.29		1		
161433	**Good identification**						**161433**
Aerococcus viridans		100.0	0.29	VPT 1			
161450	**Good identification**						**161450**
Staphylococcus kloosii		100.0	0.33	VPT 10			
161452	**Low discrimination**						**161452**
					NOV	YEP	
Aerococcus viridans		53.81	0.34	VPT 1	1	1	
Staphylococcus kloosii		46.19	0.33		99	99	
161453	**Low discrimination**						**161453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		78.57	0.42	URE 99	99		
Aerococcus viridans		21.43	0.42		1		
161463	**Good identification**						**161463**
Aerococcus viridans		100.0	0.32	VPT 1			
161472	**Low discrimination**						**161472**
					NOV	YEP	
Aerococcus viridans		82.33	0.34	VPT 1	1	1	
Staphylococcus kloosii		17.67	0.21		99	99	
161473	**Good identification**						**161473**
Aerococcus viridans		100.0	0.42	VPT 1			
161533	**Good identification**						**161533**
Aerococcus viridans		100.0	0.29	VPT 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
161543	**Low discrimination**						**161543**
					NOV		
Aerococcus viridans		68.20	0.32	VPT 1	1		
Staphylococcus cohnii subsp. urealyticum		31.80	0.14		99		
161550	**Good identification**						**161550**
Staphylococcus kloosii		100.0	0.39	VPT 10			
161552	**Low discrimination**						**161552**
					NOV	YEP	
Staphylococcus kloosii		62.50	0.39	VPT 10	99	99	
Aerococcus viridans		37.50	0.34	PHS 99	1	1	
161553	**Low discrimination**						**161553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		61.11	0.35	URE 99	99		
Aerococcus viridans		38.89	0.42		1		
161563	**Good identification**						**161563**
Aerococcus viridans		100.0	0.32	VPT 1			
161572	**Low discrimination**						**161572**
					NOV	YEP	
Aerococcus viridans		59.07	0.34	VPT 1	1	1	
Staphylococcus kloosii		24.61	0.27		99	99	
Staphylococcus arlettae		16.33	0.12		99	99	
161573	**Low discrimination**						**161573**
					NOV	YEP	
Aerococcus viridans		86.43	0.42	VPT 1	1	1	
Staphylococcus arlettae		13.57	0.14		99	99	
161613	**Good identification**						**161613**
Aerococcus viridans		100.0	0.29	VPT 1			
161633	**Good identification**						**161633**
Aerococcus viridans		100.0	0.29	VPT 1			
161641	**Good identification**						**161641**
Kocuria kristinae		100.0	0.37	GLR 1			
161652	**Good identification**						**161652**
Aerococcus viridans		100.0	0.34	VPT 1			
161653	**Low discrimination**						**161653**
					OXI	NOV	
Aerococcus viridans		84.49	0.42	VPT 1	1	1	
Staphylococcus cohnii subsp. urealyticum		9.58	0.11		1	99	
Kocuria kristinae		5.93	-.04		99	1	
161663	**Good identification**						**161663**
Aerococcus viridans		100.0	0.32	VPT 1			
161672	**Low discrimination**						**161672**
					NOV	YEP	
Staphylococcus arlettae		82.84	0.37	VPT 1	99	99	
Aerococcus viridans		17.16	0.34		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
161673	**Low discrimination**						**161673**
					NOV	YEP	
Staphylococcus arlettae		73.29	0.40	VPT 1	99	99	
Aerococcus viridans		26.72	0.42		1	1	
161713	**Good identification**						**161713**
Aerococcus viridans		100.0	0.29	VPT 1			
161733	**Low discrimination**						**161733**
					NOV	YEP	
Aerococcus viridans		61.42	0.29	VPT 1	1	1	
Staphylococcus arlettae		38.58	0.14		99	99	
161741	**Low discrimination**						**161741**
					OXI		
Kocuria kristinae		87.76	0.27	GLR 1	99		
Rothia mucilaginosa		12.24	0.05		1		
161752	**Low discrimination**						**161752**
					NOV	YEP	
Aerococcus viridans		69.39	0.34	VPT 1	1	1	
Staphylococcus arlettae		19.18	0.12		99	99	
Staphylococcus kloosii		11.44	0.19		99	99	
161753	**Low discrimination**						**161753**
					NOV	YEP	
Aerococcus viridans		86.43	0.42	VPT 1	1	1	
Staphylococcus arlettae		13.57	0.14		99	99	
161763	**Low discrimination**						**161763**
					NOV	YEP	
Aerococcus viridans		67.98	0.32	VPT 1	1	1	
Staphylococcus arlettae		32.02	0.14		99	99	
161772	**Good identification**						**161772**
					NOV	YEP	
Staphylococcus arlettae		96.47	0.53	VPT 1	99	99	
Aerococcus viridans		3.53	0.34		1	1	
161773	**Good identification**						**161773**
					NOV	YEP	
Staphylococcus arlettae		92.08	0.55	VPT 1	99	99	
Aerococcus viridans		5.92	0.42		1	1	
Staphylococcus xylosus		2.00	0.12		99	40	
162000	**Good identification**						**162000**
Kocuria rosea		97.38	0.40	BGA 1			
Kocuria varians		2.62	0.17				
162020	**Good identification**						**162020**
Kocuria rosea		97.84	0.33	BGA 1			
Kocuria varians		2.16	0.08				
162240	**Low discrimination**						**162240**
					PYR		
Staphylococcus haemolyticus		86.32	0.30	ARG 92	98		
Staphylococcus auricularis		7.03	0.07	TRE 93	40		
Staphylococcus warneri		6.64	0.15		8		
162340	**Good identification**						**162340**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus haemolyticus		100.0	0.31	ARG 92				
162350	**Good identification**							**162350**
Staphylococcus haemolyticus		100.0	0.27	ARG 92				
162440	**Low discrimination**							**162440**
					NOV	PYR		
Staphylococcus auricularis		72.11	0.29	VPT 12	1	40		
Staphylococcus cohnii subsp. cohnii		12.89	0.17	BGA 14	99	1		
Staphylococcus haemolyticus		9.53	0.12	GLR 3	2	98		
162453	**Good identification**							**162453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	URE 99				
162553	**Good identification**							**162553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	URE 99				
162640	**Low discrimination**							**162640**
					PYR			
Staphylococcus haemolyticus		83.58	0.53	ARG 92	98			
Staphylococcus warneri		9.20	0.41		8			
Staphylococcus auricularis		5.90	0.29		40			
162642	**Acceptable identification**							**162642**
					PYR			
Staphylococcus haemolyticus		92.66	0.48	ARG 92	98			
Staphylococcus warneri		6.05	0.31		8			
Staphylococcus auricularis		1.29	0.09		40			
162650	**Low discrimination**							**162650**
					NOV	PYR		
Staphylococcus haemolyticus		81.26	0.49	ARG 92	2	98		
Staphylococcus warneri		11.03	0.38		1	8		
Staphylococcus saprophyticus subsp. bovis		5.26	0.22		99	99		
162652	**Low discrimination**							**162652**
					NOV	PYR		
Staphylococcus haemolyticus		88.40	0.44	ARG 92	2	98		
Staphylococcus warneri		7.11	0.28		1	8		
Staphylococcus saprophyticus subsp. bovis		2.54	0.09		99	99		
162740	**Low discrimination**							**162740**
					PYR			
Staphylococcus haemolyticus		89.36	0.55	ARG 92	98			
Rothia mucilaginosa		4.94	0.17		75			
Staphylococcus warneri		2.75	0.31		8			
162741	**Low discrimination**							**162741**
					OXI			
Rothia mucilaginosa		72.32	0.27	GLR 1	1			
Kocuria kristinae		14.19	0.16	ESL 90	99			
Staphylococcus haemolyticus		13.49	0.23		1			
162742	**Acceptable identification**							**162742**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		91.93	0.49	ARG 92	1	98	2	
Macrococcus caseolyticus		6.39	0.14		99	80	99	
Staphylococcus warneri		1.68	0.21		1	8	1	
162750	**Low discrimination**							**162750**
					NOV	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		78.90	0.50	ARG 92	2	98	
Staphylococcus saprophyticus subsp. bovis		18.10	0.34		99	99	
Staphylococcus warneri		3.00	0.28		1	8	
162752	**Low discrimination**						**162752**
					NOV	PYR	
Staphylococcus haemolyticus		88.93	0.45	ARG 92	2	98	
Staphylococcus saprophyticus subsp. bovis		9.07	0.22		99	99	
Staphylococcus warneri		2.00	0.18		1	8	
162771	**Good identification**						**162771**
Staphylococcus xylosus		100.0	0.33	VPT 14			
162773	**Good identification**						**162773**
Staphylococcus xylosus		98.58	0.43	VPT 14			
Staphylococcus arlettae		1.42	0.14	URE 99			
163453	**Good identification**						**163453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	URE 99			
163700	**Good identification**						**163700**
Rothia mucilaginosa		100.0	0.29	GLR 1			
163701	**Good identification**						**163701**
Rothia mucilaginosa		100.0	0.39	GLR 1			
163713	**Good identification**						**163713**
Staphylococcus lentus		100.0	0.30	VPT 7			
163733	**Low discrimination**						**163733**
					OXI		
Staphylococcus lentus		85.95	0.30	VPT 7	99		
Staphylococcus xylosus		14.05	0.07	GLR 1	1		
163740	**Good identification**						**163740**
Rothia mucilaginosa		98.01	0.37	GLR 1			
Staphylococcus haemolyticus		1.99	0.13				
163741	**Good identification**						**163741**
					OXI		
Rothia mucilaginosa		99.46	0.47	GLR 1	1		
Kocuria kristinae		0.54	0.04		99		
163753	**Acceptable identification**						**163753**
					OXI		
Staphylococcus lentus		91.49	0.36	VPT 7	99		
Staphylococcus xylosus		8.51	0.09	GLR 1	1		
163773	**Low discrimination**						**163773**
					OXI		
Staphylococcus xylosus		47.44	0.30	VPT 14	1		
Staphylococcus lentus		44.34	0.36	URE 99	99		
Staphylococcus equorum		5.49	0.12		1		
164050	**Good identification**						**164050**
Staphylococcus kloosii		100.0	0.33	VPT 10			
164052	**Good identification**						**164052**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.33	VPT 10		
164150	**Good identification**					**164150**
Staphylococcus kloosii		100.0	0.39	VPT 10		
164152	**Good identification**					**164152**
Staphylococcus kloosii		100.0	0.39	VPT 10		
164172	**Good identification**					**164172**
Staphylococcus kloosii		100.0	0.27	VPT 10		
164410	**Acceptable identification**					**164410**
					YEP	
Staphylococcus kloosii		93.02	0.47	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		6.98	0.12	MLT 95	12	
164412	**Good identification**					**164412**
Staphylococcus kloosii		100.0	0.48	VPT 10		
164413	**Good identification**					**164413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	URE 99		
164430	**Good identification**					**164430**
Staphylococcus kloosii		100.0	0.35	VPT 10		
164432	**Good identification**					**164432**
Staphylococcus kloosii		100.0	0.36	VPT 10		
164440	**Low discrimination**					**164440**
					YEP	
Staphylococcus kloosii		50.68	0.33	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		49.32	0.20	MAN 99	12	
164442	**Good identification**					**164442**
Staphylococcus kloosii		100.0	0.33	VPT 10		
164443	**Good identification**					**164443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	URE 99		
164450	**Very good identification**					**164450**
					YEP	
Staphylococcus kloosii		98.77	0.74	VPT 10	99	
Staphylococcus cohnii subsp. cohnii		1.24	0.22		12	
164451	**Low discrimination**					**164451**
					YEP	
Staphylococcus cohnii subsp. urealyticum		63.76	0.34	URE 99	64	
Staphylococcus cohnii subsp. cohnii		20.76	0.23	LAC 91	12	
Staphylococcus kloosii		15.48	0.33		99	
164452	**Very good identification**					**164452**
Staphylococcus kloosii		99.61	0.74	VPT 10		
Staphylococcus cohnii subsp. urealyticum		0.39	0.13			
164453	**Very good identification**					**164453**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		97.47	0.54	URE 99		
Staphylococcus kloosii		2.54	0.33			
164470	**Very good identification**					**164470**
Staphylococcus kloosii		100.0	0.61	VPT 10		
164472	**Very good identification**					**164472**
Staphylococcus kloosii		100.0	0.62	VPT 10		
164510	**Very good identification**					**164510**
Staphylococcus kloosii		100.0	0.53	VPT 10		
164512	**Very good identification**					**164512**
Staphylococcus kloosii		100.0	0.54	VPT 10		
164513	**Good identification**					**164513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	URE 99		
164530	**Good identification**					**164530**
Staphylococcus kloosii		100.0	0.41	VPT 10		
164532	**Good identification**					**164532**
Staphylococcus kloosii		100.0	0.42	VPT 10		
164540	**Good identification**					**164540**
Staphylococcus kloosii		100.0	0.39	VPT 10		
164542	**Good identification**					**164542**
Staphylococcus kloosii		100.0	0.39	VPT 10		
164550	**Very good identification**					**164550**
Staphylococcus kloosii		100.0	0.80	VPT 10		
164551	**Low discrimination**					**164551**
Staphylococcus kloosii		52.37	0.39	VPT 10		
Staphylococcus cohnii subsp. urealyticum		47.63	0.26	MNS 1		
164552	**Very good identification**					**164552**
Staphylococcus kloosii		99.91	0.80	VPT 10		
Staphylococcus cohnii subsp. urealyticum		0.09	0.06			
164553	**Low discrimination**					**164553**
Staphylococcus cohnii subsp. urealyticum		89.46	0.47	URE 99		
Staphylococcus kloosii		10.54	0.39			
164562	**Good identification**					**164562**
Staphylococcus kloosii		100.0	0.27	VPT 10		
164570	**Very good identification**					**164570**
Staphylococcus kloosii		100.0	0.67	VPT 10		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
164572	**Very good identification**					**164572**
Staphylococcus kloosii		99.01	0.68	VPT 10		
Staphylococcus arlettae		1.00	0.15			
164610	**Good identification**					**164610**
Staphylococcus kloosii		100.0	0.27	VPT 10		
164612	**Good identification**					**164612**
Staphylococcus kloosii		100.0	0.27	VPT 10		
164650	**Very good identification**					**164650**
Staphylococcus kloosii		100.0	0.53	VPT 10		
164652	**Very good identification**					**164652**
Staphylococcus kloosii		100.0	0.54	VPT 10		
164670	**Good identification**					**164670**
Staphylococcus kloosii		100.0	0.41	VPT 10		
164672	**Low discrimination**					**164672**
Staphylococcus arlettae		77.51	0.41	VPT 1		
Staphylococcus kloosii		22.49	0.41			
164673	**Good identification**					**164673**
Staphylococcus arlettae		100.0	0.43	VPT 1		
164710	**Good identification**					**164710**
Staphylococcus kloosii		100.0	0.33	VPT 10		
164712	**Good identification**					**164712**
Staphylococcus kloosii		100.0	0.33	VPT 10		
164750	**Very good identification**					**164750**
Staphylococcus kloosii		100.0	0.59	VPT 10		
164752	**Very good identification**					**164752**
Staphylococcus kloosii		97.52	0.60	VPT 10		
Staphylococcus arlettae		2.48	0.15	SUC 9		
164770	**Acceptable identification**					**164770**
Staphylococcus kloosii		90.08	0.47	VPT 10		
Staphylococcus arlettae		9.92	0.15	SUC 9		
164772	**Good identification**					**164772**
Staphylococcus arlettae		90.96	0.56	VPT 1		
Staphylococcus kloosii		9.04	0.47			
164773	**Very good identification**					**164773**
Staphylococcus arlettae		98.39	0.59	VPT 1		
Staphylococcus xylosus		1.61	0.13			

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
165050	**Good identification**					**165050**
Staphylococcus kloosii		100.0	0.33	VPT 10		
165052	**Good identification**					**165052**
Staphylococcus kloosii		100.0	0.33	VPT 10		
165150	**Good identification**					**165150**
Staphylococcus kloosii		100.0	0.39	VPT 10		
165152	**Good identification**					**165152**
Staphylococcus kloosii		100.0	0.39	VPT 10		
165172	**Good identification**					**165172**
Staphylococcus kloosii		100.0	0.27	VPT 10		
165410	**Good identification**					**165410**
Staphylococcus kloosii		100.0	0.47	VPT 10		
165412	**Good identification**					**165412**
Staphylococcus kloosii		100.0	0.48	VPT 10		
165413	**Good identification**					**165413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	URE 99		
165430	**Good identification**					**165430**
Staphylococcus kloosii		100.0	0.35	VPT 10		
165432	**Good identification**					**165432**
Staphylococcus kloosii		100.0	0.36	VPT 10		
165440	**Good identification**					**165440**
Staphylococcus kloosii		100.0	0.33	VPT 10		
165442	**Good identification**					**165442**
Staphylococcus kloosii		100.0	0.33	VPT 10		
165450	**Very good identification**					**165450**
Staphylococcus kloosii		100.0	0.74	VPT 10		
165452	**Very good identification**					**165452**
Staphylococcus kloosii		99.85	0.74	VPT 10		
Staphylococcus cohnii subsp. urealyticum		0.15	0.05			
165453	**Acceptable identification**					**165453**
Staphylococcus cohnii subsp. urealyticum		93.73	0.46	URE 99		
Staphylococcus kloosii		6.27	0.33			
165470	**Very good identification**					**165470**
Staphylococcus kloosii		100.0	0.61	VPT 10		
165472	**Very good identification**					**165472**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.62	VPT 10		
165510	**Very good identification**					**165510**
Staphylococcus kloosii		100.0	0.53	VPT 10		
165512	**Very good identification**					**165512**
Staphylococcus kloosii		100.0	0.54	VPT 10		
165530	**Good identification**					**165530**
Staphylococcus kloosii		100.0	0.41	VPT 10		
165532	**Good identification**					**165532**
Staphylococcus kloosii		100.0	0.42	VPT 10		
165540	**Good identification**					**165540**
Staphylococcus kloosii		100.0	0.39	VPT 10		
165542	**Good identification**					**165542**
Staphylococcus kloosii		100.0	0.39	VPT 10		
165550	**Very good identification**					**165550**
Staphylococcus kloosii		100.0	0.80	VPT 10		
165551	**Low discrimination**					**165551**
Staphylococcus kloosii		73.87	0.39	VPT 10		
Staphylococcus cohnii subsp. urealyticum		26.13	0.18	MNS 1		
165552	**Very good identification**					**165552**
Staphylococcus kloosii		100.0	0.80	VPT 10		
165553	**Low discrimination**					**165553**
Staphylococcus cohnii subsp. urealyticum		76.75	0.38	URE 99		
Staphylococcus kloosii		23.25	0.39			
165562	**Good identification**					**165562**
Staphylococcus kloosii		100.0	0.27	VPT 10		
165570	**Very good identification**					**165570**
Staphylococcus kloosii		100.0	0.67	VPT 10		
165572	**Very good identification**					**165572**
Staphylococcus kloosii		99.01	0.68	VPT 10		
Staphylococcus arlettae		1.00	0.15			
165610	**Good identification**					**165610**
Staphylococcus kloosii		100.0	0.27	VPT 10		
165612	**Good identification**					**165612**
Staphylococcus kloosii		100.0	0.27	VPT 10		
165650	**Very good identification**					**165650**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.53	VPT 10		
165652	**Very good identification**					**165652**
Staphylococcus kloosii		100.0	0.54	VPT 10		
165670	**Good identification**					**165670**
Staphylococcus kloosii		100.0	0.41	VPT 10		
165672	**Low discrimination**					**165672**
Staphylococcus arlettae		77.51	0.41	VPT 1		
Staphylococcus kloosii		22.49	0.41			
165673	**Good identification**					**165673**
Staphylococcus arlettae		100.0	0.43	VPT 1		
165710	**Good identification**					**165710**
Staphylococcus kloosii		100.0	0.33	VPT 10		
165712	**Good identification**					**165712**
Staphylococcus kloosii		100.0	0.33	VPT 10		
165750	**Very good identification**					**165750**
Staphylococcus kloosii		100.0	0.59	VPT 10		
165752	**Very good identification**					**165752**
Staphylococcus kloosii		97.52	0.60	VPT 10		
Staphylococcus arlettae		2.48	0.15	SUC 9		
165770	**Acceptable identification**					**165770**
Staphylococcus kloosii		90.08	0.47	VPT 10		
Staphylococcus arlettae		9.92	0.15	SUC 9		
165772	**Good identification**					**165772**
Staphylococcus arlettae		90.96	0.56	VPT 1		
Staphylococcus kloosii		9.04	0.47			
165773	**Very good identification**					**165773**
Staphylococcus arlettae		99.59	0.59	VPT 1		
Staphylococcus xylosus		0.41	0.00			
166001	**Good identification**					**166001**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86		
166101	**Good identification**					**166101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.33	ARG 86		
166401	**Good identification**					**166401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	ARG 86		
166450	**Good identification**					**166450**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Staphylococcus kloosii	100.0	0.33	VPT 10		
166452	**Good identification**					**166452**
	Staphylococcus kloosii	100.0	0.33	VPT 10		
166453	**Good identification**					**166453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	URE 99		
166501	**Good identification**					**166501**
	Staphylococcus schleiferi subsp. schleiferi	100.0	0.43	ARG 86		
166550	**Low discrimination**					**166550**
	Staphylococcus kloosii	79.03	0.39	VPT 10		
	Staphylococcus saprophyticus subsp. bovis	20.97	0.09	NIT 1		
166552	**Good identification**					**166552**
	Staphylococcus kloosii	100.0	0.39	VPT 10		
166553	**Good identification**					**166553**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.31	URE 99		
166572	**Good identification**					**166572**
	Staphylococcus kloosii	100.0	0.27	VPT 10		
166773	**Acceptable identification**					**166773**
	Staphylococcus xylosus	92.90	0.31	VPT 14		
	Staphylococcus arlettae	7.10	0.18	URE 99		
167450	**Good identification**					**167450**
	Staphylococcus kloosii	100.0	0.33	VPT 10		
167452	**Good identification**					**167452**
	Staphylococcus kloosii	100.0	0.33	VPT 10		
167453	**Good identification**					**167453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.30	URE 99		
167550	**Good identification**					**167550**
	Staphylococcus kloosii	100.0	0.39	VPT 10		
167552	**Good identification**					**167552**
	Staphylococcus kloosii	100.0	0.39	VPT 10		
167572	**Good identification**					**167572**
	Staphylococcus kloosii	100.0	0.27	VPT 10		
167771	**Low discrimination**					**167771**
	Staphylococcus gallinarum	76.79	0.27	VPT 1		
	Staphylococcus xylosus	9.89	0.10	URE 92		
	Staphylococcus equorum	6.74	0.08			
167773	**Low discrimination**					**167773**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					OXI	
Staphylococcus gallinarum		70.83	0.35	VPT 1	1	
Staphylococcus xylosus		10.61	0.19	URE 92	1	
Staphylococcus lentus		8.79	0.23		99	
172641	**Good identification**					**172641**
Staphylococcus lugdunensis		100.0	0.30	GLR 2		
172643	**Good identification**					**172643**
Staphylococcus lugdunensis		100.0	0.49	GLR 2		
172741	**Good identification**					**172741**
Staphylococcus lugdunensis		100.0	0.30	GLR 2		
172743	**Good identification**					**172743**
Staphylococcus lugdunensis		100.0	0.49	GLR 2		
174450	**Good identification**					**174450**
Staphylococcus kloosii		100.0	0.33	VPT 10		
174452	**Good identification**					**174452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
174550	**Good identification**					**174550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
174552	**Good identification**					**174552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
174572	**Good identification**					**174572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
175450	**Good identification**					**175450**
Staphylococcus kloosii		100.0	0.33	VPT 10		
175452	**Good identification**					**175452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
175550	**Good identification**					**175550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
175552	**Good identification**					**175552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
175572	**Good identification**					**175572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
200000	**Low discrimination**					**200000**
					OXI	
Dermacoccus nishinomiyaensis		62.46	1.00		93	
Micrococcus sp.		33.50	0.85		89	
Kocuria varians		3.60	0.82		13	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
200001	**Low discrimination**						**200001**
					OXI		
Micrococcus sp.		41.77	0.50	MNS 2	89		
Dermacoccus nishinomiyaensis		38.55	0.59		93		
Kocuria varians		19.14	0.60		13		
200002	**Low discrimination**						**200002**
					OXI		
Kocuria varians		51.07	0.71		13		
Dermacoccus nishinomiyaensis		31.72	0.59		93		
Micrococcus sp.		17.01	0.44		89		
200003	**Low discrimination**						**200003**
					OXI		
Kocuria varians		86.93	0.49	MNS 8	13		
Micrococcus sp.		6.80	0.09		89		
Dermacoccus nishinomiyaensis		6.27	0.18		93		
200010	**Low discrimination**						**200010**
					YEP		
Macrococcus equipersicus		60.70	0.84		99		
Macrococcus bovicus		29.39	0.76		99		
Dermacoccus nishinomiyaensis		6.16	0.59		1		
200011	**Low discrimination**						**200011**
					OXI	NOV	
Macrococcus equipersicus		51.99	0.43	MNS 1	99	99	
Macrococcus bovicus		25.17	0.35		99	99	
Staphylococcus capitis subsp. capitis		8.29	0.23		1	1	
200012	**Low discrimination**						**200012**
					NOV	OXI	
Macrococcus equipersicus		55.45	0.43	LAC 1	99	99	
Macrococcus bovicus		26.85	0.35		99	99	
Kocuria varians		9.06	0.30		1	13	
200020	**Low discrimination**						**200020**
					OXI		
Kocuria varians		55.68	0.73		13		
Dermacoccus nishinomiyaensis		27.76	0.59		93		
Micrococcus sp.		14.89	0.44		89		
200021	**Low discrimination**						**200021**
					OXI		
Kocuria varians		87.94	0.51	MNS 8	13		
Micrococcus sp.		5.52	0.09		89		
Dermacoccus nishinomiyaensis		5.09	0.18		93		
200022	**Very good identification**						**200022**
					OXI		
Kocuria varians		97.33	0.62		13		
Dermacoccus nishinomiyaensis		1.74	0.18		93		
Micrococcus sp.		0.93	0.03		89		
200023	**Good identification**						**200023**
Kocuria varians		100.0	0.40	MNS 8			
200030	**Low discrimination**						**200030**
					NOV	OXI	
Macrococcus equipersicus		52.50	0.43	XYL 1	99	99	
Macrococcus bovicus		25.42	0.35		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		10.68	0.32		1	13	
200040	**Low discrimination**						**200040**
					NOV	OXI	
Micrococcus sp.		51.34	0.62	MLT 7	1	89	
Macrococcus equipersicus		18.03	0.67		99	99	
Kocuria varians		16.10	0.69		1	13	
200041	**Low discrimination**						**200041**
					NOV	OXI	
Kocuria varians		49.56	0.47	MNS 8	1	13	
Micrococcus sp.		37.09	0.27		1	89	
Macrococcus equipersicus		6.45	0.26		99	99	
200042	**Low discrimination**						**200042**
					NOV	OXI	
Kocuria varians		83.10	0.57		1	13	
Micrococcus sp.		9.49	0.21		1	89	
Macrococcus equipersicus		3.33	0.26		99	99	
200043	**Good identification**						**200043**
					NOV	OXI	
Kocuria varians		95.26	0.36	MNS 8	1	13	
Micrococcus sp.		2.55	-.14		1	89	
Staphylococcus hominis subsp. novobiosepticus		2.19	-.08		99	1	
200050	**Low discrimination**						**200050**
					NOV		
Macrococcus equipersicus		70.39	0.93		99		
Macrococcus bovicus		29.42	0.84		99		
Micrococcus sp.		0.11	0.21		1		
200051	**Low discrimination**						**200051**
					OXI	YEP	
Macrococcus equipersicus		69.57	0.52	MNS 1	99	99	
Macrococcus bovicus		29.07	0.43		99	99	
Staphylococcus cohnii subsp. cohnii		0.86	0.18		1	12	
200052	**Low discrimination**						**200052**
					OXI		
Macrococcus equipersicus		68.42	0.52	LAC 1	99		
Macrococcus bovicus		28.59	0.43		99		
Staphylococcus saprophyticus subsp. saprophyticus		1.66	0.08		1		
200060	**Low discrimination**						**200060**
					NOV	OXI	
Kocuria varians		87.00	0.59		1	13	
Micrococcus sp.		7.98	0.21		1	89	
Macrococcus equipersicus		2.80	0.26		99	99	
200061	**Good identification**						**200061**
					OXI		
Kocuria varians		97.89	0.38	MNS 8	13		
Micrococcus sp.		2.11	-.14		89		
200062	**Good identification**						**200062**
Kocuria varians		100.0	0.48				
200070	**Low discrimination**						**200070**
					NOV	OXI	
Macrococcus equipersicus		69.72	0.52	XYL 1	99	99	
Macrococcus bovicus		29.13	0.43		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		1.15	0.18		1	13	
200100	**Low discrimination**						**200100**
					OXI		
Dermacoccus nishinomiyaensis		66.87	0.85		93		
Kocuria varians		30.36	0.86		13		
Micrococcus sp.		1.90	0.44		89		
200101	**Low discrimination**						**200101**
					OXI		
Kocuria varians		78.52	0.64	MNS 8	13		
Dermacoccus nishinomiyaensis		20.09	0.44		93		
Micrococcus sp.		1.15	0.09		89		
200102	**Good identification**						**200102**
					OXI		
Kocuria varians		92.42	0.75		13		
Dermacoccus nishinomiyaensis		7.29	0.44		93		
Micrococcus sp.		0.21	0.03		89		
200103	**Good identification**						**200103**
					OXI		
Kocuria varians		99.09	0.53	MNS 8	13		
Dermacoccus nishinomiyaensis		0.91	0.03		93		
200110	**Low discrimination**						**200110**
					YEP		
Macrococcus equipersicus		64.47	0.77		99		
Macrococcus bovicus		31.22	0.69		99		
Dermacoccus nishinomiyaensis		2.91	0.44		1		
200111	**Low discrimination**						**200111**
					NOV	OXI	
Macrococcus equipersicus		58.63	0.36	MNS 1	99	99	
Macrococcus bovicus		28.39	0.28		99	99	
Kocuria varians		10.33	0.23		1	13	
200112	**Low discrimination**						**200112**
					NOV	OXI	
Macrococcus equipersicus		47.60	0.36	LAC 1	99	99	
Kocuria varians		27.21	0.33		1	13	
Macrococcus bovicus		23.05	0.28		99	99	
200120	**Good identification**						**200120**
					OXI		
Kocuria varians		93.82	0.77		13		
Dermacoccus nishinomiyaensis		5.94	0.44		93		
Micrococcus sp.		0.17	0.03		89		
200121	**Very good identification**						**200121**
					OXI		
Kocuria varians		99.27	0.55	MNS 8	13		
Dermacoccus nishinomiyaensis		0.73	0.03		93		
200122	**Very good identification**						**200122**
					OXI		
Kocuria varians		99.77	0.65		13		
Dermacoccus nishinomiyaensis		0.23	0.03		93		
200123	**Good identification**						**200123**
Kocuria varians		100.0	0.43	MNS 8			
200130	**Low discrimination**						**200130**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	OXI		
Macrococcus equipersicus		44.62	0.36	XYL 1	99	99		
Kocuria varians		31.77	0.35		1	13		
Macrococcus bovicus		21.60	0.28		99	99		
200140	**Low discrimination**							**200140**
					OXI	YEP	NOV	
Kocuria varians		67.64	0.72		13	80	1	
Macrococcus equipersicus		21.64	0.59		99	99	99	
Dermacoccus nishinomiyaensis		6.86	0.44		93	1	80	
200141	**Acceptable identification**							**200141**
					NOV	OXI	YEP	
Kocuria varians		92.76	0.50	MNS 8	1	13	80	
Macrococcus equipersicus		3.45	0.18		99	99	99	
Staphylococcus hominis subsp. novobiosepticus		2.23	0.07		99	1	1	
200142	**Very good identification**							**200142**
					NOV	OXI	YEP	
Kocuria varians		96.67	0.61		1	13	80	
Staphylococcus hominis subsp. novobiosepticus		1.55	0.14		99	1	1	
Macrococcus equipersicus		1.11	0.18		99	99	99	
200143	**Good identification**							**200143**
					NOV			
Kocuria varians		95.61	0.39	MNS 8	1			
Staphylococcus hominis subsp. novobiosepticus		4.39	0.02		99			
200150	**Low discrimination**							**200150**
					NOV	OXI		
Macrococcus equipersicus		70.43	0.86		99	99		
Macrococcus bovicus		29.43	0.76		99	99		
Kocuria varians		0.12	0.31		1	13		
200151	**Low discrimination**							**200151**
					NOV	OXI		
Macrococcus equipersicus		69.82	0.45	MNS 1	99	99		
Macrococcus bovicus		29.18	0.35		99	99		
Kocuria varians		1.00	0.09		1	13		
200152	**Low discrimination**							**200152**
					NOV	OXI		
Macrococcus equipersicus		68.29	0.45	LAC 1	99	99		
Macrococcus bovicus		28.54	0.35		99	99		
Kocuria varians		3.17	0.20		1	13		
200160	**Very good identification**							**200160**
					OXI	YEP	NOV	
Kocuria varians		98.80	0.63		13	80	1	
Macrococcus equipersicus		0.91	0.18		99	99	99	
Dermacoccus nishinomiyaensis		0.29	0.03		93	1	80	
200161	**Good identification**							**200161**
Kocuria varians		100.0	0.41	MNS 8				
200162	**Good identification**							**200162**
Kocuria varians		100.0	0.52					
200163	**Good identification**							**200163**
Kocuria varians		100.0	0.30	MNS 8				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
200170	**Low discrimination**						**200170**
					NOV	OXI	
Macrococcus equipersicus		67.77	0.45	XYL 1	99	99	
Macrococcus bovicus		28.32	0.35		99	99	
Kocuria varians		3.92	0.22		1	13	
200200	**Low discrimination**						**200200**
					OXI		
Dermacoccus nishinomiyaensis		50.78	0.75	SUC 6	93		
Micrococcus sp.		42.19	0.65		89		
Kocuria varians		6.26	0.64		13		
200201	**Low discrimination**						**200201**
					OXI		
Micrococcus sp.		42.83	0.30	SUC 9	89		
Kocuria varians		27.05	0.43	MNS 2	13		
Dermacoccus nishinomiyaensis		25.51	0.34		93		
200202	**Low discrimination**						**200202**
					OXI		
Kocuria varians		64.10	0.53	SUC 12	13		
Dermacoccus nishinomiyaensis		18.64	0.34		93		
Micrococcus sp.		15.48	0.23		89		
200203	**Low discrimination**						**200203**
					OXI	PYR	
Kocuria varians		84.12	0.31	SUC 12	13	10	
Kocuria kristinae		8.00	0.02	MNS 8	99	93	
Micrococcus sp.		4.77	-.11		89	63	
200210	**Low discrimination**						**200210**
					YEP		
Macrococcus bovicus		51.43	0.64		99		
Macrococcus equipersicus		42.02	0.64		99		
Dermacoccus nishinomiyaensis		2.75	0.34		1		
200211	**Low discrimination**						**200211**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		55.48	0.29	URE 1	1	1	
Macrococcus bovicus		22.67	0.23	NIT 89	99	99	
Macrococcus equipersicus		18.52	0.23		99	99	
200220	**Low discrimination**						**200220**
					OXI		
Kocuria varians		68.99	0.55	SUC 12	13		
Dermacoccus nishinomiyaensis		16.10	0.34		93		
Micrococcus sp.		13.38	0.23		89		
200221	**Good identification**						**200221**
					OXI		
Kocuria varians		95.65	0.33	SUC 12	13		
Micrococcus sp.		4.35	-.11	MNS 8	89		
200222	**Good identification**						**200222**
Kocuria varians		100.0	0.44	SUC 12			
200230	**Low discrimination**						**200230**
Staphylococcus vitulinus		65.82	0.29	URE 1			
Macrococcus bovicus		16.26	0.23	NIT 91			
Macrococcus equipersicus		13.29	0.23				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
200240	**Low discrimination**							**200240**
					NOV	OXI		
Staphylococcus hominis subsp. novobiosepticus		33.45	0.51	VPT 90	99	1		
Micrococcus sp.		25.46	0.41	NIT 90	1	89		
Kocuria varians		11.01	0.51		1	13		
200241	**Low discrimination**							**200241**
					NOV	OXI	PYR	
Staphylococcus hominis subsp. novobiosepticus		57.99	0.39	VPT 90	99	1	1	
Kocuria kristinae		29.20	0.34	NIT 90	1	99	93	
Kocuria varians		6.64	0.29		1	13	10	
200242	**Low discrimination**							**200242**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		58.53	0.45	VPT 90	99			
Staphylococcus hominis subsp. hominis		20.36	0.47	NIT 90	1			
Kocuria varians		10.09	0.40		1			
200243	**Low discrimination**							**200243**
					NOV	OXI	PYR	
Staphylococcus hominis subsp. novobiosepticus		72.73	0.33	VPT 90	99	1	1	
Kocuria kristinae		17.00	0.22	NIT 90	1	99	93	
Kocuria varians		4.36	0.18		1	13	10	
200250	**Low discrimination**							**200250**
					OXI	NOV		
Macrococcus bovicus		50.16	0.72		99	99		
Macrococcus equipersicus		47.49	0.73		99	99		
Staphylococcus warneri		1.70	0.53		1	1		
200251	**Low discrimination**							**200251**
					OXI	NOV		
Macrococcus bovicus		43.93	0.31	MNS 1	99	99		
Macrococcus equipersicus		41.59	0.32		99	99		
Staphylococcus capitis subsp. ureolyticus		5.17	0.00		1	1		
200252	**Low discrimination**							**200252**
					OXI	NOV		
Staphylococcus saprophyticus subsp. saprophyticus		38.96	0.29	TRE 99	1	99		
Staphylococcus warneri		18.57	0.43		1	1		
Macrococcus bovicus		16.57	0.31		99	99		
200260	**Low discrimination**							**200260**
					NOV	OXI		
Kocuria varians		82.43	0.42	SUC 12	1	13		
Staphylococcus hominis subsp. novobiosepticus		7.20	0.10		99	1		
Micrococcus sp.		5.48	0.00		1	89		
200262	**Low discrimination**							**200262**
					NOV			
Kocuria varians		81.65	0.30	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		13.62	0.04		99			
Staphylococcus hominis subsp. hominis		4.74	0.06		1			
200270	**Low discrimination**							**200270**
					OXI	NOV		
Macrococcus bovicus		50.49	0.31	XYL 1	99	99		
Macrococcus equipersicus		47.80	0.32		99	99		
Staphylococcus warneri		1.72	0.12		1	1		
200300	**Low discrimination**							**200300**
					OXI			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		48.69	0.61	SUC 6	93			
Kocuria varians		47.23	0.68		13			
Micrococcus sp.		2.15	0.23		89			
200301	**Low discrimination**							**200301**
					OXI	NOV		
Kocuria varians		83.82	0.46	SUC 12	13	1		
Dermacoccus nishinomiyaensis		10.04	0.20	MNS 8	93	80		
Staphylococcus hominis subsp. novobiosepticus		3.25	0.07		1	99		
200302	**Good identification**							**200302**
					OXI	NOV		
Kocuria varians		93.63	0.57	SUC 12	13	1		
Dermacoccus nishinomiyaensis		3.46	0.20		93	80		
Staphylococcus hominis subsp. novobiosepticus		2.41	0.14		1	99		
200303	**Acceptable identification**							**200303**
					NOV			
Kocuria varians		93.11	0.35	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		6.89	0.02	MNS 8	99			
200310	**Low discrimination**							**200310**
					YEP			
Macrococcus bovicus		53.50	0.56		99			
Macrococcus equipersicus		43.71	0.56		99			
Dermacoccus nishinomiyaensis		1.27	0.20		1			
200320	**Very good identification**							**200320**
					OXI			
Kocuria varians		97.12	0.59	SUC 12	13			
Dermacoccus nishinomiyaensis		2.88	0.20		93			
200321	**Good identification**							**200321**
Kocuria varians		100.0	0.37	SUC 12				
200322	**Good identification**							**200322**
Kocuria varians		100.0	0.47	SUC 12				
200340	**Low discrimination**							**200340**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		76.57	0.61	VPT 90	99			
Kocuria varians		12.60	0.55	NIT 90	1			
Staphylococcus hominis subsp. hominis		4.41	0.47		1			
200341	**Low discrimination**							**200341**
					NOV	OXI	PYR	
Staphylococcus hominis subsp. novobiosepticus		88.63	0.48	VPT 90	99	1	1	
Kocuria varians		5.07	0.33	NIT 90	1	13	10	
Kocuria kristinae		4.96	0.25		1	99	93	
200342	**Low discrimination**							**200342**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		80.99	0.55	VPT 90	99			
Staphylococcus hominis subsp. hominis		10.17	0.48	NIT 90	1			
Kocuria varians		6.98	0.43		1			
200343	**Low discrimination**							**200343**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		89.71	0.43	VPT 90	99			
Staphylococcus epidermidis		4.35	0.24	NIT 90	1			
Kocuria varians		2.69	0.21		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
200350	**Low discrimination**						**200350**
					OXI	NOV	
Macrococcus bovicus		49.94	0.64		99	99	
Macrococcus equipersicus		47.28	0.65		99	99	
Staphylococcus warneri		1.25	0.42		1	1	
200352	**Low discrimination**						**200352**
					OXI	NOV	
Staphylococcus hominis subsp. hominis		24.19	0.31	NIT 99	1	1	
Macrococcus bovicus		18.53	0.23	MAN 13	99	99	
Macrococcus equipersicus		17.54	0.24		99	99	
200360	**Low discrimination**						**200360**
					NOV		
Kocuria varians		84.40	0.45	SUC 12	1		
Staphylococcus hominis subsp. novobiosepticus		14.75	0.20		99		
Staphylococcus hominis subsp. hominis		0.85	0.05		1		
200362	**Low discrimination**						**200362**
					NOV		
Kocuria varians		72.70	0.34	SUC 12	1		
Staphylococcus hominis subsp. novobiosepticus		24.25	0.14		99		
Staphylococcus hominis subsp. hominis		3.05	0.07		1		
200400	**Low discrimination**						**200400**
					NOV	YEP	
Dermacoccus nishinomiyaensis		37.15	0.59	TRE 1	80	1	
Micrococcus sp.		19.92	0.44		1	41	
Macrococcus equipersicus		19.28	0.58		99	99	
200401	**Low discrimination**						**200401**
					NOV	OXI	
Staphylococcus cohnii subsp. cohnii		54.90	0.47	URE 1	99	1	
Kocuria varians		25.09	0.37		1	13	
Micrococcus sp.		7.35	0.09		1	89	
200402	**Low discrimination**						**200402**
					OXI		
Kocuria varians		82.38	0.48	TRE 7	13		
Dermacoccus nishinomiyaensis		6.87	0.18		93		
Micrococcus sp.		3.68	0.03		89		
200410	**Low discrimination**						**200410**
					OXI	YEP	
Macrococcus equipersicus		57.44	0.84		99	99	
Macrococcus bovicus		41.72	0.80		99	99	
Staphylococcus cohnii subsp. cohnii		0.56	0.48		1	12	
200411	**Low discrimination**						**200411**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		37.38	0.49	URE 1	1	12	
Macrococcus equipersicus		35.78	0.43		99	99	
Macrococcus bovicus		25.99	0.39		99	99	
200412	**Low discrimination**						**200412**
					OXI		
Macrococcus equipersicus		53.98	0.43	LAC 1	99		
Macrococcus bovicus		39.21	0.39		99		
Staphylococcus saprophyticus subsp. saprophyticus		3.55	0.08		1		
200413	**Low discrimination**						**200413**
					NOV	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus cohnii subsp. urealyticum		82.02	0.36	GLR 99	99	53			
Aerococcus viridans		8.35	0.27		1	95			
Staphylococcus cohnii subsp. cohnii		3.63	0.08		99	1			
200420	**Low discrimination**								**200420**
					OXI				
Kocuria varians		87.15	0.50	TRE 7	13				
Dermacoccus nishinomiyaensis		5.83	0.18		93				
Micrococcus sp.		3.13	0.03		89				
200421	**Acceptable identification**								**200421**
					NOV				
Kocuria varians		94.08	0.28	TRE 7	1				
Staphylococcus cohnii subsp. cohnii		5.92	0.05	MNS 8	99				
200422	**Good identification**								**200422**
Kocuria varians		100.0	0.38	TRE 7					
200430	**Low discrimination**								**200430**
					NOV	OXI			
Macrococcus equipersicus		57.10	0.43	XYL 1	99	99			
Macrococcus bovicus		41.47	0.39		99	99			
Kocuria varians		0.87	0.09		1	13			
200433	**Good identification**								**200433**
Aerococcus viridans		100.0	0.27	URE 1					
200440	**Low discrimination**								**200440**
					OXI	YEP			
Macrococcus equipersicus		66.05	0.67	MAN 95	99	99			
Staphylococcus cohnii subsp. cohnii		11.25	0.56		1	12			
Macrococcus bovicus		7.95	0.47		99	99			
200441	**Low discrimination**								**200441**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		85.29	0.57	URE 1	1	99	1	12	
Macrococcus equipersicus		4.67	0.26		99	99	1	99	
Kocuria kristinae		4.13	0.20		99	1	93	15	
200442	**Low discrimination**								**200442**
					NOV				
Staphylococcus warneri		35.37	0.46	SUC 95	1				
Staphylococcus saprophyticus subsp. saprophyticus		21.91	0.21		99				
Kocuria varians		21.60	0.34		1				
200443	**Low discrimination**								**200443**
					OXI	NOV			
Staphylococcus cohnii subsp. urealyticum		52.22	0.31	GLR 99	1	99			
Aerococcus viridans		12.65	0.30	MAN 91	1	1			
Kocuria kristinae		12.30	0.07		99	1			
200450	**Low discrimination**								**200450**
					OXI	YEP			
Macrococcus equipersicus		60.69	0.93		99	99			
Macrococcus bovicus		38.04	0.88		99	99			
Staphylococcus cohnii subsp. cohnii		0.69	0.58		1	12			
200451	**Low discrimination**								**200451**
					OXI	YEP			
Staphylococcus cohnii subsp. cohnii		41.66	0.59	URE 1	1	12			
Macrococcus equipersicus		34.05	0.52		99	99			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		21.34	0.47		99	99		
200452	**Low discrimination**							**200452**
					OXI			
Staphylococcus saprophyticus subsp. saprophyticus		56.16	0.49	SUC 91	1			
Macrococcus equipersicus		23.35	0.52		99			
Macrococcus bovicus		14.64	0.47		99			
200453	**Low discrimination**							**200453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		84.11	0.52	GLR 99	99	53		
Aerococcus viridans		6.05	0.39		1	95		
Staphylococcus saprophyticus subsp. saprophyticus		4.09	0.08		99	7		
200460	**Low discrimination**							**200460**
					NOV	OXI	YEP	
Kocuria varians		62.82	0.36	TRE 7	1	13	80	
Macrococcus equipersicus		26.87	0.26		99	99	99	
Staphylococcus cohnii subsp. cohnii		4.58	0.15		99	1	12	
200463	**Low discrimination**							**200463**
					PYR			
Aerococcus viridans		79.87	0.30	URE 1	95			
Kocuria varians		20.13	0.03	ESL 90	10			
200470	**Low discrimination**							**200470**
					OXI			
Macrococcus equipersicus		60.08	0.52	XYL 1	99			
Macrococcus bovicus		37.66	0.47		99			
Staphylococcus kloosii		1.11	0.34		1			
200473	**Low discrimination**							**200473**
					NOV			
Aerococcus viridans		82.40	0.39	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		11.58	0.11	ESL 90	99			
Staphylococcus xylosus		6.02	-.04		99			
200500	**Low discrimination**							**200500**
					OXI	YEP	NOV	
Kocuria varians		57.56	0.63	TRE 7	13	80	1	
Macrococcus equipersicus		19.86	0.51		99	99	99	
Dermacoccus nishinomiyaensis		17.01	0.44		93	1	80	
200501	**Low discrimination**							**200501**
					NOV	OXI	YEP	
Kocuria varians		80.88	0.41	TRE 7	1	13	80	
Staphylococcus cohnii subsp. cohnii		13.11	0.27	MNS 8	99	1	12	
Macrococcus equipersicus		3.24	0.09		99	99	99	
200502	**Good identification**							**200502**
					OXI	YEP	NOV	
Kocuria varians		97.38	0.51	TRE 7	13	80	1	
Macrococcus equipersicus		1.20	0.09		99	99	99	
Dermacoccus nishinomiyaensis		1.03	0.03		93	1	80	
200503	**Low discrimination**							**200503**
					NOV	PYR		
Kocuria varians		89.71	0.30	TRE 7	1	10		
Staphylococcus cohnii subsp. urealyticum		5.71	0.08	MNS 8	99	53		
Aerococcus viridans		4.58	0.17		1	95		
200510	**Low discrimination**							**200510**
					OXI	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus equipersicus		57.67	0.77		99	99		
Macrococcus bovicus		41.89	0.73		99	99		
Staphylococcus cohnii subsp. cohnii		0.15	0.28		1	12		
200511	**Low discrimination**							**200511**
					OXI	YEP		
Macrococcus equipersicus		49.49	0.36	MNS 1	99	99		
Macrococcus bovicus		35.94	0.31		99	99		
Staphylococcus cohnii subsp. cohnii		13.40	0.29		1	12		
200512	**Low discrimination**							**200512**
					OXI			
Macrococcus equipersicus		54.20	0.36	LAC 1	99			
Macrococcus bovicus		39.36	0.31		99			
Staphylococcus kloosii		2.79	0.27		1			
200513	**Low discrimination**							**200513**
					NOV			
Staphylococcus cohnii subsp. urealyticum		80.80	0.29	GLR 99	99			
Aerococcus viridans		19.20	0.27		1			
200520	**Very good identification**							**200520**
					OXI	YEP	NOV	
Kocuria varians		98.19	0.53	TRE 7	13	80	1	
Macrococcus equipersicus		0.97	0.09		99	99	99	
Dermacoccus nishinomiyaensis		0.83	0.03		93	1	80	
200521	**Good identification**							**200521**
Kocuria varians		100.0	0.32	TRE 7				
200522	**Good identification**							**200522**
Kocuria varians		100.0	0.42	TRE 7				
200530	**Low discrimination**							**200530**
					NOV	OXI		
Macrococcus equipersicus		56.18	0.36	XYL 1	99	99		
Macrococcus bovicus		40.81	0.31		99	99		
Kocuria varians		3.01	0.12		1	13		
200533	**Good identification**							**200533**
Aerococcus viridans		100.0	0.27	URE 1				
200540	**Low discrimination**							**200540**
					NOV	OXI		
Macrococcus equipersicus		67.42	0.59	MAN 95	99	99		
Kocuria varians		15.86	0.49		1	13		
Macrococcus bovicus		8.11	0.39		99	99		
200541	**Low discrimination**							**200541**
					NOV	OXI	YEP	
Staphylococcus cohnii subsp. cohnii		54.38	0.37	URE 1	99	1	12	
Kocuria varians		23.26	0.27	GAL 10	1	13	80	
Macrococcus equipersicus		11.48	0.18		99	99	99	
200542	**Low discrimination**							**200542**
					NOV	OXI		
Kocuria varians		59.07	0.38	TRE 7	1	13		
Staphylococcus warneri		20.36	0.35		1	1		
Macrococcus equipersicus		8.99	0.18		99	99		
200550	**Low discrimination**							**200550**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	PYR		
Macrococcus equipersicus		60.79	0.86		99	1		
Macrococcus bovicus		38.11	0.80		99	1		
Staphylococcus saprophyticus subsp. bovis		0.50	0.43		1	99		
200551	**Low discrimination**							**200551**
					OXI	YEP		
Macrococcus equipersicus		45.33	0.45	MNS 1	99	99		
Macrococcus bovicus		28.41	0.39		99	99		
Staphylococcus cohnii subsp. cohnii		14.38	0.39		1	12		
200552	**Low discrimination**							**200552**
					OXI			
Macrococcus equipersicus		35.68	0.45	LAC 1	99			
Macrococcus bovicus		22.37	0.39		99			
Staphylococcus saprophyticus subsp. saprophyticus		17.41	0.27		1			
200553	**Low discrimination**							**200553**
					NOV			
Staphylococcus cohnii subsp. urealyticum		76.79	0.44	GLR 99	99			
Aerococcus viridans		12.88	0.39		1			
Staphylococcus saprophyticus subsp. bovis		7.81	0.18		99			
200560	**Low discrimination**							**200560**
					NOV	OXI		
Kocuria varians		87.96	0.40	TRE 7	1	13		
Macrococcus equipersicus		10.75	0.18		99	99		
Macrococcus bovicus		1.29	-.02		99	99		
200562	**Acceptable identification**							**200562**
					PYR			
Kocuria varians		91.03	0.29	TRE 7	10			
Aerococcus viridans		8.98	0.22		95			
200563	**Low discrimination**							**200563**
					PYR			
Aerococcus viridans		72.57	0.30	URE 1	95			
Kocuria varians		27.43	0.07	ESL 90	10			
200570	**Low discrimination**							**200570**
					OXI			
Macrococcus equipersicus		58.19	0.45	XYL 1	99			
Macrococcus bovicus		36.47	0.39		99			
Staphylococcus kloosii		4.87	0.40		1			
200572	**Low discrimination**							**200572**
					OXI	NOV	YEP	
Staphylococcus kloosii		63.37	0.41	PHS 99	1	99	99	
Aerococcus viridans		25.14	0.32		1	1	1	
Macrococcus equipersicus		7.07	0.04		99	99	99	
200573	**Low discrimination**							**200573**
					NOV			
Aerococcus viridans		57.52	0.39	URE 1	1			
Staphylococcus xylosus		42.49	0.16	ESL 90	99			
200600	**Low discrimination**							**200600**
					OXI	NOV		
Staphylococcus warneri		28.36	0.56	MLT 95	1	1		
Macrococcus carouselicus		14.44	0.34		99	99		
Dermacoccus nishinomiyaensis		13.85	0.34		93	80		
200601	**Acceptable identification**							**200601**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	PYR	
Kocuria kristinae		92.65	0.41	VPT 90	99	93	
Kocuria varians		3.42	0.19	URE 10	13	10	
Staphylococcus warneri		1.79	0.21	MLT 90	1	8	
200602	**Low discrimination**						**200602**
Staphylococcus warneri		63.73	0.46	MLT 95			
Kocuria varians		24.18	0.30				
Staphylococcus hominis subsp. hominis		4.55	0.16				
200603	**Low discrimination**						**200603**
					OXI	PYR	
Kocuria kristinae		83.37	0.29	VPT 90	99	93	
Staphylococcus simulans		9.06	0.14	URE 10	1	70	
Kocuria varians		3.47	0.08	MLT 90	13	10	
200610	**Low discrimination**						**200610**
					OXI	NOV	
Macrococcus bovicus		61.09	0.68		99	99	
Macrococcus equipersicus		33.27	0.64		99	99	
Staphylococcus warneri		3.23	0.53		1	1	
200611	**Low discrimination**						**200611**
					NOV	PYR	
Macrococcus bovicus		49.32	0.27	MNS 1	99	1	
Macrococcus equipersicus		26.86	0.23		99	1	
Kocuria kristinae		15.68	0.13		1	93	
200612	**Low discrimination**						**200612**
					OXI	NOV	
Staphylococcus saprophyticus subsp. saprophyticus		49.76	0.29	MLT 99	1	99	
Staphylococcus warneri		23.72	0.43		1	1	
Macrococcus bovicus		13.60	0.27		99	99	
200613	**Low discrimination**						**200613**
					OXI		
Staphylococcus simulans		76.57	0.27	ARG 92	1		
Aerococcus viridans		12.35	0.27	BGA 90	1		
Kocuria kristinae		7.34	0.00	NIT 96	99		
200620	**Acceptable identification**						**200620**
Kocuria varians		93.98	0.32	SUC 12			
Staphylococcus warneri		6.03	0.14	TRE 7			
200633	**Low discrimination**						**200633**
					NOV		
Aerococcus viridans		70.53	0.27	URE 1	1		
Staphylococcus xylosus		29.47	-.01	ESL 90	99		
200640	**Low discrimination**						**200640**
					PYR		
Staphylococcus warneri		77.21	0.82		8		
Staphylococcus hominis subsp. hominis		8.16	0.56		1		
Staphylococcus petrasii subsp. pragensis		5.36	0.33		99		
200641	**Good identification**						**200641**
					OXI	PYR	
Kocuria kristinae		94.79	0.61	VPT 90	99	93	
Staphylococcus warneri		3.87	0.47	URE 10	1	8	
Staphylococcus hominis subsp. hominis		0.41	0.21		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
200642	**Low discrimination**							**200642**
					NOV			
Staphylococcus warneri		57.98	0.72		1			
Staphylococcus hominis subsp. hominis		21.58	0.58		1			
Staphylococcus saprophyticus subsp. saprophyticus		19.12	0.41		99			
200643	**Acceptable identification**							**200643**
					OXI	PYR		
Kocuria kristinae		90.78	0.48	VPT 90	99	93		
Staphylococcus warneri		4.94	0.37	URE 10	1	8		
Staphylococcus hominis subsp. hominis		1.84	0.23		1	1		
200650	**Low discrimination**							**200650**
					OXI	NOV		
Macrococcus bovicus		42.05	0.75		99	99		
Macrococcus equipersicus		26.54	0.73		99	99		
Staphylococcus warneri		18.10	0.79		1	1		
200651	**Low discrimination**							**200651**
					OXI	NOV	PYR	
Kocuria kristinae		26.19	0.32	VPT 90	99	1	93	
Macrococcus bovicus		21.35	0.34	URE 10	99	99	1	
Staphylococcus warneri		18.57	0.44	MAN 4	1	1	8	
200652	**Low discrimination**							**200652**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		89.64	0.70		99			
Staphylococcus warneri		8.20	0.69		1			
Staphylococcus hominis subsp. hominis		0.63	0.41		1			
200653	**Low discrimination**							**200653**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		55.18	0.29	MNS 1	1	99	7	
Kocuria kristinae		10.79	0.20		99	1	93	
Staphylococcus warneri		10.20	0.34		1	1	8	
200660	**Low discrimination**							**200660**
Staphylococcus warneri		70.86	0.41	XYL 1				
Kocuria varians		12.77	0.18					
Staphylococcus hominis subsp. hominis		7.49	0.15					
200662	**Low discrimination**							**200662**
					NOV			
Staphylococcus warneri		49.51	0.31	XYL 1	1			
Staphylococcus hominis subsp. hominis		18.42	0.16		1			
Staphylococcus saprophyticus subsp. saprophyticus		16.33	0.00		99			
200663	**Low discrimination**							**200663**
					OXI	NOV		
Aerococcus viridans		44.68	0.30	URE 1	1	1		
Kocuria kristinae		43.44	0.07	ESL 90	99	1		
Staphylococcus xylosus		11.88	-.03		1	99		
200670	**Low discrimination**							**200670**
					OXI	NOV		
Macrococcus bovicus		42.36	0.34	XYL 1	99	99		
Macrococcus equipersicus		26.74	0.32		99	99		
Staphylococcus warneri		18.23	0.38		1	1		
200672	**Low discrimination**							**200672**
					NOV	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. saprophyticus		84.15	0.29	XYL 1	99	7	
Staphylococcus warneri		7.70	0.28		1	8	
Aerococcus viridans		5.28	0.32		1	95	
200700	**Low discrimination**						**200700**
					NOV	OXI	
Kocuria varians		43.15	0.45	SUC 12	1	13	
Staphylococcus warneri		20.26	0.45	TRE 7	1	1	
Macrococcus carouselicus		13.99	0.27		99	99	
200701	**Low discrimination**						**200701**
					OXI	PYR	
Kocuria kristinae		84.43	0.31	VPT 90	99	93	
Kocuria varians		14.03	0.23	URE 10	13	10	
Staphylococcus warneri		1.55	0.10	MLT 90	1	8	
200702	**Low discrimination**						**200702**
Kocuria varians		59.15	0.34	SUC 12			
Staphylococcus warneri		32.82	0.35	TRE 7			
Staphylococcus hominis subsp. hominis		8.04	0.17				
200710	**Low discrimination**						**200710**
					OXI	NOV	
Macrococcus bovicus		61.93	0.60		99	99	
Macrococcus equipersicus		33.73	0.56		99	99	
Staphylococcus warneri		2.41	0.42		1	1	
200712	**Low discrimination**						**200712**
					OXI	NOV	
Staphylococcus warneri		33.57	0.33	MLT 95	1	1	
Macrococcus bovicus		26.12	0.19		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		19.39	0.07		1	99	
200713	**Low discrimination**						**200713**
					COA		
Aerococcus viridans		42.61	0.27	URE 1	1		
Staphylococcus simulans		22.98	0.05	ESL 90	1		
Staphylococcus intermedius		18.75	0.01		99		
200720	**Good identification**						**200720**
Kocuria varians		100.0	0.36	SUC 12			
200740	**Low discrimination**						**200740**
					OXI	NOV	
Staphylococcus warneri		65.46	0.72		1	1	
Staphylococcus hominis subsp. hominis		23.73	0.57		1	1	
Macrococcus equipersicus		4.97	0.39		99	99	
200741	**Acceptable identification**						**200741**
					OXI	PYR	
Kocuria kristinae		93.43	0.51	VPT 90	99	93	
Staphylococcus warneri		3.61	0.37	URE 10	1	8	
Staphylococcus hominis subsp. hominis		1.31	0.22		1	1	
200742	**Low discrimination**						**200742**
					NOV		
Staphylococcus hominis subsp. hominis		52.46	0.58	NIT 99	1		
Staphylococcus warneri		41.09	0.62		1		
Staphylococcus saprophyticus subsp. saprophyticus		3.73	0.19		99		
200743	**Low discrimination**						**200743**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	PYR		
Kocuria kristinae		84.39	0.39	VPT 90	99	93		
Staphylococcus hominis subsp. hominis		5.55	0.23	URE 10	1	1		
Staphylococcus warneri		4.35	0.27		1	8		
200750	**Low discrimination**							**200750**
					OXI	NOV		
Macrococcus bovicus		44.65	0.68		99	99		
Macrococcus equipersicus		28.18	0.65		99	99		
Staphylococcus warneri		14.16	0.69		1	1		
200751	**Low discrimination**							**200751**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		58.92	0.43	NIT 99	1	99	99	
Macrococcus bovicus		11.35	0.27		99	99	1	
Kocuria kristinae		10.83	0.23		99	1	93	
200752	**Low discrimination**							**200752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		60.57	0.48	GAL 8	99	7		
Staphylococcus warneri		20.13	0.59		1	8		
Staphylococcus saprophyticus subsp. bovis		9.98	0.43		99	99		
200753	**Low discrimination**							**200753**
					NOV			
Staphylococcus saprophyticus subsp. bovis		37.08	0.31	NIT 99	99			
Staphylococcus xylosus		18.67	0.21		99			
Aerococcus viridans		15.30	0.39		1			
200760	**Low discrimination**							**200760**
Staphylococcus warneri		45.08	0.30	XYL 1				
Kocuria varians		38.59	0.22					
Staphylococcus hominis subsp. hominis		16.34	0.16					
200770	**Low discrimination**							**200770**
					OXI	NOV		
Macrococcus bovicus		44.12	0.27	XYL 1	99	99		
Macrococcus equipersicus		27.84	0.24		99	99		
Staphylococcus warneri		13.99	0.28		1	1		
200771	**Acceptable identification**							**200771**
					NOV	PYR		
Staphylococcus xylosus		92.57	0.33	GLR 95	99	65		
Staphylococcus equorum		3.51	0.05	NIT 89	99	5		
Aerococcus viridans		2.09	0.20		1	95		
200772	**Low discrimination**							**200772**
					NOV	PYR		
Staphylococcus arlettae		45.82	0.31	URE 1	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		20.55	0.07	GLR 92	99	7		
Aerococcus viridans		14.82	0.32		1	95		
200773	**Low discrimination**							**200773**
					NOV	YEP		
Staphylococcus xylosus		81.26	0.43	GLR 95	99	40		
Staphylococcus arlettae		10.17	0.34	NIT 89	99	99		
Aerococcus viridans		5.79	0.39		1	1		
201000	**Low discrimination**							**201000**
					NOV	YEP		
Macrococcus equipersicus		40.30	0.65	MAN 95	99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		36.55	0.59		80	1		
Micrococcus sp.		19.60	0.44		1	41		
201002	**Low discrimination**							**201002**
					OXI	YEP	NOV	
Kocuria varians		37.89	0.30	ESL 1	13	80	1	
Macrococcus equipersicus		25.95	0.24		99	99	99	
Dermacoccus nishinomiyaensis		23.54	0.18		93	1	80	
201010	**Very good identification**							**201010**
Macrococcus equipersicus		97.37	0.91					
Macrococcus bovicus		2.47	0.57					
Macrococcus carouselicus		0.10	0.24					
201011	**Good identification**							**201011**
Macrococcus equipersicus		97.53	0.50	MNS 1				
Macrococcus bovicus		2.47	0.16					
201012	**Good identification**							**201012**
Macrococcus equipersicus		97.53	0.50	LAC 1				
Macrococcus bovicus		2.47	0.16					
201020	**Low discrimination**							**201020**
					OXI	YEP	NOV	
Kocuria varians		43.18	0.32	ESL 1	13	80	1	
Macrococcus equipersicus		23.74	0.24		99	99	99	
Dermacoccus nishinomiyaensis		21.53	0.18		93	1	80	
201030	**Good identification**							**201030**
Macrococcus equipersicus		96.94	0.50	XYL 1				
Macrococcus bovicus		2.45	0.16					
Staphylococcus vitulinus		0.61	-.03					
201040	**Very good identification**							**201040**
					NOV	OXI		
Macrococcus equipersicus		97.53	0.74	MAN 95	99	99		
Micrococcus sp.		1.32	0.21		1	89		
Kocuria varians		0.41	0.28		1	13		
201041	**Acceptable identification**							**201041**
					NOV	OXI		
Macrococcus equipersicus		93.99	0.33	MAN 95	99	99		
Kocuria varians		3.44	0.06	MNS 1	1	13		
Micrococcus sp.		2.57	-.14		1	89		
201042	**Low discrimination**							**201042**
					NOV	OXI		
Macrococcus equipersicus		89.40	0.33	MAN 95	99	99		
Kocuria varians		10.60	0.16	LAC 1	1	13		
201050	**Very good identification**							**201050**
Macrococcus equipersicus		97.86	1.00					
Macrococcus bovicus		2.14	0.64					
201051	**Very good identification**							**201051**
Macrococcus equipersicus		97.86	0.59	MNS 1				
Macrococcus bovicus		2.14	0.23					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
201052	**Very good identification**							**201052**
					OXI	NOV		
Macrococcus equipersicus		97.53	0.59	LAC 1	99	99		
Macrococcus bovicus		2.13	0.23		99	99		
Aerococcus viridans		0.34	0.25		1	1		
201060	**Low discrimination**							**201060**
					NOV	OXI		
Macrococcus equipersicus		87.13	0.33	MAN 95	99	99		
Kocuria varians		12.87	0.18	XYL 1	1	13		
201070	**Very good identification**							**201070**
Macrococcus equipersicus		97.86	0.59	XYL 1				
Macrococcus bovicus		2.14	0.23					
201073	**Good identification**							**201073**
Aerococcus viridans		100.0	0.33	URE 1				
201100	**Low discrimination**							**201100**
					OXI	YEP	NOV	
Macrococcus equipersicus		61.27	0.57	MAN 95	99	99	99	
Dermacoccus nishinomiyaensis		24.70	0.44		93	1	80	
Kocuria varians		11.21	0.45		13	80	1	
201102	**Low discrimination**							**201102**
					OXI	YEP	NOV	
Kocuria varians		78.46	0.33	ESL 1	13	80	1	
Macrococcus equipersicus		15.35	0.16		99	99	99	
Dermacoccus nishinomiyaensis		6.19	0.03		93	1	80	
201110	**Very good identification**							**201110**
Macrococcus equipersicus		97.41	0.84					
Macrococcus bovicus		2.47	0.49					
Macrococcus carouselicus		0.10	0.16					
201111	**Good identification**							**201111**
Macrococcus equipersicus		97.53	0.43	MNS 1				
Macrococcus bovicus		2.47	0.08					
201112	**Good identification**							**201112**
Macrococcus equipersicus		97.53	0.43	LAC 1				
Macrococcus bovicus		2.47	0.08					
201120	**Low discrimination**							**201120**
					OXI	YEP	NOV	
Kocuria varians		81.94	0.35	ESL 1	13	80	1	
Macrococcus equipersicus		12.87	0.16		99	99	99	
Dermacoccus nishinomiyaensis		5.19	0.03		93	1	80	
201130	**Good identification**							**201130**
Macrococcus equipersicus		97.53	0.43	XYL 1				
Macrococcus bovicus		2.47	0.08					
201140	**Very good identification**							**201140**
					NOV	OXI		
Macrococcus equipersicus		97.88	0.66	MAN 95	99	99		
Kocuria varians		1.45	0.31		1	13		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		0.41	0.16		99	99		
201150	**Very good identification**							**201150**
Macrococcus equipersicus		97.86	0.92					
Macrococcus bovicus		2.14	0.57					
201151	**Good identification**							**201151**
Macrococcus equipersicus		97.86	0.51	MNS 1				
Macrococcus bovicus		2.14	0.16					
201152	**Good identification**							**201152**
					OXI	NOV		
Macrococcus equipersicus		97.10	0.51	LAC 1	99	99		
Macrococcus bovicus		2.12	0.16		99	99		
Aerococcus viridans		0.78	0.25		1	1		
201153	**Low discrimination**							**201153**
					OXI	NOV		
Aerococcus viridans		64.92	0.33	URE 1	1	1		
Macrococcus equipersicus		35.08	0.10	TRE 95	99	99		
201170	**Good identification**							**201170**
Macrococcus equipersicus		97.86	0.51	XYL 1				
Macrococcus bovicus		2.14	0.16					
201173	**Good identification**							**201173**
Aerococcus viridans		100.0	0.33	URE 1				
201200	**Low discrimination**							**201200**
					YEP			
Macrococcus equipersicus		36.16	0.44	SUC 9	99			
Macrococcus carouselicus		22.05	0.34	MAN 95	67			
Dermacoccus nishinomiyaensis		21.16	0.34		1			
201210	**Good identification**							**201210**
Macrococcus equipersicus		90.25	0.70	SUC 9				
Macrococcus bovicus		5.78	0.44					
Staphylococcus vitulinus		2.13	0.29					
201211	**Acceptable identification**							**201211**
Macrococcus equipersicus		93.99	0.29	SUC 9				
Macrococcus bovicus		6.02	0.03	MNS 1				
201212	**Acceptable identification**							**201212**
Macrococcus equipersicus		93.99	0.29	SUC 9				
Macrococcus bovicus		6.02	0.03	LAC 1				
201230	**Low discrimination**							**201230**
Staphylococcus vitulinus		85.56	0.38	URE 1				
Macrococcus equipersicus		13.57	0.29	NIT 91				
Macrococcus bovicus		0.87	0.03					
201240	**Low discrimination**							**201240**
					NOV	OXI	YEP	
Macrococcus equipersicus		84.00	0.53	SUC 9	99	99	99	
Staphylococcus hominis subsp. novobiosepticus		12.86	0.29	MAN 95	99	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Micrococcus sp.		1.14	0.00		1	89	41		
201250	**Good identification**								**201250**
					OXI	NOV			
Macrococcus equipersicus		94.75	0.79	SUC 9	99	99			
Macrococcus bovicus		5.23	0.52		99	99			
Staphylococcus warneri		0.02	0.12		1	1			
201251	**Acceptable identification**								**201251**
					NOV	PYR			
Macrococcus equipersicus		93.70	0.38	SUC 9	99	1			
Macrococcus bovicus		5.18	0.11	MNS 1	99	1			
Kocuria kristinae		1.13	-.06		1	93			
201252	**Acceptable identification**								**201252**
					OXI	NOV			
Macrococcus equipersicus		91.75	0.38	SUC 9	99	99			
Macrococcus bovicus		5.07	0.11	LAC 1	99	99			
Aerococcus viridans		3.19	0.25		1	1			
201253	**Good identification**								**201253**
Aerococcus viridans		100.0	0.33	URE 1					
201270	**Acceptable identification**								**201270**
Macrococcus equipersicus		92.70	0.38	SUC 9					
Macrococcus bovicus		5.12	0.11	XYL 1					
Staphylococcus vitulinus		2.18	-.03						
201273	**Good identification**								**201273**
Aerococcus viridans		100.0	0.33	URE 1					
201300	**Low discrimination**								**201300**
					YEP				
Macrococcus equipersicus		46.48	0.37	SUC 9	99				
Macrococcus carouselicus		28.35	0.27	MAN 95	67				
Dermacoccus nishinomiyaensis		12.09	0.20		1				
201310	**Good identification**								**201310**
Macrococcus equipersicus		92.21	0.63	SUC 9					
Macrococcus bovicus		5.90	0.37						
Macrococcus carouselicus		1.89	0.23						
201340	**Low discrimination**								**201340**
					NOV	OXI	YEP		
Staphylococcus hominis subsp. novobiosepticus		50.80	0.39	VPT 90	99	1	1		
Macrococcus equipersicus		47.39	0.45	ESL 8	99	99	99		
Kocuria varians		0.97	0.13	NIT 90	1	13	80		
201341	**Low discrimination**								**201341**
					OXI	NOV	YEP	PYR	
Staphylococcus hominis subsp. novobiosepticus		83.43	0.27	VPT 90	1	99	1	1	
Kocuria kristinae		13.42	0.12	ESL 8	99	1	15	93	
Macrococcus equipersicus		3.15	0.04	NIT 90	99	99	99	1	
201342	**Good identification**								**201342**
					OXI	NOV	YEP		
Staphylococcus hominis subsp. novobiosepticus		95.96	0.33	VPT 90	1	99	1		
Macrococcus equipersicus		1.68	0.04	ESL 8	99	99	99		
Staphylococcus hominis subsp. hominis		1.40	0.07	NIT 90	1	1	49		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
201350	**Good identification**						**201350**
					OXI	YEP	
Micrococcus equipersicus		94.72	0.72	SUC 9	99	99	
Micrococcus bovicus		5.23	0.44		99	99	
Staphylococcus hominis subsp. novobiosepticus		0.05	-.02		1	1	
201351	**Acceptable identification**						**201351**
Micrococcus equipersicus		94.77	0.31	SUC 9			
Micrococcus bovicus		5.23	0.03	MNS 1			
201352	**Low discrimination**						**201352**
					OXI	NOV	
Micrococcus equipersicus		85.72	0.31	SUC 9	99	99	
Aerococcus viridans		6.94	0.25	LAC 1	1	1	
Micrococcus bovicus		4.73	0.03		99	99	
201353	**Good identification**						**201353**
Aerococcus viridans		100.0	0.33	URE 1			
201370	**Acceptable identification**						**201370**
Micrococcus equipersicus		94.77	0.31	SUC 9			
Micrococcus bovicus		5.23	0.03	XYL 1			
201400	**Low discrimination**						**201400**
					YEP		
Micrococcus equipersicus		67.05	0.65	MAN 95	99		
Micrococcus carouselicus		31.21	0.53		67		
Dermacoccus nishinomiyaensis		0.61	0.18		1		
201403	**Low discrimination**						**201403**
					NOV		
Aerococcus viridans		88.82	0.37	URE 1	1		
Staphylococcus cohnii subsp. urealyticum		11.18	0.07		99		
201410	**Good identification**						**201410**
Micrococcus equipersicus		94.90	0.91				
Micrococcus bovicus		3.60	0.60				
Micrococcus carouselicus		1.49	0.49				
201411	**Acceptable identification**						**201411**
Micrococcus equipersicus		93.08	0.50	MNS 1			
Micrococcus bovicus		3.54	0.19				
Micrococcus carouselicus		1.46	0.08				
201412	**Acceptable identification**						**201412**
					OXI	NOV	
Micrococcus equipersicus		90.65	0.50	LAC 1	99	99	
Aerococcus viridans		4.00	0.39		1	1	
Micrococcus bovicus		3.44	0.19		99	99	
201413	**Low discrimination**						**201413**
					OXI	NOV	
Aerococcus viridans		65.68	0.47	URE 1	1	1	
Staphylococcus cohnii subsp. urealyticum		27.87	0.28		1	99	
Micrococcus equipersicus		6.45	0.09		99	99	
201422	**Good identification**						**201422**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.29	URE 1				
201423	**Good identification**							**201423**
Aerococcus viridans		100.0	0.37	URE 1				
201430	**Acceptable identification**							**201430**
Macrococcus equipersicus		94.16	0.50	XYL 1				
Macrococcus bovicus		3.58	0.19					
Macrococcus carouselicus		1.48	0.08					
201431	**Low discrimination**							**201431**
					OXI	NOV		
Aerococcus viridans		53.09	0.27	URE 1	1	1		
Macrococcus equipersicus		46.91	0.09	LAC 90	99	99		
201432	**Low discrimination**							**201432**
					OXI	NOV		
Aerococcus viridans		81.36	0.39	URE 1	1	1		
Macrococcus equipersicus		18.64	0.09		99	99		
201433	**Good identification**							**201433**
Aerococcus viridans		100.0	0.47	URE 1				
201440	**Very good identification**							**201440**
Macrococcus equipersicus		99.00	0.74	MAN 95				
Macrococcus bovicus		0.62	0.27					
Macrococcus carouselicus		0.17	0.12					
201441	**Low discrimination**							**201441**
					OXI	NOV	PYR	
Macrococcus equipersicus		77.04	0.33	MAN 95	99	99	1	
Aerococcus viridans		8.25	0.30	MNS 1	1	1	95	
Kocuria kristinae		8.02	0.07		99	1	93	
201442	**Low discrimination**							**201442**
					OXI	NOV		
Macrococcus equipersicus		69.56	0.33	MAN 95	99	99		
Aerococcus viridans		28.74	0.42	LAC 1	1	1		
Staphylococcus piscifermentans		1.71	0.11		1	1		
201443	**Low discrimination**							**201443**
					OXI	NOV		
Aerococcus viridans		82.96	0.49	URE 1	1	1		
Staphylococcus cohnii subsp. urealyticum		14.80	0.23		1	99		
Kocuria kristinae		2.24	-.05		99	1		
201450	**Very good identification**							**201450**
					OXI			
Macrococcus equipersicus		96.77	1.00		99			
Macrococcus bovicus		3.17	0.68		99			
Staphylococcus kloosii		0.03	0.46		1			
201451	**Good identification**							**201451**
					OXI	NOV		
Macrococcus equipersicus		93.94	0.59	MNS 1	99	99		
Macrococcus bovicus		3.08	0.27		99	99		
Aerococcus viridans		1.59	0.39		1	1		
201452	**Low discrimination**							**201452**
					OXI	NOV	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus equipersicus		87.18	0.59	LAC 1	99	99	99	
Aerococcus viridans		5.69	0.51		1	1	1	
Staphylococcus kloosii		3.28	0.47		1	99	99	
201453	**Low discrimination**							**201453**
					OXI	NOV		
Aerococcus viridans		59.97	0.59	URE 1	1	1		
Staphylococcus cohnii subsp. urealyticum		36.05	0.43		1	99		
Macrococcus equipersicus		3.98	0.18		99	99		
201460	**Good identification**							**201460**
					OXI	NOV		
Macrococcus equipersicus		95.61	0.33	MAN 95	99	99		
Aerococcus viridans		4.39	0.22	XYL 1	1	1		
201461	**Good identification**							**201461**
Aerococcus viridans		100.0	0.30	URE 1				
201462	**Good identification**							**201462**
Aerococcus viridans		100.0	0.42	URE 1				
201463	**Good identification**							**201463**
Aerococcus viridans		100.0	0.49	URE 1				
201470	**Very good identification**							**201470**
					OXI			
Macrococcus equipersicus		95.36	0.59	XYL 1	99			
Macrococcus bovicus		3.13	0.27		99			
Staphylococcus kloosii		0.83	0.34		1			
201471	**Low discrimination**							**201471**
					OXI	NOV		
Aerococcus viridans		62.61	0.39	URE 1	1	1		
Macrococcus equipersicus		37.39	0.18	LAC 90	99	99		
201472	**Low discrimination**							**201472**
					OXI	NOV	YEP	
Aerococcus viridans		76.98	0.51	URE 1	1	1	1	
Macrococcus equipersicus		11.92	0.18		99	99	99	
Staphylococcus kloosii		11.11	0.35		1	99	99	
201473	**Very good identification**							**201473**
Aerococcus viridans		100.0	0.59	URE 1				
201500	**Low discrimination**							**201500**
					NOV	OXI		
Macrococcus equipersicus		67.02	0.57	MAN 95	99	99		
Macrococcus carouselicus		31.19	0.45		99	99		
Kocuria varians		0.92	0.22		1	13		
201503	**Good identification**							**201503**
Aerococcus viridans		100.0	0.37	URE 1				
201510	**Good identification**							**201510**
Macrococcus equipersicus		94.88	0.84					
Macrococcus bovicus		3.60	0.53					
Macrococcus carouselicus		1.49	0.41					
201511	**Acceptable identification**							**201511**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Macrococcus equipersicus		92.57	0.43	MNS 1	99	99	
Macrococcus bovicus		3.52	0.12		99	99	
Aerococcus viridans		2.47	0.27		1	1	
201512	**Low discrimination**						**201512**
					OXI	NOV	
Macrococcus equipersicus		84.69	0.43	LAC 1	99	99	
Aerococcus viridans		8.71	0.39		1	1	
Macrococcus bovicus		3.22	0.12		99	99	
201513	**Low discrimination**						**201513**
					OXI	NOV	
Aerococcus viridans		81.71	0.47	URE 1	1	1	
Staphylococcus cohnii subsp. urealyticum		14.86	0.20		1	99	
Macrococcus equipersicus		3.44	0.01		99	99	
201522	**Good identification**						**201522**
Aerococcus viridans		100.0	0.29	URE 1			
201523	**Good identification**						**201523**
Aerococcus viridans		100.0	0.37	URE 1			
201530	**Acceptable identification**						**201530**
Macrococcus equipersicus		93.89	0.43	XYL 1			
Macrococcus bovicus		3.57	0.12				
Macrococcus carouselicus		1.47	0.00				
201531	**Low discrimination**						**201531**
					OXI	NOV	
Aerococcus viridans		72.53	0.27	URE 1	1	1	
Macrococcus equipersicus		27.47	0.01	LAC 90	99	99	
201532	**Acceptable identification**						**201532**
					OXI	NOV	
Aerococcus viridans		91.06	0.39	URE 1	1	1	
Macrococcus equipersicus		8.94	0.01		99	99	
201533	**Good identification**						**201533**
Aerococcus viridans		100.0	0.47	URE 1			
201540	**Very good identification**						**201540**
Macrococcus equipersicus		98.93	0.66	MAN 95			
Macrococcus bovicus		0.62	0.19				
Macrococcus carouselicus		0.17	0.04				
201543	**Acceptable identification**						**201543**
					NOV		
Aerococcus viridans		92.90	0.49	URE 1	1		
Staphylococcus cohnii subsp. urealyticum		7.10	0.15		99		
201550	**Very good identification**						**201550**
					OXI		
Macrococcus equipersicus		96.65	0.92		99		
Macrococcus bovicus		3.17	0.60		99		
Staphylococcus kloosii		0.15	0.52		1		
201551	**Acceptable identification**						**201551**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
Macrococcus equipersicus		92.48	0.51	MNS 1	99	99			
Aerococcus viridans		3.65	0.39		1	1			
Macrococcus bovicus		3.03	0.19		99	99			
201552	**Low discrimination**								**201552**
					OXI	NOV	YEP		
Macrococcus equipersicus		73.77	0.51	LAC 1	99	99	99		
Staphylococcus kloosii		12.58	0.53		1	99	99		
Aerococcus viridans		11.23	0.51		1	1	1		
201553	**Low discrimination**								**201553**
					OXI	NOV			
Aerococcus viridans		77.76	0.59	URE 1	1	1			
Staphylococcus cohnii subsp. urealyticum		20.03	0.36		1	99			
Macrococcus equipersicus		2.21	0.10		99	99			
201561	**Good identification**								**201561**
Aerococcus viridans		100.0	0.30	URE 1					
201562	**Good identification**								**201562**
Aerococcus viridans		100.0	0.42	URE 1					
201563	**Good identification**								**201563**
Aerococcus viridans		100.0	0.49	URE 1					
201570	**Acceptable identification**								**201570**
					OXI				
Macrococcus equipersicus		91.82	0.51	XYL 1	99				
Staphylococcus kloosii		3.61	0.40		1				
Macrococcus bovicus		3.01	0.19		99				
201571	**Low discrimination**								**201571**
					OXI	NOV	YEP	PYR	
Aerococcus viridans		69.31	0.39	URE 1	1	1	1	95	
Macrococcus equipersicus		17.74	0.10	LAC 90	99	99	99	1	
Staphylococcus equorum		12.95	0.05		1	99	1	5	
201572	**Low discrimination**								**201572**
					OXI	NOV	YEP		
Aerococcus viridans		74.27	0.51	URE 1	1	1	1		
Staphylococcus kloosii		20.80	0.41		1	99	99		
Macrococcus equipersicus		4.93	0.10		99	99	99		
201573	**Good identification**								**201573**
					NOV	PYR			
Aerococcus viridans		93.12	0.59	URE 1	1	95			
Staphylococcus equorum		4.97	0.14		99	5			
Staphylococcus xylosus		1.91	0.04		99	65			
201600	**Good identification**								**201600**
Macrococcus carouselicus		90.21	0.59	URE 1					
Macrococcus equipersicus		9.44	0.44						
Macrococcus bovicus		0.17	0.07						
201601	**Low discrimination**								**201601**
					NOV	PYR			
Kocuria kristinae		73.51	0.29	VPT 90	1	93			
Macrococcus carouselicus		22.43	0.18	URE 10	99	1			
Macrococcus equipersicus		2.35	0.03	MLT 90	99	1			

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
201610	**Low discrimination**							**201610**
Micrococcus equipersicus		69.85	0.70	SUC 9				
Micrococcus carouselicus		22.46	0.55					
Micrococcus bovicus		6.71	0.48					
201611	**Low discrimination**							**201611**
					OXI	NOV		
Micrococcus equipersicus		60.88	0.29	SUC 9	99	99		
Micrococcus carouselicus		19.58	0.14	MNS 1	99	99		
Aerococcus viridans		7.04	0.27		1	1		
201612	**Low discrimination**							**201612**
					OXI	NOV		
Micrococcus equipersicus		53.67	0.29	SUC 9	99	99		
Aerococcus viridans		23.93	0.39	LAC 1	1	1		
Micrococcus carouselicus		17.26	0.14		99	99		
201613	**Good identification**							**201613**
Aerococcus viridans		100.0	0.47	URE 1				
201622	**Good identification**							**201622**
Aerococcus viridans		100.0	0.29	URE 1				
201623	**Good identification**							**201623**
Aerococcus viridans		100.0	0.37	URE 1				
201630	**Low discrimination**							**201630**
Staphylococcus vitulinus		70.73	0.33	URE 1				
Micrococcus equipersicus		19.95	0.29	NIT 91				
Micrococcus carouselicus		6.41	0.14					
201631	**Low discrimination**							**201631**
					OXI	NOV	PYR	
Aerococcus viridans		49.18	0.27	URE 1	1	1	95	
Staphylococcus equorum		35.58	0.05	LAC 90	1	99	5	
Staphylococcus vitulinus		15.24	-.08		99	99	1	
201632	**Acceptable identification**							**201632**
					OXI	NOV		
Aerococcus viridans		92.56	0.39	URE 1	1	1		
Staphylococcus vitulinus		7.44	-.08		99	99		
201633	**Low discrimination**							**201633**
					NOV	PYR		
Aerococcus viridans		82.87	0.47	URE 1	1	95		
Staphylococcus equorum		17.13	0.14		99	5		
201640	**Low discrimination**							**201640**
					OXI	NOV		
Micrococcus equipersicus		85.52	0.53	SUC 9	99	99		
Staphylococcus warneri		7.27	0.41	MAN 95	1	1		
Micrococcus carouselicus		3.05	0.18		99	99		
201641	**Good identification**							**201641**
					OXI	NOV	PYR	
Kocuria kristinae		98.04	0.48	VPT 90	99	1	93	
Aerococcus viridans		1.02	0.30	URE 10	1	1	95	
Micrococcus equipersicus		0.94	0.12		99	99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
201642	**Low discrimination**							**201642**
					PYR			
Aerococcus viridans		39.45	0.42	URE 1	95			
Staphylococcus warneri		26.50	0.31		8			
Staphylococcus hominis subsp. hominis		9.86	0.16		1			
201643	**Low discrimination**							**201643**
					OXI			
Kocuria kristinae		72.78	0.36	VPT 90	99			
Aerococcus viridans		27.22	0.49	URE 10	1			
201650	**Good identification**							**201650**
					OXI	NOV		
Macrococcus equipersicus		91.69	0.79	SUC 9	99	99		
Macrococcus bovicus		7.60	0.56		99	99		
Staphylococcus warneri		0.30	0.38		1	1		
201651	**Low discrimination**							**201651**
					OXI	NOV	PYR	
Macrococcus equipersicus		64.97	0.38	SUC 9	99	99	1	
Kocuria kristinae		14.85	0.20	MNS 1	99	1	93	
Aerococcus viridans		11.11	0.39		1	1	95	
201652	**Low discrimination**							**201652**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		38.25	0.29	ESL 1	1	99	7	
Macrococcus equipersicus		32.72	0.38		99	99	1	
Aerococcus viridans		21.58	0.51		1	1	95	
201653	**Good identification**							**201653**
					OXI	NOV		
Aerococcus viridans		94.72	0.59	URE 1	1	1		
Kocuria kristinae		3.52	0.08		99	1		
Staphylococcus cohnii subsp. urealyticum		1.76	0.12		1	99		
201661	**Low discrimination**							**201661**
					OXI			
Aerococcus viridans		50.70	0.30	URE 1	1			
Kocuria kristinae		49.30	0.07	LAC 90	99			
201662	**Good identification**							**201662**
Aerococcus viridans		100.0	0.42	URE 1				
201663	**Good identification**							**201663**
					OXI			
Aerococcus viridans		97.37	0.49	URE 1	1			
Kocuria kristinae		2.63	-.05		99			
201670	**Low discrimination**							**201670**
					OXI	NOV		
Macrococcus equipersicus		85.52	0.38	SUC 9	99	99		
Macrococcus bovicus		7.09	0.15	XYL 1	99	99		
Aerococcus viridans		6.27	0.32		1	1		
201672	**Acceptable identification**							**201672**
					NOV	YEP		
Aerococcus viridans		94.28	0.51	URE 1	1	1		
Staphylococcus arlettae		5.72	0.15		99	99		
201673	**Low discrimination**							**201673**
					NOV	PYR		
Aerococcus viridans		62.67	0.59	URE 1	1	95		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus equorum		32.75	0.34		99	5		
Staphylococcus xylosus		2.42	0.10		99	65		
201700	**Acceptable identification**							**201700**
					OXI	NOV		
Macrococcus carouselicus		90.01	0.51	URE 1	99	99		
Macrococcus equipersicus		9.42	0.37		99	99		
Staphylococcus piscifermentans		0.22	0.14		1	1		
201703	**Low discrimination**							**201703**
					OXI			
Aerococcus viridans		71.63	0.37	URE 1	1			
Kocuria kristinae		28.37	0.07		99			
201710	**Low discrimination**							**201710**
Macrococcus equipersicus		70.49	0.63	SUC 9				
Macrococcus carouselicus		22.66	0.47					
Macrococcus bovicus		6.77	0.40					
201712	**Low discrimination**							**201712**
					OXI	NOV		
Aerococcus viridans		41.00	0.39	URE 1	1	1		
Macrococcus equipersicus		39.42	0.22		99	99		
Macrococcus carouselicus		12.67	0.06		99	99		
201713	**Acceptable identification**							**201713**
					OXI	NOV		
Aerococcus viridans		91.12	0.47	URE 1	1	1		
Staphylococcus lentus		8.88	0.12		99	99		
201722	**Good identification**							**201722**
Aerococcus viridans		100.0	0.29	URE 1				
201723	**Good identification**							**201723**
Aerococcus viridans		100.0	0.37	URE 1				
201732	**Good identification**							**201732**
Aerococcus viridans		100.0	0.39	URE 1				
201733	**Low discrimination**							**201733**
					NOV	PYR		
Staphylococcus equorum		63.25	0.34	GLR 90	99	5		
Aerococcus viridans		30.26	0.47	NIT 92	1	95		
Staphylococcus xylosus		3.55	0.07	MLT 91	99	65		
201740	**Low discrimination**							**201740**
					OXI	NOV		
Macrococcus equipersicus		85.05	0.45	SUC 9	99	99		
Staphylococcus warneri		5.33	0.30	MAN 95	1	1		
Macrococcus carouselicus		3.04	0.10		99	99		
201741	**Good identification**							**201741**
					OXI	NOV	PYR	
Kocuria kristinae		95.10	0.39	VPT 90	99	1	93	
Aerococcus viridans		2.96	0.30	URE 10	1	1	95	
Macrococcus equipersicus		1.17	0.04		99	99	1	
201742	**Low discrimination**							**201742**
					PYR			
Aerococcus viridans		56.24	0.42	URE 1	95			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		15.23	0.17		1			
Staphylococcus warneri		11.93	0.21		8			
201743	**Low discrimination**							**201743**
					OXI			
Aerococcus viridans		52.87	0.49	URE 1	1			
Kocuria kristinae		47.13	0.26		99			
201750	**Good identification**							**201750**
					OXI	NOV		
Macrococcus equipersicus		91.57	0.72	SUC 9	99	99		
Macrococcus bovicus		7.59	0.48		99	99		
Staphylococcus warneri		0.22	0.28		1	1		
201751	**Low discrimination**							**201751**
					OXI	NOV		
Macrococcus equipersicus		47.11	0.31	SUC 9	99	99		
Aerococcus viridans		18.80	0.39	MNS 1	1	1		
Staphylococcus sciuri		12.08	0.19		99	99		
201752	**Low discrimination**							**201752**
					OXI	NOV	PYR	
Aerococcus viridans		48.45	0.51	URE 1	1	1	95	
Macrococcus equipersicus		31.48	0.31		99	99	1	
Staphylococcus saprophyticus subsp. saprophyticus		7.47	0.07		1	99	7	
201753	**Low discrimination**							**201753**
					OXI	NOV	PYR	
Aerococcus viridans		85.57	0.59	URE 1	1	1	95	
Staphylococcus equorum		4.57	0.14		1	99	5	
Staphylococcus lentus		4.23	0.18		99	99	6	
201761	**Low discrimination**							**201761**
					OXI	NOV	PYR	
Aerococcus viridans		47.16	0.30	URE 1	1	1	95	
Staphylococcus equorum		26.43	0.05	LAC 90	1	99	5	
Kocuria kristinae		15.28	-.03		99	1	93	
201762	**Good identification**							**201762**
Aerococcus viridans		100.0	0.42	URE 1				
201763	**Low discrimination**							**201763**
					NOV	PYR		
Aerococcus viridans		80.99	0.49	URE 1	1	95		
Staphylococcus equorum		12.97	0.14		99	5		
Staphylococcus xylosus		6.05	0.06		99	65		
201770	**Low discrimination**							**201770**
					OXI	NOV		
Macrococcus equipersicus		73.11	0.31	SUC 9	99	99		
Aerococcus viridans		12.50	0.32	XYL 1	1	1		
Macrococcus bovicus		6.06	0.07		99	99		
201771	**Low discrimination**							**201771**
					NOV	PYR		
Staphylococcus equorum		85.84	0.47	GLR 90	99	5		
Staphylococcus xylosus		5.72	0.21	NIT 92	99	65		
Aerococcus viridans		4.64	0.39		1	95		
201772	**Low discrimination**							**201772**
					NOV	YEP	PYR	
Aerococcus viridans		66.86	0.51	URE 1	1	1	95	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus arlettae		22.97	0.31		99	99	1	
Staphylococcus equorum		8.33	0.14		99	1	5	
201773	**Low discrimination**							**201773**
					NOV	PYR		
Staphylococcus equorum		75.66	0.55	GLR 90	99	5		
Aerococcus viridans		14.32	0.59	NIT 92	1	95		
Staphylococcus xylosus		5.58	0.30		99	65		
202000	**Low discrimination**							**202000**
					OXI			
Dermacoccus nishinomiyaensis		63.62	0.95		93			
Kocuria varians		31.84	0.96		13			
Micrococcus sp.		3.19	0.59		89			
202001	**Low discrimination**							**202001**
					OXI			
Kocuria varians		77.82	0.75	MNS 8	13			
Dermacoccus nishinomiyaensis		18.06	0.54		93			
Kocuria rosea		1.95	0.33		35			
202002	**Good identification**							**202002**
					OXI			
Kocuria varians		92.86	0.85		13			
Dermacoccus nishinomiyaensis		6.65	0.54		93			
Micrococcus sp.		0.33	0.18		89			
202003	**Very good identification**							**202003**
					OXI			
Kocuria varians		99.18	0.63	MNS 8	13			
Dermacoccus nishinomiyaensis		0.82	0.13		93			
202010	**Low discrimination**							**202010**
					OXI	YEP	NOV	
Dermacoccus nishinomiyaensis		34.44	0.54	MAN 1	93	1	80	
Macrococcus bovicus		32.48	0.57		99	99	99	
Kocuria varians		17.24	0.55		13	80	1	
202011	**Low discrimination**							**202011**
Staphylococcus capitis subsp. capitis		74.82	0.42	URE 1				
Kocuria varians		14.25	0.34					
Kocuria rosea		3.92	0.13					
202012	**Low discrimination**							**202012**
					OXI	YEP	NOV	
Kocuria varians		85.62	0.44	MAN 1	13	80	1	
Dermacoccus nishinomiyaensis		6.13	0.13		93	1	80	
Macrococcus bovicus		5.78	0.16		99	99	99	
202020	**Good identification**							**202020**
					OXI			
Kocuria varians		90.03	0.87		13			
Dermacoccus nishinomiyaensis		5.17	0.54		93			
Kocuria rosea		4.54	0.51		35			
202021	**Very good identification**							**202021**
					OXI			
Kocuria varians		96.42	0.65	MNS 8	13			
Kocuria rosea		2.94	0.25		35			
Dermacoccus nishinomiyaensis		0.64	0.13		93			
202022	**Very good identification**							**202022**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI		
Kocuria varians		99.62	0.76		13		
Dermacoccus nishinomiyaensis		0.21	0.13		93		
Kocuria rosea		0.18	0.10		35		
202023	**Very good identification**						**202023**
Kocuria varians		100.0	0.54	MNS 8			
202030	**Low discrimination**						**202030**
					OXI		
Kocuria varians		57.93	0.46	MAN 1	13		
Kocuria rosea		32.10	0.32		35		
Dermacoccus nishinomiyaensis		3.33	0.13		93		
202032	**Good identification**						**202032**
Kocuria varians		100.0	0.35	MAN 1			
202040	**Low discrimination**						**202040**
					OXI		
Kocuria varians		86.35	0.83		13		
Dermacoccus nishinomiyaensis		7.94	0.54		93		
Micrococcus sp.		2.97	0.36		89		
202041	**Very good identification**						**202041**
					OXI	NOV	
Kocuria varians		95.34	0.61	MNS 8	13	1	
Staphylococcus hominis subsp. novobiosepticus		2.12	0.17		1	99	
Dermacoccus nishinomiyaensis		1.02	0.13		93	80	
202042	**Good identification**						**202042**
					NOV		
Kocuria varians		92.80	0.72		1		
Staphylococcus hominis subsp. hominis		5.23	0.47		1		
Staphylococcus hominis subsp. novobiosepticus		1.37	0.24		99		
202043	**Acceptable identification**						**202043**
					NOV		
Kocuria varians		94.26	0.50	MNS 8	1		
Staphylococcus hominis subsp. novobiosepticus		3.99	0.12		99		
Staphylococcus hominis subsp. hominis		1.25	0.12		1		
202050	**Low discrimination**						**202050**
					NOV	OXI	
Macrococcus bovicus		76.60	0.64	NIT 10	99	99	
Macrococcus equipersicus		16.66	0.52		99	99	
Kocuria varians		3.82	0.42		1	13	
202052	**Low discrimination**						**202052**
					NOV	OXI	
Kocuria varians		32.42	0.30	MAN 1	1	13	
Staphylococcus hominis subsp. hominis		27.03	0.30		1	1	
Macrococcus bovicus		23.26	0.23		99	99	
202060	**Very good identification**						**202060**
					OXI		
Kocuria varians		99.37	0.74		13		
Dermacoccus nishinomiyaensis		0.26	0.13		93		
Kocuria rosea		0.23	0.10		35		
202061	**Good identification**						**202061**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.52	MNS 8				
202062	**Very good identification**							**202062**
Kocuria varians		99.84	0.62					
Staphylococcus hominis subsp. hominis		0.16	0.06					
202063	**Good identification**							**202063**
Kocuria varians		100.0	0.40	MNS 8				
202070	**Low discrimination**							**202070**
					NOV	OXI		
Kocuria varians		58.79	0.32	MAN 1	1	13		
Macrococcus bovicus		33.85	0.23		99	99		
Macrococcus equipersicus		7.36	0.11		99	99		
202100	**Low discrimination**							**202100**
					OXI			
Kocuria varians		79.69	1.00		13			
Dermacoccus nishinomiyaensis		20.22	0.80		93			
Micrococcus sp.		0.05	0.18		89			
202101	**Very good identification**							**202101**
					OXI			
Kocuria varians		97.14	0.78	MNS 8	13			
Dermacoccus nishinomiyaensis		2.86	0.39		93			
202102	**Very good identification**							**202102**
					OXI			
Kocuria varians		99.08	0.89		13			
Dermacoccus nishinomiyaensis		0.90	0.39		93			
Staphylococcus hominis subsp. hominis		0.01	0.07		1			
202103	**Very good identification**							**202103**
					OXI			
Kocuria varians		99.90	0.67	MNS 8	13			
Dermacoccus nishinomiyaensis		0.11	-.02		93			
202110	**Low discrimination**							**202110**
					OXI	YEP	NOV	
Kocuria varians		52.67	0.59	MAN 1	13	80	1	
Macrococcus bovicus		28.36	0.49		99	99	99	
Dermacoccus nishinomiyaensis		13.37	0.39		93	1	80	
202111	**Low discrimination**							**202111**
					NOV	OXI		
Kocuria varians		88.70	0.37	MAN 1	1	13		
Macrococcus bovicus		5.55	0.08	MNS 8	99	99		
Staphylococcus capitis subsp. capitis		3.14	0.01		1	1		
202112	**Good identification**							**202112**
					OXI	YEP	NOV	
Kocuria varians		97.24	0.48	MAN 1	13	80	1	
Macrococcus bovicus		1.88	0.08		99	99	99	
Dermacoccus nishinomiyaensis		0.88	-.02		93	1	80	
202120	**Very good identification**							**202120**
					OXI			
Kocuria varians		99.24	0.91		13			
Dermacoccus nishinomiyaensis		0.72	0.39		93			
Kocuria rosea		0.03	0.10		35			
202121	**Very good identification**							**202121**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Kocuria varians		99.92	0.69	MNS 8	13			
Dermacoccus nishinomiyaensis		0.09	-.02		93			
202122	**Very good identification**							**202122**
					OXI			
Kocuria varians		99.97	0.79		13			
Dermacoccus nishinomiyaensis		0.03	-.02		93			
202123	**Very good identification**							**202123**
Kocuria varians		100.0	0.58	MNS 8				
202130	**Good identification**							**202130**
					OXI	YEP	NOV	
Kocuria varians		97.77	0.50	MAN 1	13	80	1	
Macrococcus bovicus		1.51	0.08		99	99	99	
Dermacoccus nishinomiyaensis		0.71	-.02		93	1	80	
202131	**Good identification**							**202131**
Kocuria varians		100.0	0.28	MAN 1				
202132	**Good identification**							**202132**
Kocuria varians		100.0	0.38	MAN 1				
202140	**Very good identification**							**202140**
					OXI	NOV		
Kocuria varians		96.26	0.86		13	1		
Staphylococcus hominis subsp. novobiosepticus		1.49	0.39		1	99		
Dermacoccus nishinomiyaensis		1.12	0.39		93	80		
202141	**Very good identification**							**202141**
					NOV			
Kocuria varians		95.30	0.65	MNS 8	1			
Staphylococcus hominis subsp. novobiosepticus		4.23	0.27		99			
Staphylococcus hominis subsp. hominis		0.22	0.12		1			
202142	**Good identification**							**202142**
					NOV			
Kocuria varians		93.35	0.75		1			
Staphylococcus hominis subsp. hominis		3.80	0.48		1			
Staphylococcus hominis subsp. novobiosepticus		2.75	0.34		99			
202143	**Good identification**							**202143**
					NOV			
Kocuria varians		90.17	0.53	MNS 8	1			
Staphylococcus hominis subsp. novobiosepticus		7.64	0.21		99			
Staphylococcus epidermidis		1.33	0.14		1			
202150	**Low discrimination**							**202150**
					OXI	PYR		
Macrococcus bovicus		62.48	0.57	NIT 10	99	1		
Macrococcus equipersicus		13.59	0.45		99	1		
Staphylococcus saprophyticus subsp. bovis		11.05	0.43		1	99		
202151	**Low discrimination**							**202151**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		61.66	0.31	TRE 99	99	99	1	
Kocuria varians		21.19	0.24		1	10	13	
Macrococcus bovicus		14.08	0.16		99	1	99	
202152	**Low discrimination**							**202152**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Kocuria varians		35.99	0.34	MAN 1	1	10	
Staphylococcus saprophyticus subsp. bovis		32.30	0.31		99	99	
Staphylococcus hominis subsp. hominis		21.67	0.31		1	1	
202160	**Very good identification**						**202160**
					NOV		
Kocuria varians		99.89	0.77		1		
Staphylococcus hominis subsp. novobiosepticus		0.04	-.02		99		
Staphylococcus hominis subsp. hominis		0.03	0.05		1		
202161	**Very good identification**						**202161**
Kocuria varians		100.0	0.55	MNS 8			
202162	**Very good identification**						**202162**
					NOV		
Kocuria varians		99.80	0.66		1		
Staphylococcus hominis subsp. hominis		0.12	0.07		1		
Staphylococcus hominis subsp. novobiosepticus		0.09	-.07		99		
202163	**Good identification**						**202163**
Kocuria varians		100.0	0.44	MNS 8			
202170	**Low discrimination**						**202170**
					NOV	OXI	
Kocuria varians		81.34	0.36	MAN 1	1	13	
Macrococcus bovicus		13.38	0.16		99	99	
Macrococcus equipersicus		2.91	0.04		99	99	
202200	**Low discrimination**						**202200**
					OXI		
Kocuria varians		48.08	0.79	SUC 12	13		
Dermacoccus nishinomiyaensis		44.97	0.70		93		
Micrococcus sp.		3.50	0.38		89		
202201	**Low discrimination**						**202201**
					OXI		
Kocuria varians		80.37	0.57	SUC 12	13		
Dermacoccus nishinomiyaensis		8.73	0.29	MNS 8	93		
Staphylococcus capitis subsp. capitis		4.37	0.24		1		
202202	**Low discrimination**						**202202**
					OXI		
Kocuria varians		86.59	0.67	SUC 12	13		
Staphylococcus hominis subsp. hominis		7.85	0.47		1		
Dermacoccus nishinomiyaensis		2.90	0.29		93		
202203	**Low discrimination**						**202203**
					NOV		
Kocuria varians		85.75	0.45	SUC 12	1		
Staphylococcus hominis subsp. novobiosepticus		5.85	0.12	MNS 8	99		
Staphylococcus simulans		4.36	0.16		1		
202210	**Low discrimination**						**202210**
					NOV	OXI	
Staphylococcus vitulinus		42.04	0.41	URE 1	99	99	
Macrococcus bovicus		30.55	0.44		99	99	
Kocuria varians		8.84	0.38		1	13	
202211	**Good identification**						**202211**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus capitis subsp. capitis		96.02	0.47	URE 1				
Kocuria varians		1.34	0.16					
Staphylococcus capitis subsp. ureolyticus		1.05	-.02					
202212	**Low discrimination**							**202212**
Staphylococcus hominis subsp. hominis		41.62	0.30	MAN 13				
Kocuria varians		31.01	0.26	MLT 99				
Staphylococcus simulans		16.21	0.18					
202220	**Good identification**							**202220**
					OXI			
Kocuria varians		93.08	0.69	SUC 12	13			
Kocuria rosea		3.82	0.32		35			
Dermacoccus nishinomiyaensis		2.50	0.29		93			
202221	**Good identification**							**202221**
Kocuria varians		97.58	0.47	SUC 12				
Kocuria rosea		2.42	0.05	MNS 8				
202222	**Very good identification**							**202222**
Kocuria varians		99.74	0.58	SUC 12				
Staphylococcus hominis subsp. hominis		0.26	0.06					
202223	**Good identification**							**202223**
Kocuria varians		100.0	0.36	SUC 12				
202230	**Good identification**							**202230**
					NOV	OXI		
Staphylococcus vitulinus		95.93	0.50	URE 1	99	99		
Kocuria varians		2.62	0.28		1	13		
Kocuria rosea		1.18	0.12		1	35		
202240	**Low discrimination**							**202240**
					NOV			
Staphylococcus hominis subsp. hominis		59.17	0.87		1			
Staphylococcus hominis subsp. novobiosepticus		33.72	0.71		99			
Kocuria varians		6.02	0.65		1			
202241	**Low discrimination**							**202241**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		81.27	0.58	VPT 90	99			
Staphylococcus hominis subsp. hominis		11.64	0.52		1			
Kocuria varians		5.05	0.43		1			
202242	**Low discrimination**							**202242**
					NOV			
Staphylococcus hominis subsp. hominis		77.44	0.88		1			
Staphylococcus hominis subsp. novobiosepticus		20.24	0.65		99			
Kocuria varians		1.89	0.54		1			
202243	**Low discrimination**							**202243**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		67.82	0.53	VPT 90	99			
Staphylococcus hominis subsp. hominis		21.18	0.54		1			
Staphylococcus epidermidis		6.26	0.39		1			
202250	**Low discrimination**							**202250**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		69.89	0.70	MAN 13	1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		17.65	0.52		99	1	99	
Staphylococcus saprophyticus subsp. bovis		5.35	0.43		1	99	99	
202252	**Very good identification**							**202252**
					PYR	NOV		
Staphylococcus hominis subsp. hominis		95.04	0.71	MAN 13	1	1		
Staphylococcus hominis subsp. novobiosepticus		1.68	0.24		1	99		
Staphylococcus saprophyticus subsp. bovis		1.55	0.31		99	99		
202253	**Low discrimination**							**202253**
					COA			
Staphylococcus capitis subsp. ureolyticus		86.64	0.44	ARG 99	1			
Staphylococcus hominis subsp. hominis		5.51	0.37		1			
Staphylococcus delphini		3.96	0.18		99			
202260	**Low discrimination**							**202260**
					NOV			
Kocuria varians		69.27	0.56	SUC 12	1			
Staphylococcus hominis subsp. hominis		19.57	0.46		1			
Staphylococcus hominis subsp. novobiosepticus		11.16	0.30		99			
202261	**Low discrimination**							**202261**
					NOV			
Kocuria varians		65.39	0.34	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		30.27	0.17	MNS 8	99			
Staphylococcus hominis subsp. hominis		4.34	0.11		1			
202262	**Low discrimination**							**202262**
					NOV			
Staphylococcus hominis subsp. hominis		47.36	0.47	XYL 1	1			
Kocuria varians		40.27	0.44		1			
Staphylococcus hominis subsp. novobiosepticus		12.38	0.24		99			
202270	**Low discrimination**							**202270**
					OXI	NOV		
Staphylococcus hominis subsp. hominis		50.74	0.29	MAN 13	1	1		
Staphylococcus vitulinus		20.43	0.09	XYL 1	99	99		
Macrococcus bovicus		12.81	0.11		99	99		
202272	**Acceptable identification**							**202272**
Staphylococcus hominis subsp. hominis		94.57	0.30	MAN 13				
Kocuria varians		5.44	0.03	XYL 1				
202300	**Low discrimination**							**202300**
					OXI	NOV		
Kocuria varians		86.18	0.82	SUC 12	13	1		
Dermacoccus nishinomiyaensis		10.24	0.55		93	80		
Staphylococcus hominis subsp. novobiosepticus		2.14	0.39		1	99		
202301	**Good identification**							**202301**
					OXI	NOV		
Kocuria varians		91.38	0.60	SUC 12	13	1		
Staphylococcus hominis subsp. novobiosepticus		6.53	0.27	MNS 8	1	99		
Dermacoccus nishinomiyaensis		1.26	0.14		93	80		
202302	**Low discrimination**							**202302**
					NOV			
Kocuria varians		89.42	0.71	SUC 12	1			
Staphylococcus hominis subsp. hominis		5.86	0.48		1			
Staphylococcus hominis subsp. novobiosepticus		4.24	0.34		99			
202303	**Low discrimination**							**202303**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Kocuria varians		77.36	0.49	SUC 12	1		
Staphylococcus hominis subsp. novobiosepticus		10.55	0.21	MNS 8	99		
Staphylococcus hyicus		4.73	0.14		1		
202310	**Low discrimination**						**202310**
					NOV	OXI	
Kocuria varians		39.27	0.41	SUC 12	1	13	
Macrococcus bovicus		38.76	0.37	MAN 1	99	99	
Staphylococcus hominis subsp. hominis		9.15	0.30		1	1	
202312	**Low discrimination**						**202312**
					NOV	PYR	
Kocuria varians		48.20	0.30	SUC 12	1	10	
Staphylococcus hominis subsp. hominis		46.71	0.31	MAN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		2.81	0.02		99	99	
202320	**Very good identification**						**202320**
					OXI	NOV	
Kocuria varians		99.54	0.73	SUC 12	13	1	
Dermacoccus nishinomiyaensis		0.34	0.14		93	80	
Staphylococcus hominis subsp. novobiosepticus		0.07	-.02		1	99	
202321	**Good identification**						**202321**
Kocuria varians		100.0	0.51	SUC 12			
202322	**Very good identification**						**202322**
					NOV		
Kocuria varians		99.68	0.62	SUC 12	1		
Staphylococcus hominis subsp. hominis		0.19	0.07		1		
Staphylococcus hominis subsp. novobiosepticus		0.14	-.07		99		
202323	**Good identification**						**202323**
Kocuria varians		100.0	0.40	SUC 12			
202330	**Low discrimination**						**202330**
					NOV	OXI	
Kocuria varians		80.24	0.32	SUC 12	1	13	
Staphylococcus vitulinus		19.76	0.09	MAN 1	99	99	
202340	**Low discrimination**						**202340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		57.83	0.80	VPT 90	99		
Staphylococcus hominis subsp. hominis		36.64	0.88		1		
Kocuria varians		5.16	0.69		1		
202341	**Low discrimination**						**202341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		89.81	0.68	VPT 90	99		
Staphylococcus hominis subsp. hominis		4.65	0.53		1		
Kocuria varians		2.79	0.47		1		
202342	**Low discrimination**						**202342**
					NOV		
Staphylococcus hominis subsp. hominis		56.33	0.89		1		
Staphylococcus hominis subsp. novobiosepticus		40.77	0.75		99		
Kocuria varians		1.91	0.57		1		
202343	**Low discrimination**						**202343**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		76.48	0.62	VPT 90	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		13.34	0.55		1			
Staphylococcus hominis subsp. hominis		8.63	0.54		1			
202350	**Low discrimination**							**202350**
					PYR	NOV		
Staphylococcus hominis subsp. hominis		65.92	0.71	MAN 13	1	1		
Staphylococcus saprophyticus subsp. bovis		18.64	0.55		99	99		
Staphylococcus hominis subsp. novobiosepticus		7.03	0.39		1	99		
202351	**Low discrimination**							**202351**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		58.21	0.43	TRE 99	99	99		
Staphylococcus hominis subsp. novobiosepticus		21.96	0.27		1	99		
Staphylococcus hominis subsp. hominis		16.80	0.36		1	1		
202352	**Low discrimination**							**202352**
					PYR	NOV		
Staphylococcus hominis subsp. hominis		89.59	0.72	MAN 13	1	1		
Staphylococcus saprophyticus subsp. bovis		5.40	0.43		99	99		
Staphylococcus hominis subsp. novobiosepticus		4.38	0.34		1	99		
202353	**Low discrimination**							**202353**
					COA	NOV	PYR	
Staphylococcus hominis subsp. hominis		28.42	0.37	MAN 13	1	1	1	
Staphylococcus saprophyticus subsp. bovis		20.96	0.31	MNS 2	1	99	99	
Staphylococcus delphini		18.82	0.18		99	1	15	
202360	**Low discrimination**							**202360**
					NOV			
Kocuria varians		65.52	0.59	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		21.10	0.39		99			
Staphylococcus hominis subsp. hominis		13.37	0.47		1			
202361	**Low discrimination**							**202361**
					NOV			
Kocuria varians		49.31	0.37	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		45.66	0.27	MNS 8	99			
Staphylococcus hominis subsp. hominis		2.36	0.12		1			
202362	**Low discrimination**							**202362**
					NOV			
Kocuria varians		40.34	0.48	SUC 12	1			
Staphylococcus hominis subsp. hominis		34.27	0.48		1			
Staphylococcus hominis subsp. novobiosepticus		24.81	0.34		99			
202370	**Low discrimination**							**202370**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		54.93	0.30	MAN 13	1	1		
Kocuria varians		18.19	0.18	XYL 1	1	10		
Staphylococcus saprophyticus subsp. bovis		15.53	0.14		99	99		
202372	**Low discrimination**							**202372**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		84.12	0.31	MAN 13	1	1		
Kocuria varians		6.70	0.07	XYL 1	1	10		
Staphylococcus saprophyticus subsp. bovis		5.07	0.02		99	99		
202373	**Good identification**							**202373**
Staphylococcus xylosus		100.0	0.33	GLR 95				
202400	**Low discrimination**							**202400**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		73.00	0.73	TRE 7	13		
Dermacoccus nishinomiyaensis		19.57	0.54		93		
Staphylococcus auricularis		3.88	0.46		1		
202401	**Low discrimination**						**202401**
					NOV		
Kocuria varians		88.27	0.51	TRE 7	1		
Staphylococcus auricularis		6.00	0.26	MNS 8	1		
Staphylococcus cohnii subsp. cohnii		2.98	0.23		99		
202402	**Good identification**						**202402**
Kocuria varians		94.96	0.62	TRE 7			
Staphylococcus auricularis		1.99	0.26				
Staphylococcus piscifermentans		1.30	0.32				
202403	**Acceptable identification**						**202403**
Kocuria varians		90.94	0.40	TRE 7			
Staphylococcus simulans		6.63	0.14	MNS 8			
Staphylococcus auricularis		2.43	0.07				
202410	**Low discrimination**						**202410**
					OXI	PYR	
Macrococcus bovicus		82.71	0.60	NIT 10	99	1	
Macrococcus equipersicus		10.35	0.43		99	1	
Staphylococcus saprophyticus subsp. bovis		2.44	0.31		1	99	
202413	**Low discrimination**						**202413**
					NOV		
Staphylococcus simulans		62.25	0.27	ARG 92	1		
Staphylococcus cohnii subsp. urealyticum		17.39	0.21	BGA 90	99		
Staphylococcus felis		14.27	0.04	SUC 96	1		
202420	**Very good identification**						**202420**
					OXI		
Kocuria varians		98.43	0.64	TRE 7	13		
Dermacoccus nishinomiyaensis		0.76	0.13		93		
Kocuria rosea		0.67	0.10		35		
202421	**Good identification**						**202421**
Kocuria varians		100.0	0.42	TRE 7			
202422	**Good identification**						**202422**
Kocuria varians		100.0	0.53	TRE 7			
202423	**Good identification**						**202423**
Kocuria varians		100.0	0.31	TRE 7			
202440	**Low discrimination**						**202440**
Kocuria varians		40.87	0.60	TRE 7			
Staphylococcus hominis subsp. hominis		23.28	0.56				
Staphylococcus auricularis		23.11	0.53				
202441	**Low discrimination**						**202441**
					NOV		
Kocuria varians		39.35	0.38	TRE 7	1		
Staphylococcus auricularis		28.43	0.34	MNS 8	1		
Staphylococcus cohnii subsp. cohnii		19.15	0.34		99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
202442	**Low discrimination**					**202442**
Staphylococcus hominis subsp. hominis		61.66	0.58	SUC 99		
Kocuria varians		26.01	0.48			
Staphylococcus auricularis		5.79	0.34			
202443	**Low discrimination**					**202443**
Kocuria varians		42.96	0.27	TRE 7		
Staphylococcus hominis subsp. hominis		23.90	0.23	MNS 8		
Staphylococcus auricularis		12.22	0.14			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
202450	**Low discrimination**							**202450**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		50.91	0.72		1	99		
Macrococcus bovicus		40.70	0.68		99	1		
Macrococcus equipersicus		5.90	0.52		99	1		
202451	**Good identification**							**202451**
					OXI	PYR	YEP	
Staphylococcus saprophyticus subsp. bovis		92.04	0.59		1	99	80	
Staphylococcus cohnii subsp. cohnii		3.93	0.36		1	1	12	
Macrococcus bovicus		2.97	0.27		99	1	99	
202452	**Low discrimination**							**202452**
					OXI	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		86.02	0.59		1	99	99	
Staphylococcus hominis subsp. hominis		6.82	0.41		1	1	1	
Macrococcus bovicus		2.78	0.27		99	1	99	
202453	**Low discrimination**							**202453**
					NOV			
Staphylococcus saprophyticus subsp. bovis		82.20	0.47		99			
Staphylococcus cohnii subsp. urealyticum		13.45	0.36		99			
Staphylococcus simulans		3.82	0.19		1			
202460	**Good identification**							**202460**
Kocuria varians		96.84	0.50	TRE 7				
Staphylococcus hominis subsp. hominis		1.59	0.15					
Staphylococcus auricularis		1.57	0.12					
202461	**Good identification**							**202461**
Kocuria varians		100.0	0.29	TRE 7				
202462	**Acceptable identification**							**202462**
Kocuria varians		93.62	0.39	TRE 7				
Staphylococcus hominis subsp. hominis		6.38	0.16					
202470	**Low discrimination**							**202470**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		50.38	0.31	XYL 1	1	99		
Macrococcus bovicus		40.27	0.27		99	1		
Macrococcus equipersicus		5.84	0.11		99	1		
202500	**Very good identification**							**202500**
					OXI			
Kocuria varians		95.77	0.77	TRE 7	13			
Dermacoccus nishinomiyaensis		3.26	0.39		93			
Staphylococcus auricularis		0.38	0.26		1			
202501	**Very good identification**							**202501**
					OXI			
Kocuria varians		98.86	0.55	TRE 7	13			
Staphylococcus auricularis		0.50	0.07	MNS 8	1			
Dermacoccus nishinomiyaensis		0.39	-.02		93			
202502	**Very good identification**							**202502**
Kocuria varians		98.01	0.66	TRE 7				
Staphylococcus piscifermentans		1.35	0.36					
Staphylococcus hominis subsp. hominis		0.37	0.17					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
202503	**Good identification**							**202503**
Kocuria varians		95.51	0.44	TRE 7				
Staphylococcus felis		2.13	-.02	MNS 8				
Staphylococcus hyicus		1.23	-.05					
202510	**Low discrimination**							**202510**
					OXI	PYR		
Macrococcus bovicus		66.17	0.53	NIT 10	99	1		
Staphylococcus saprophyticus subsp. bovis		18.21	0.43		1	99		
Macrococcus equipersicus		8.28	0.36		99	1		
202511	**Low discrimination**							**202511**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		76.82	0.31	MLT 99	99	99	1	
Macrococcus bovicus		11.28	0.12		99	1	99	
Kocuria varians		9.05	0.14		1	10	13	
202512	**Low discrimination**							**202512**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		57.10	0.31	MLT 99	99	99		
Kocuria varians		21.82	0.25		1	10		
Staphylococcus piscifermentans		12.70	0.28		1	15		
202520	**Very good identification**							**202520**
					OXI			
Kocuria varians		99.90	0.68	TRE 7	13			
Dermacoccus nishinomiyaensis		0.10	-.02		93			
202521	**Good identification**							**202521**
Kocuria varians		100.0	0.46	TRE 7				
202522	**Very good identification**							**202522**
Kocuria varians		100.0	0.56	TRE 7				
202523	**Good identification**							**202523**
Kocuria varians		100.0	0.34	TRE 7				
202530	**Low discrimination**							**202530**
					NOV	PYR	OXI	
Kocuria varians		71.77	0.26	TRE 7	1	10	13	
Macrococcus bovicus		22.14	0.12	MAN 1	99	1	99	
Staphylococcus saprophyticus subsp. bovis		6.09	0.02		99	99	1	
202540	**Low discrimination**							**202540**
					NOV	PYR		
Kocuria varians		59.95	0.63	TRE 7	1	10		
Staphylococcus hominis subsp. hominis		24.66	0.57		1	1		
Staphylococcus saprophyticus subsp. bovis		8.14	0.43		99	99		
202541	**Low discrimination**							**202541**
					NOV	PYR		
Kocuria varians		63.43	0.42	TRE 7	1	10		
Staphylococcus saprophyticus subsp. bovis		24.77	0.31	MNS 8	99	99		
Staphylococcus hominis subsp. hominis		6.12	0.22		1	1		
202542	**Low discrimination**							**202542**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		56.61	0.58	SUC 99	1	1		
Kocuria varians		33.06	0.52		1	10		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. bovis		3.98	0.31		99	99		
202543	**Low discrimination**							**202543**
					NOV	PYR		
Kocuria varians		53.73	0.30	TRE 7	1	10		
Staphylococcus hominis subsp. hominis		21.59	0.23	MNS 8	1	1		
Staphylococcus saprophyticus subsp. bovis		18.60	0.18		99	99		
202550	**Good identification**							**202550**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		90.46	0.84		1	99		
Macrococcus bovicus		7.75	0.60		99	1		
Macrococcus equipersicus		1.12	0.45		99	1		
202551	**Very good identification**							**202551**
					OXI	PYR	YEP	
Staphylococcus saprophyticus subsp. bovis		99.33	0.72		1	99	80	
Macrococcus bovicus		0.34	0.19		99	1	99	
Staphylococcus cohnii subsp. cohnii		0.12	0.16		1	1	12	
202552	**Very good identification**							**202552**
					OXI	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		97.05	0.72		1	99	99	
Staphylococcus hominis subsp. hominis		2.08	0.41		1	1	1	
Macrococcus bovicus		0.34	0.19		99	1	99	
202553	**Very good identification**							**202553**
Staphylococcus saprophyticus subsp. bovis		97.28	0.59					
Staphylococcus cohnii subsp. urealyticum		1.71	0.29					
Staphylococcus xylosus		0.84	0.13					
202560	**Very good identification**							**202560**
					NOV	PYR		
Kocuria varians		98.45	0.54	TRE 7	1	10		
Staphylococcus hominis subsp. hominis		1.16	0.16		1	1		
Staphylococcus saprophyticus subsp. bovis		0.39	0.02		99	99		
202561	**Acceptable identification**							**202561**
					NOV			
Kocuria varians		94.21	0.32	TRE 7	1			
Staphylococcus xylosus		5.79	0.01	MNS 8	99			
202562	**Good identification**							**202562**
Kocuria varians		95.31	0.43	TRE 7				
Staphylococcus hominis subsp. hominis		4.69	0.17					
202570	**Low discrimination**							**202570**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		88.95	0.43	XYL 1	99	99	1	
Macrococcus bovicus		7.62	0.19		99	1	99	
Kocuria varians		2.32	0.13		1	10	13	
202571	**Low discrimination**							**202571**
Staphylococcus saprophyticus subsp. bovis		53.59	0.31	XYL 1				
Staphylococcus xylosus		46.42	0.26					
202572	**Good identification**							**202572**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		97.14	0.31	XYL 1	99	99		
Kocuria varians		2.86	0.02		1	10		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
202573	**Low discrimination**						**202573**
					COA	NOV	
Staphylococcus xylosus		74.39	0.35	GLR 95	1	99	
Staphylococcus lutrae		18.06	0.18	SUC 95	99	1	
Staphylococcus saprophyticus subsp. bovis		7.54	0.18		1	99	
202600	**Low discrimination**						**202600**
Kocuria varians		37.37	0.55	SUC 12			
Staphylococcus hominis subsp. hominis		34.27	0.56	TRE 7			
Staphylococcus auricularis		13.45	0.45				
202601	**Low discrimination**						**202601**
					OXI	PYR	
Kocuria varians		44.45	0.34	SUC 12	13	10	
Staphylococcus auricularis		20.45	0.26	TRE 7	1	40	
Kocuria kristinae		18.56	0.18	MNS 8	99	93	
202602	**Low discrimination**						**202602**
Staphylococcus hominis subsp. hominis		67.16	0.58	MLT 99			
Kocuria varians		17.60	0.44				
Staphylococcus simulans		5.87	0.32				
202603	**Low discrimination**						**202603**
					YEP		
Staphylococcus simulans		71.00	0.43	ARG 92	1		
Staphylococcus chromogenes		7.90	0.20	BGA 90	95		
Staphylococcus hyicus		6.75	0.14		1		
202610	**Low discrimination**						**202610**
					OXI	PYR	
Macrococcus bovicus		39.23	0.48	NIT 10	99	1	
Staphylococcus vitulinus		20.25	0.36		99	1	
Staphylococcus saprophyticus subsp. bovis		18.51	0.43		1	99	
202611	**Low discrimination**						**202611**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		64.02	0.31	MLT 99	99	99	
Staphylococcus capitis subsp. capitis		13.29	0.13		1	1	
Staphylococcus simulans		9.93	0.14		1	70	
202612	**Low discrimination**						**202612**
					NOV	PYR	
Staphylococcus simulans		52.71	0.44	ARG 92	1	70	
Staphylococcus hominis subsp. hominis		22.55	0.41	BGA 90	1	1	
Staphylococcus saprophyticus subsp. bovis		11.50	0.31		99	99	
202613	**Very good identification**						**202613**
					NOV		
Staphylococcus simulans		95.41	0.55	ARG 92	1		
Staphylococcus felis		1.85	0.11	BGA 90	1		
Staphylococcus saprophyticus subsp. bovis		1.55	0.18		99		
202620	**Low discrimination**						**202620**
					NOV	OXI	
Kocuria varians		69.36	0.46	SUC 12	1	13	
Staphylococcus muscae		26.16	0.18	TRE 7	1	1	
Staphylococcus vitulinus		1.93	0.04		99	99	
202622	**Acceptable identification**						**202622**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		90.11	0.35	SUC 12			
Staphylococcus hominis subsp. hominis		9.89	0.16	TRE 7			
202630	**Good identification**						**202630**
					NOV	OXI	
Staphylococcus vitulinus		98.59	0.45	URE 1	99	99	
Macrococcus bovicus		0.71	0.07		99	99	
Kocuria varians		0.36	0.05		1	13	
202640	**Good identification**						**202640**
Staphylococcus hominis subsp. hominis		94.33	0.97				
Staphylococcus warneri		3.87	0.74				
Staphylococcus auricularis		0.87	0.53				
202641	**Low discrimination**						**202641**
					OXI	PYR	
Staphylococcus hominis subsp. hominis		73.12	0.62	MNS 2	1	1	
Kocuria kristinae		12.90	0.38		99	93	
Staphylococcus auricularis		3.68	0.33		1	40	
202642	**Very good identification**						**202642**
Staphylococcus hominis subsp. hominis		98.44	0.99				
Staphylococcus warneri		1.15	0.64				
Staphylococcus auricularis		0.09	0.33				
202643	**Low discrimination**						**202643**
					OXI	PYR	
Staphylococcus hominis subsp. hominis		84.73	0.64	MNS 2	1	1	
Staphylococcus simulans		5.51	0.35		1	70	
Kocuria kristinae		3.18	0.25		99	93	
202650	**Low discrimination**						**202650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		63.98	0.84		99	99	
Staphylococcus hominis subsp. hominis		26.72	0.80		1	1	
Staphylococcus warneri		5.31	0.71		1	8	
202651	**Very good identification**						**202651**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		95.43	0.72		99	99	
Staphylococcus hominis subsp. hominis		3.25	0.45		1	1	
Staphylococcus warneri		0.65	0.37		1	8	
202652	**Low discrimination**						**202652**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		62.50	0.82	MAN 13	1	1	
Staphylococcus saprophyticus subsp. bovis		31.87	0.72		99	99	
Staphylococcus warneri		3.52	0.62		1	8	
202653	**Low discrimination**						**202653**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		67.57	0.59		99	99	
Staphylococcus simulans		18.83	0.48		1	70	
Staphylococcus hominis subsp. hominis		10.82	0.47		1	1	
202660	**Low discrimination**						**202660**
Staphylococcus hominis subsp. hominis		87.88	0.56	XYL 1			
Kocuria varians		7.39	0.33				
Staphylococcus warneri		3.60	0.33				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
202662	**Very good identification**							**202662**
Staphylococcus hominis subsp. hominis		96.91	0.58	XYL 1				
Kocuria varians		1.96	0.21					
Staphylococcus warneri		1.13	0.23					
202670	**Low discrimination**							**202670**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		63.27	0.43	XYL 1	99	99		
Staphylococcus hominis subsp. hominis		26.43	0.39		1	1		
Staphylococcus warneri		5.25	0.30		1	8		
202671	**Low discrimination**							**202671**
					PYR			
Staphylococcus xylosus		58.12	0.31	GLR 95	65			
Staphylococcus saprophyticus subsp. bovis		35.71	0.31	GAL 91	99			
Staphylococcus equorum		3.13	0.07		5			
202672	**Low discrimination**							**202672**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		62.89	0.41	MAN 13	1	1		
Staphylococcus saprophyticus subsp. bovis		32.07	0.31	XYL 1	99	99		
Staphylococcus warneri		3.55	0.20		1	8		
202673	**Low discrimination**							**202673**
					PYR			
Staphylococcus xylosus		88.79	0.41	GLR 95	65			
Staphylococcus saprophyticus subsp. bovis		4.79	0.18	GAL 91	99			
Staphylococcus equorum		4.32	0.15		5			
202700	**Low discrimination**							**202700**
Kocuria varians		55.68	0.59	SUC 12				
Staphylococcus hominis subsp. hominis		36.88	0.57	TRE 7				
Staphylococcus warneri		2.30	0.38					
202701	**Low discrimination**							**202701**
					YEP			
Kocuria varians		51.74	0.37	SUC 12	80			
Staphylococcus chromogenes		13.88	0.20	TRE 7	95			
Staphylococcus hyicus		11.86	0.14	MNS 8	1			
202702	**Low discrimination**							**202702**
Staphylococcus hominis subsp. hominis		68.64	0.58	MLT 99				
Kocuria varians		24.90	0.48					
Staphylococcus piscifermentans		3.76	0.39					
202703	**Low discrimination**							**202703**
					YEP			
Staphylococcus hyicus		44.06	0.36	ARG 96	1			
Staphylococcus chromogenes		40.37	0.39	PHS 92	95			
Kocuria varians		4.72	0.26		80			
202710	**Low discrimination**							**202710**
					OXI	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		70.58	0.55	MLT 99	1	99	99	
Macrococcus bovicus		16.03	0.40		99	1	99	
Staphylococcus hominis subsp. hominis		7.98	0.40		1	1	1	
202711	**Acceptable identification**							**202711**
					NOV			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		90.20	0.43	MLT 99	99		
Staphylococcus chromogenes		2.72	0.08		1		
Staphylococcus xylosus		2.43	0.07		99		
202712	**Low discrimination**						**202712**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		56.12	0.43	MLT 99	99	99	
Staphylococcus hominis subsp. hominis		29.81	0.41		1	1	
Staphylococcus simulans		5.59	0.22		1	70	
202713	**Low discrimination**						**202713**
					NOV	YEP	
Staphylococcus simulans		24.27	0.33	ARG 92	1	1	
Staphylococcus chromogenes		19.70	0.28	BGA 90	1	95	
Staphylococcus saprophyticus subsp. bovis		18.18	0.31	GAL 8	99	80	
202720	**Good identification**						**202720**
Kocuria varians		98.13	0.50	SUC 12			
Staphylococcus hominis subsp. hominis		1.87	0.16	TRE 7			
202721	**Low discrimination**						**202721**
					NOV		
Kocuria varians		87.61	0.28	SUC 12	1		
Staphylococcus xylosus		12.40	0.04	TRE 7	99		
202722	**Acceptable identification**						**202722**
Kocuria varians		92.66	0.38	SUC 12			
Staphylococcus hominis subsp. hominis		7.34	0.17	TRE 7			
202731	**Acceptable identification**						**202731**
					PYR		
Staphylococcus xylosus		90.63	0.29	GLR 95	65		
Staphylococcus equorum		6.41	0.07	MLT 93	5		
Staphylococcus saprophyticus subsp. bovis		2.96	0.02		99		
202733	**Acceptable identification**						**202733**
					OXI		
Staphylococcus xylosus		91.78	0.38	GLR 95	1		
Staphylococcus equorum		5.87	0.15	MLT 93	1		
Staphylococcus lentus		2.35	0.12		99		
202740	**Very good identification**						**202740**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		96.55	0.98		1	1	
Staphylococcus saprophyticus subsp. bovis		1.29	0.55		99	99	
Staphylococcus warneri		1.15	0.64		1	8	
202741	**Low discrimination**						**202741**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		73.24	0.63	MNS 2	1	1	
Staphylococcus saprophyticus subsp. bovis		11.97	0.43		99	99	
Staphylococcus hominis subsp. novobiosepticus		4.52	0.27		1	99	
202742	**Very good identification**						**202742**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		98.96	0.99		1	1	
Staphylococcus warneri		0.34	0.54		1	8	
Staphylococcus saprophyticus subsp. bovis		0.28	0.43		99	99	
202743	**Low discrimination**						**202743**
					NOV	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		78.98	0.65	MNS 2	1	1	
Staphylococcus chromogenes		10.56	0.39		1	45	
Staphylococcus saprophyticus subsp. bovis		2.75	0.31		99	99	
202750	**Low discrimination**						**202750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		88.84	0.96		99	99	
Staphylococcus hominis subsp. hominis		10.05	0.81		1	1	
Staphylococcus warneri		0.58	0.61		1	8	
202751	**Very good identification**						**202751**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.53	0.84		99	99	
Staphylococcus hominis subsp. hominis		0.91	0.46		1	1	
Staphylococcus xylosus		0.36	0.30		99	65	
202752	**Low discrimination**						**202752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		64.83	0.84		99	99	
Staphylococcus hominis subsp. hominis		34.44	0.82		1	1	
Staphylococcus warneri		0.57	0.51		1	8	
202753	**Low discrimination**						**202753**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		89.66	0.72		99	99	
Staphylococcus hominis subsp. hominis		3.89	0.48		1	1	
Staphylococcus xylosus		3.69	0.40		99	65	
202760	**Low discrimination**						**202760**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		87.20	0.57	XYL 1	1	1	
Kocuria varians		10.16	0.36		1	10	
Staphylococcus saprophyticus subsp. bovis		1.16	0.14		99	99	
202761	**Low discrimination**						**202761**
					NOV		
Staphylococcus xylosus		71.91	0.27	GLR 95	99		
Staphylococcus hominis subsp. hominis		16.92	0.22	MAN 94	1		
Kocuria varians		8.40	0.14		1		
202762	**Very good identification**						**202762**
Staphylococcus hominis subsp. hominis		96.28	0.58	XYL 1			
Kocuria varians		2.69	0.25				
Staphylococcus warneri		0.33	0.13				
202763	**Low discrimination**						**202763**
					COA	NOV	
Staphylococcus xylosus		81.65	0.37	GLR 95	1	99	
Staphylococcus lutrae		8.42	0.12	MAN 94	99	1	
Staphylococcus hominis subsp. hominis		7.93	0.23		1	1	
202770	**Low discrimination**						**202770**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		88.07	0.55	XYL 1	99	99	
Staphylococcus hominis subsp. hominis		9.96	0.40		1	1	
Staphylococcus xylosus		0.92	0.11		99	65	
202771	**Low discrimination**						**202771**
					PYR		
Staphylococcus xylosus		76.13	0.52	GLR 95	65		
Staphylococcus saprophyticus subsp. bovis		18.50	0.43		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus equorum		4.10	0.27		5			
202772	**Low discrimination**							**202772**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		60.31	0.43	XYL 1	99	99		
Staphylococcus hominis subsp. hominis		32.03	0.41		1	1		
Staphylococcus xylosus		7.13	0.20		99	65		
202773	**Good identification**							**202773**
					PYR			
Staphylococcus xylosus		91.57	0.61	GLR 95	65			
Staphylococcus equorum		4.46	0.36		5			
Staphylococcus saprophyticus subsp. bovis		1.96	0.31		99			
203000	**Low discrimination**							**203000**
					OXI			
Dermacoccus nishinomiyaensis		62.70	0.54	ESL 1	93			
Kocuria varians		31.38	0.55		13			
Micrococcus sp.		3.15	0.18		89			
203001	**Low discrimination**							**203001**
					OXI			
Kocuria varians		81.16	0.34	ESL 1	13			
Dermacoccus nishinomiyaensis		18.84	0.13	MNS 8	93			
203002	**Acceptable identification**							**203002**
					OXI			
Kocuria varians		92.52	0.44	ESL 1	13			
Dermacoccus nishinomiyaensis		6.62	0.13		93			
Staphylococcus piscifermentans		0.86	0.11		1			
203010	**Low discrimination**							**203010**
					YEP			
Macrococcus equipersicus		73.81	0.50	NIT 1	99			
Macrococcus bovicus		20.56	0.37		99			
Dermacoccus nishinomiyaensis		1.98	0.13		1			
203020	**Acceptable identification**							**203020**
					OXI			
Kocuria varians		90.27	0.46	ESL 1	13			
Dermacoccus nishinomiyaensis		5.19	0.13		93			
Kocuria rosea		4.55	0.10		35			
203022	**Good identification**							**203022**
Kocuria varians		100.0	0.35	ESL 1				
203040	**Low discrimination**							**203040**
					OXI	YEP	NOV	
Kocuria varians		58.27	0.42	ESL 1	13	80	1	
Macrococcus equipersicus		28.40	0.33		99	99	99	
Dermacoccus nishinomiyaensis		5.36	0.13		93	1	80	
203042	**Low discrimination**							**203042**
					NOV			
Kocuria varians		81.60	0.30	ESL 1	1			
Staphylococcus hominis subsp. novobiosepticus		10.35	0.02		99			
Staphylococcus hominis subsp. hominis		4.60	0.06		1			
203050	**Low discrimination**							**203050**
Macrococcus equipersicus		80.62	0.59	NIT 1				
Macrococcus bovicus		19.38	0.45					

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
203060	**Good identification**							**203060**
Kocuria varians		100.0	0.32	ESL 1				
203100	**Low discrimination**							**203100**
					OXI	YEP	NOV	
Kocuria varians		78.81	0.59	ESL 1	13	80	1	
Dermacoccus nishinomiyaensis		20.00	0.39		93	1	80	
Macrococcus equipersicus		0.89	0.16		99	99	99	
203101	**Good identification**							**203101**
					OXI			
Kocuria varians		97.14	0.37	ESL 1	13			
Dermacoccus nishinomiyaensis		2.86	-.02	MNS 8	93			
203102	**Good identification**							**203102**
					OXI			
Kocuria varians		98.20	0.48	ESL 1	13			
Staphylococcus piscifermentans		0.91	0.14		1			
Dermacoccus nishinomiyaensis		0.89	-.02		93			
203110	**Low discrimination**							**203110**
					NOV	OXI		
Macrococcus equipersicus		74.77	0.43	NIT 1	99	99		
Macrococcus bovicus		20.82	0.30		99	99		
Kocuria varians		3.52	0.18		1	13		
203120	**Good identification**							**203120**
					OXI			
Kocuria varians		99.28	0.50	ESL 1	13			
Dermacoccus nishinomiyaensis		0.72	-.02		93			
203121	**Good identification**							**203121**
Kocuria varians		100.0	0.28	ESL 1				
203122	**Good identification**							**203122**
Kocuria varians		100.0	0.38	ESL 1				
203140	**Low discrimination**							**203140**
					NOV	OXI	YEP	
Kocuria varians		76.84	0.45	ESL 1	1	13	80	
Macrococcus equipersicus		10.70	0.25		99	99	99	
Staphylococcus hominis subsp. novobiosepticus		10.21	0.17		99	1	1	
203142	**Low discrimination**							**203142**
					NOV			
Kocuria varians		74.81	0.34	ESL 1	1			
Staphylococcus hominis subsp. novobiosepticus		18.97	0.12		99			
Staphylococcus piscifermentans		3.17	0.14		1			
203150	**Low discrimination**							**203150**
					NOV	OXI		
Macrococcus equipersicus		80.13	0.51	NIT 1	99	99		
Macrococcus bovicus		19.26	0.37		99	99		
Kocuria varians		0.31	0.04		1	13		
203160	**Good identification**							**203160**
Kocuria varians		100.0	0.36	ESL 1				
203200	**Low discrimination**							**203200**
					OXI	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		40.67	0.38	ESL 1	13	1		
Dermacoccus nishinomiyaensis		38.04	0.29	SUC 12	93	80		
Staphylococcus vitulinus		5.28	0.09		99	99		
203210	**Acceptable identification**							**203210**
Staphylococcus vitulinus		92.78	0.50	URE 1				
Macrococcus equipersicus		3.94	0.29					
Macrococcus bovicus		2.77	0.25					
203220	**Low discrimination**							**203220**
					NOV	OXI		
Kocuria varians		50.03	0.28	ESL 1	1	13		
Staphylococcus vitulinus		49.97	0.18	SUC 12	99	99		
203230	**Very good identification**							**203230**
Staphylococcus vitulinus		100.0	0.59	URE 1				
203240	**Low discrimination**							**203240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		81.06	0.49	VPT 90	99			
Staphylococcus hominis subsp. hominis		16.52	0.46	ESL 8	1			
Kocuria varians		1.68	0.24		1			
203241	**Good identification**							**203241**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		96.53	0.36	VPT 90	1	99	1	
Staphylococcus hominis subsp. hominis		1.61	0.11	ESL 8	1	1	1	
Kocuria kristinae		1.17	-.01		99	1	93	
203242	**Low discrimination**							**203242**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		68.48	0.43	VPT 90	99			
Staphylococcus hominis subsp. hominis		30.43	0.47	ESL 8	1			
Kocuria varians		0.74	0.13		1			
203243	**Good identification**							**203243**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		96.50	0.31	VPT 90	99			
Staphylococcus hominis subsp. hominis		3.50	0.12	ESL 8	1			
203250	**Low discrimination**							**203250**
					OXI	NOV		
Macrococcus equipersicus		49.91	0.38	NIT 1	99	99		
Macrococcus bovicus		30.32	0.32	SUC 9	99	99		
Staphylococcus hominis subsp. hominis		10.92	0.29		1	1		
203252	**Low discrimination**							**203252**
					NOV			
Staphylococcus hominis subsp. hominis		86.80	0.30	ESL 1	1			
Staphylococcus hominis subsp. novobiosepticus		13.20	0.02	MAN 13	99			
203300	**Low discrimination**							**203300**
					OXI	NOV		
Kocuria varians		71.47	0.41	ESL 1	13	1		
Staphylococcus hominis subsp. novobiosepticus		15.28	0.17	SUC 12	1	99		
Dermacoccus nishinomiyaensis		8.49	0.14		93	80		
203302	**Low discrimination**							**203302**
					NOV			
Kocuria varians		63.45	0.30	ESL 1	1			
Staphylococcus hominis subsp. novobiosepticus		25.90	0.12	SUC 12	99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		6.49	0.18		1		
203320	**Good identification**						**203320**
Kocuria varians		100.0	0.32	ESL 1			
203340	**Good identification**						**203340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		92.09	0.59	VPT 90	99		
Staphylococcus hominis subsp. hominis		6.78	0.47	ESL 8	1		
Kocuria varians		0.96	0.28		1		
203341	**Good identification**						**203341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.45	0.46	VPT 90	99		
Staphylococcus hominis subsp. hominis		0.59	0.12	ESL 8	1		
Kocuria varians		0.36	0.06		1		
203342	**Low discrimination**						**203342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		85.39	0.53	VPT 90	99		
Staphylococcus hominis subsp. hominis		13.70	0.48	ESL 8	1		
Kocuria varians		0.46	0.16		1		
203343	**Good identification**						**203343**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		96.77	0.41	VPT 90	99		
Staphylococcus epidermidis		1.96	0.14	ESL 8	1		
Staphylococcus hominis subsp. hominis		1.27	0.13		1		
203350	**Low discrimination**						**203350**
					OXI	NOV	
Micrococcus equipersicus		34.81	0.31	NIT 1	99	99	
Micrococcus bovicus		21.15	0.25	SUC 9	99	99	
Staphylococcus hominis subsp. hominis		19.25	0.30		1	1	
203352	**Low discrimination**						**203352**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		65.50	0.31	ESL 1	1	1	
Staphylococcus hominis subsp. novobiosepticus		27.59	0.12	MAN 13	1	99	
Staphylococcus saprophyticus subsp. bovis		3.94	0.02		99	99	
203371	**Low discrimination**						**203371**
					OXI		
Staphylococcus equorum		59.17	0.27	GLR 90	1		
Staphylococcus sciuri		20.23	0.21	TRE 99	99		
Staphylococcus xylosus		11.45	0.11		1		
203373	**Low discrimination**						**203373**
Staphylococcus equorum		71.02	0.36	GLR 90			
Staphylococcus xylosus		15.21	0.21	TRE 99			
Staphylococcus gallinarum		6.51	0.12				
203400	**Low discrimination**						**203400**
					NOV	OXI	
Staphylococcus piscifermentans		40.09	0.48	ARG 99	1	1	
Micrococcus carouselicus		26.96	0.28		99	99	
Kocuria varians		17.44	0.32		1	13	
203402	**Acceptable identification**						**203402**
Staphylococcus piscifermentans		92.44	0.52	ARG 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		7.56	0.21					
203410	**Low discrimination**							**203410**
Macrococcus equipersicus		61.40	0.50	NIT 1				
Macrococcus bovicus		25.65	0.41					
Macrococcus carouselicus		6.08	0.24					
203412	**Acceptable identification**							**203412**
					OXI	NOV		
Staphylococcus piscifermentans		91.18	0.44	ARG 99	1	1		
Macrococcus equipersicus		6.22	0.09		99	99		
Macrococcus bovicus		2.60	0.00		99	99		
203440	**Low discrimination**							**203440**
					NOV	OXI		
Staphylococcus piscifermentans		53.62	0.48	ARG 99	1	1		
Macrococcus equipersicus		33.15	0.33		99	99		
Kocuria varians		5.12	0.19		1	13		
203442	**Acceptable identification**							**203442**
Staphylococcus piscifermentans		94.29	0.52	ARG 99				
Staphylococcus hominis subsp. hominis		4.01	0.16					
Kocuria varians		1.69	0.07					
203450	**Low discrimination**							**203450**
					OXI	PYR		
Macrococcus equipersicus		69.54	0.59	NIT 1	99	1		
Macrococcus bovicus		25.07	0.48		99	1		
Staphylococcus saprophyticus subsp. bovis		2.85	0.31		1	99		
203452	**Low discrimination**							**203452**
					OXI	NOV	PYR	
Staphylococcus piscifermentans		69.52	0.44	ARG 99	1	1	15	
Staphylococcus saprophyticus subsp. bovis		13.02	0.18		1	99	99	
Macrococcus equipersicus		12.83	0.18		99	99	1	
203453	**Low discrimination**							**203453**
					NOV			
Staphylococcus cohnii subsp. urealyticum		79.75	0.28	GLR 99	99			
Staphylococcus saprophyticus subsp. bovis		12.66	0.06		99			
Aerococcus viridans		7.60	0.18		1			
203500	**Low discrimination**							**203500**
					NOV	OXI		
Staphylococcus piscifermentans		58.57	0.52	ARG 99	1	1		
Kocuria varians		25.48	0.36		1	13		
Macrococcus carouselicus		11.26	0.20		99	99		
203502	**Good identification**							**203502**
Staphylococcus piscifermentans		92.44	0.55	ARG 99				
Kocuria varians		7.56	0.25					
203510	**Low discrimination**							**203510**
					OXI	NOV		
Macrococcus equipersicus		53.41	0.43	NIT 1	99	99		
Macrococcus bovicus		22.31	0.33		99	99		
Staphylococcus piscifermentans		18.44	0.44		1	1		
203512	**Good identification**							**203512**
					OXI	NOV		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		98.09	0.48	ARG 99	1	1		
Macrococcus equipersicus		1.91	0.01		99	99		
203520	**Acceptable identification**							**203520**
Kocuria varians		93.80	0.26	ESL 1				
Staphylococcus piscifermentans		6.20	0.11	TRE 7				
203540	**Low discrimination**							**203540**
					NOV	OXI		
Staphylococcus piscifermentans		74.82	0.52	ARG 99	1	1		
Macrococcus equipersicus		13.22	0.25		99	99		
Kocuria varians		7.14	0.22		1	13		
203542	**Very good identification**							**203542**
Staphylococcus piscifermentans		95.36	0.55	ARG 99				
Staphylococcus hominis subsp. hominis		2.93	0.17					
Kocuria varians		1.71	0.11					
203550	**Low discrimination**							**203550**
					OXI	PYR		
Macrococcus equipersicus		53.45	0.51	NIT 1	99	1		
Staphylococcus saprophyticus subsp. bovis		20.45	0.43		1	99		
Macrococcus bovicus		19.27	0.41		99	1		
203551	**Low discrimination**							**203551**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		75.38	0.31	ESL 1	1	99		
Staphylococcus sciuri		9.89	0.13		99	1		
Macrococcus equipersicus		7.96	0.10		99	1		
203552	**Low discrimination**							**203552**
					OXI	NOV	PYR	
Staphylococcus piscifermentans		63.65	0.48	ARG 99	1	1	15	
Staphylococcus saprophyticus subsp. bovis		31.79	0.31		1	99	99	
Macrococcus equipersicus		3.36	0.10		99	99	1	
203571	**Low discrimination**							**203571**
					OXI			
Staphylococcus equorum		66.19	0.27	GLR 90	1			
Staphylococcus xylosus		16.17	0.13	SUC 99	1			
Staphylococcus sciuri		7.39	0.11		99			
203573	**Low discrimination**							**203573**
Staphylococcus equorum		68.53	0.36	GLR 90				
Staphylococcus xylosus		18.54	0.23	SUC 99				
Staphylococcus gallinarum		6.29	0.12					
203600	**Low discrimination**							**203600**
					NOV	OXI		
Staphylococcus piscifermentans		48.69	0.52	ARG 99	1	1		
Macrococcus carouselicus		44.32	0.34		99	99		
Kocuria varians		1.93	0.14		1	13		
203602	**Very good identification**							**203602**
Staphylococcus piscifermentans		96.54	0.55	ARG 99				
Staphylococcus hominis subsp. hominis		2.74	0.16					
Kocuria varians		0.72	0.03					
203610	**Low discrimination**							**203610**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Staphylococcus vitulinus		69.30	0.45	URE 1	99	99	
Macrococcus carouselicus		10.62	0.30		99	99	
Staphylococcus piscifermentans		7.82	0.44		1	1	
203611	**Acceptable identification**						**203611**
Staphylococcus sciuri		93.70	0.36	URE 1			
Staphylococcus vitulinus		6.30	0.04				
203612	**Acceptable identification**						**203612**
					OXI	NOV	
Staphylococcus piscifermentans		90.85	0.48	ARG 99	1	1	
Staphylococcus vitulinus		5.42	0.04		99	99	
Staphylococcus simulans		2.10	0.03		1	1	
203630	**Very good identification**						**203630**
Staphylococcus vitulinus		99.48	0.54	URE 1			
Staphylococcus sciuri		0.52	0.15				
203631	**Low discrimination**						**203631**
					OXI		
Staphylococcus sciuri		52.54	0.34	URE 1	99		
Staphylococcus equorum		33.60	0.27		1		
Staphylococcus vitulinus		12.65	0.13		99		
203633	**Low discrimination**						**203633**
					OXI		
Staphylococcus equorum		84.78	0.35	GLR 90	1		
Staphylococcus sciuri		8.39	0.18	GAL 91	99		
Staphylococcus lentus		3.47	0.12	MLT 91	99		
203640	**Low discrimination**						**203640**
Staphylococcus hominis subsp. hominis		73.56	0.56	ESL 1			
Staphylococcus piscifermentans		20.49	0.52				
Staphylococcus warneri		3.01	0.33				
203642	**Low discrimination**						**203642**
Staphylococcus hominis subsp. hominis		72.74	0.58	ESL 1			
Staphylococcus piscifermentans		25.89	0.55				
Staphylococcus warneri		0.85	0.23				
203650	**Low discrimination**						**203650**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		31.02	0.43	ESL 1	1	99	
Macrococcus equipersicus		18.71	0.38		99	1	
Macrococcus bovicus		17.05	0.36		99	1	
203651	**Low discrimination**						**203651**
					OXI	PYR	
Staphylococcus sciuri		84.26	0.48	URE 1	99	1	
Staphylococcus saprophyticus subsp. bovis		12.59	0.31		1	99	
Staphylococcus equorum		1.10	0.07		1	5	
203652	**Low discrimination**						**203652**
					NOV	PYR	
Staphylococcus piscifermentans		38.44	0.48	ARG 99	1	15	
Staphylococcus hominis subsp. hominis		37.65	0.41		1	1	
Staphylococcus saprophyticus subsp. bovis		19.20	0.31		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
203653	**Low discrimination**						**203653**
					OXI	PYR	
Staphylococcus sciuri		54.57	0.32	URE 1	99	1	
Staphylococcus saprophyticus subsp. bovis		12.53	0.18	LAC 14	1	99	
Staphylococcus equorum		11.29	0.15		1	5	
203670	**Low discrimination**						**203670**
					OXI		
Staphylococcus sciuri		58.03	0.27	URE 1	99		
Staphylococcus vitulinus		30.06	0.13	MNS 89	99		
Staphylococcus equorum		8.15	0.07		1		
203671	**Low discrimination**						**203671**
					OXI		
Staphylococcus equorum		60.81	0.48	GLR 90	1		
Staphylococcus sciuri		35.38	0.46	GAL 91	99		
Staphylococcus xylosus		2.85	0.19		1		
203673	**Good identification**						**203673**
					OXI		
Staphylococcus equorum		90.25	0.56	GLR 90	1		
Staphylococcus xylosus		4.69	0.28	GAL 91	1		
Staphylococcus sciuri		3.32	0.30		99		
203700	**Low discrimination**						**203700**
					OXI	NOV	
Staphylococcus piscifermentans		69.79	0.55	ARG 99	1	1	
Macrococcus carouselicus		18.15	0.27		99	99	
Rothia mucilaginosa		7.47	0.15		1	1	
203702	**Very good identification**						**203702**
Staphylococcus piscifermentans		97.28	0.59	ARG 99			
Staphylococcus hominis subsp. hominis		1.99	0.17				
Kocuria varians		0.72	0.07				
203710	**Low discrimination**						**203710**
					OXI	NOV	
Staphylococcus piscifermentans		46.86	0.48	ARG 99	1	1	
Macrococcus carouselicus		18.18	0.23		99	99	
Staphylococcus sciuri		9.99	0.24		99	99	
203711	**Low discrimination**						**203711**
					OXI		
Staphylococcus sciuri		56.55	0.42	URE 1	99		
Staphylococcus lentus		41.35	0.41		99		
Staphylococcus equorum		1.44	0.07		1		
203712	**Acceptable identification**						**203712**
					OXI	NOV	
Staphylococcus piscifermentans		93.75	0.51	ARG 99	1	1	
Staphylococcus lentus		2.82	0.12		99	99	
Staphylococcus sciuri		2.17	0.07		99	99	
203713	**Acceptable identification**						**203713**
					OXI		
Staphylococcus lentus		90.91	0.53	URE 1	99		
Staphylococcus sciuri		5.71	0.26		99		
Staphylococcus equorum		2.29	0.15		1		
203731	**Low discrimination**						**203731**
					OXI		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus equorum		61.10	0.48	GLR 90	1			
Staphylococcus sciuri		18.34	0.40	MLT 91	99			
Staphylococcus lentus		17.78	0.41		99			
203733	**Low discrimination**							**203733**
					OXI			
Staphylococcus equorum		68.07	0.56	GLR 90	1			
Staphylococcus lentus		27.31	0.53	MLT 91	99			
Staphylococcus xylosus		2.69	0.26		1			
203740	**Low discrimination**							**203740**
Staphylococcus hominis subsp. hominis		64.00	0.57	ESL 1				
Staphylococcus piscifermentans		24.69	0.55					
Rothia mucilaginosa		6.16	0.23					
203741	**Low discrimination**							**203741**
					OXI	NOV	PYR	
Rothia mucilaginosa		73.65	0.33	VPT 95	1	1	75	
Staphylococcus sciuri		8.46	0.23	URE 1	99	99	1	
Kocuria kristinae		6.99	0.15		99	1	93	
203742	**Low discrimination**							**203742**
					NOV			
Staphylococcus hominis subsp. hominis		65.84	0.58	ESL 1	1			
Staphylococcus piscifermentans		32.45	0.59		1			
Staphylococcus hominis subsp. novobiosepticus		1.31	0.12		99			
203750	**Low discrimination**							**203750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		67.94	0.55	ESL 1	99	99		
Staphylococcus piscifermentans		8.50	0.48		1	15		
Staphylococcus hominis subsp. hominis		7.69	0.40		1	1		
203751	**Low discrimination**							**203751**
					OXI	PYR		
Staphylococcus sciuri		55.24	0.54	URE 1	99	1		
Staphylococcus lentus		21.80	0.48		99	6		
Staphylococcus saprophyticus subsp. bovis		17.01	0.43		1	99		
203752	**Low discrimination**							**203752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		41.63	0.43	ESL 1	99	99		
Staphylococcus piscifermentans		31.26	0.51		1	15		
Staphylococcus hominis subsp. hominis		22.11	0.41		1	1		
203753	**Low discrimination**							**203753**
					OXI			
Staphylococcus lentus		72.97	0.59	URE 1	99			
Staphylococcus equorum		9.15	0.36		1			
Staphylococcus sciuri		8.49	0.38		99			
203761	**Low discrimination**							**203761**
					OXI			
Staphylococcus equorum		37.32	0.27	GLR 90	1			
Staphylococcus gallinarum		35.20	0.24	MAN 99	1			
Staphylococcus sciuri		12.76	0.21		99			
203763	**Low discrimination**							**203763**
Staphylococcus equorum		42.95	0.36	GLR 90				
Staphylococcus gallinarum		38.57	0.32	MAN 99				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		14.09	0.24				
203770	**Low discrimination**						**203770**
					OXI	PYR	
Staphylococcus sciuri		49.68	0.33	URE 1	99	1	
Staphylococcus equorum		36.36	0.27	MNS 89	1	5	
Staphylococcus saprophyticus subsp. bovis		6.63	0.14		1	99	
203771	**Low discrimination**						**203771**
					OXI		
Staphylococcus equorum		76.12	0.68	GLR 90	1		
Staphylococcus sciuri		8.50	0.52		99		
Staphylococcus gallinarum		7.33	0.45		1		
203772	**Low discrimination**						**203772**
					OXI		
Staphylococcus equorum		74.53	0.36	GLR 90	1		
Staphylococcus gallinarum		6.84	0.12	MNS 99	1		
Staphylococcus sciuri		6.45	0.17		99		
203773	**Low discrimination**						**203773**
					OXI		
Staphylococcus equorum		81.21	0.77	GLR 90	1		
Staphylococcus gallinarum		7.45	0.53		1		
Staphylococcus lentus		6.54	0.59		99		
204000	**Low discrimination**						**204000**
					OXI		
Dermacoccus nishinomiyaensis		89.67	0.74	PHS 5	93		
Micrococcus sp.		9.23	0.44		89		
Kocuria varians		0.99	0.41		13		
204001	**Low discrimination**						**204001**
					OXI		
Dermacoccus nishinomiyaensis		76.74	0.33	PHS 5	93		
Micrococcus sp.		15.96	0.09	MNS 1	89		
Kocuria varians		7.31	0.19		13		
204002	**Low discrimination**						**204002**
					OXI		
Dermacoccus nishinomiyaensis		70.83	0.33	PHS 5	93		
Kocuria varians		21.89	0.30	LAC 1	13		
Micrococcus sp.		7.29	0.03		89		
204010	**Low discrimination**						**204010**
					YEP		
Macrococcus equipersicus		48.13	0.43	PHS 1	99		
Dermacoccus nishinomiyaensis		25.44	0.33		1		
Macrococcus bovicus		23.31	0.35		99		
204020	**Low discrimination**						**204020**
					OXI		
Dermacoccus nishinomiyaensis		67.21	0.33	PHS 5	93		
Kocuria varians		25.87	0.32	XYL 1	13		
Micrococcus sp.		6.92	0.03		89		
204040	**Low discrimination**						**204040**
					COA	OXI	
Dermacoccus nishinomiyaensis		37.62	0.33	PHS 5	1	93	
Micrococcus sp.		28.85	0.21	MLT 1	1	89	
Staphylococcus aureus subsp. anaerobius		13.55	0.16		99	1	
204050	**Low discrimination**						**204050**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Macrococcus equipersicus		67.04	0.52	PHS 1	99			
Macrococcus bovicus		28.01	0.43		99			
Staphylococcus kloosii		4.95	0.46		1			
204052	**Low discrimination**							**204052**
					OXI			
Staphylococcus kloosii		84.82	0.47	TRE 99	1			
Macrococcus equipersicus		10.71	0.11		99			
Macrococcus bovicus		4.48	0.02		99			
204070	**Low discrimination**							**204070**
					OXI			
Staphylococcus kloosii		56.31	0.34	TRE 99	1			
Macrococcus equipersicus		30.81	0.11		99			
Macrococcus bovicus		12.88	0.02		99			
204072	**Good identification**							**204072**
Staphylococcus kloosii		100.0	0.35	TRE 99				
204100	**Good identification**							**204100**
					OXI			
Dermacoccus nishinomiyaensis		91.34	0.59	PHS 5	93			
Kocuria varians		7.96	0.45		13			
Micrococcus sp.		0.50	0.03		89			
204102	**Low discrimination**							**204102**
					OXI			
Kocuria varians		70.87	0.33	PHS 1	13			
Dermacoccus nishinomiyaensis		29.13	0.18		93			
204110	**Low discrimination**							**204110**
					YEP			
Macrococcus equipersicus		55.96	0.36	PHS 1	99			
Macrococcus bovicus		27.09	0.28		99			
Dermacoccus nishinomiyaensis		13.14	0.18		1			
204112	**Good identification**							**204112**
Staphylococcus kloosii		100.0	0.27	TRE 99				
204120	**Low discrimination**							**204120**
					OXI			
Kocuria varians		75.19	0.35	PHS 1	13			
Dermacoccus nishinomiyaensis		24.81	0.18		93			
204140	**Low discrimination**							**204140**
					OXI	YEP	NOV	
Kocuria varians		54.11	0.31	PHS 1	13	80	1	
Dermacoccus nishinomiyaensis		28.57	0.18		93	1	80	
Macrococcus equipersicus		17.31	0.18		99	99	99	
204150	**Low discrimination**							**204150**
					OXI			
Macrococcus equipersicus		57.07	0.45	PHS 1	99			
Macrococcus bovicus		23.85	0.35		99			
Staphylococcus kloosii		19.09	0.52		1			
204152	**Very good identification**							**204152**
					OXI			
Staphylococcus kloosii		97.29	0.53	TRE 99	1			
Macrococcus equipersicus		2.71	0.04		99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
204170	**Low discrimination**							**204170**
					OXI			
Staphylococcus kloosii		89.22	0.40	TRE 99	1			
Macrococcus equipersicus		10.78	0.04		99			
204172	**Good identification**							**204172**
Staphylococcus kloosii		100.0	0.41	TRE 99				
204200	**Low discrimination**							**204200**
					COA	OXI		
Dermacoccus nishinomiyaensis		80.64	0.49	PHS 5	1	93		
Micrococcus sp.		12.86	0.23	SUC 6	1	89		
Staphylococcus aureus subsp. anaerobius		4.60	0.16		99	1		
204240	**Very good identification**							**204240**
					COA	NOV	OXI	
Staphylococcus aureus subsp. anaerobius		98.88	0.57	URE 1	99	1	1	
Staphylococcus hominis subsp. novobiosepticus		0.28	0.10		1	99	1	
Micrococcus sp.		0.21	0.00		1	1	89	
204250	**Low discrimination**							**204250**
					OXI	COA		
Macrococcus bovicus		36.71	0.31	PHS 1	99	1		
Macrococcus equipersicus		34.75	0.32		99	1		
Staphylococcus aureus subsp. anaerobius		24.74	0.16		1	99		
204300	**Low discrimination**							**204300**
					OXI			
Dermacoccus nishinomiyaensis		84.31	0.34	PHS 5	93			
Kocuria varians		15.70	0.27	SUC 6	13			
204341	**Low discrimination**							**204341**
					NOV	YEP		
Staphylococcus epidermidis		79.13	0.29	VPT 96	1	1		
Staphylococcus hominis subsp. novobiosepticus		17.09	0.07	NIT 97	99	1		
Staphylococcus chromogenes		3.78	-.02		1	95		
204342	**Low discrimination**							**204342**
					NOV			
Staphylococcus epidermidis		71.37	0.34	VPT 96	1			
Staphylococcus hominis subsp. novobiosepticus		19.47	0.14	NIT 97	99			
Staphylococcus hominis subsp. hominis		7.49	0.17		1			
204343	**Low discrimination**							**204343**
					NOV	YEP		
Staphylococcus epidermidis		83.51	0.39	VPT 96	1	1		
Staphylococcus chromogenes		11.35	0.17	NIT 97	1	95		
Staphylococcus hominis subsp. novobiosepticus		3.07	0.02		99	1		
204352	**Good identification**							**204352**
Staphylococcus kloosii		100.0	0.32	SUC 9				
204400	**Low discrimination**							**204400**
					OXI	NOV		
Dermacoccus nishinomiyaensis		68.47	0.33	PHS 5	93	80		
Staphylococcus cohnii subsp. cohnii		9.71	0.25	TRE 1	1	99		
Micrococcus sp.		7.05	0.03		89	1		
204410	**Low discrimination**							**204410**
					OXI			
Staphylococcus kloosii		36.35	0.61	MLT 95	1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		34.94	0.43		99		
Macrococcus bovicus		25.38	0.39		99		
204411	**Low discrimination**						**204411**
					YEP		
Staphylococcus cohnii subsp. cohnii		62.00	0.28	URE 1	12		
Staphylococcus cohnii subsp. urealyticum		21.23	0.19	PHS 9	64		
Staphylococcus kloosii		6.31	0.20		99		
204412	**Very good identification**						**204412**
					OXI		
Staphylococcus kloosii		98.48	0.62	MLT 95	1		
Macrococcus equipersicus		0.88	0.02		99		
Macrococcus bovicus		0.64	-.02		99		
204413	**Low discrimination**						**204413**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		88.62	0.40	GLR 99	99	64	
Staphylococcus felis		8.56	0.04		1	1	
Staphylococcus kloosii		2.82	0.21		99	99	
204430	**Acceptable identification**						**204430**
					OXI		
Staphylococcus kloosii		93.72	0.49	MLT 95	1		
Macrococcus equipersicus		3.64	0.02		99		
Macrococcus bovicus		2.64	-.02		99		
204432	**Good identification**						**204432**
Staphylococcus kloosii		100.0	0.50	MLT 95			
204440	**Low discrimination**						**204440**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		38.32	0.35	URE 1	1	12	
Staphylococcus kloosii		32.25	0.46	PHS 9	1	99	
Macrococcus equipersicus		22.98	0.26		99	99	
204441	**Acceptable identification**						**204441**
Staphylococcus cohnii subsp. cohnii		92.84	0.36	URE 1			
Staphylococcus cohnii subsp. urealyticum		7.16	0.14	PHS 9			
204442	**Good identification**						**204442**
Staphylococcus kloosii		100.0	0.47	MAN 99			
204443	**Good identification**						**204443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	GLR 99			
204450	**Low discrimination**						**204450**
					OXI		
Staphylococcus kloosii		80.52	0.88		1		
Macrococcus equipersicus		11.02	0.52		99		
Macrococcus bovicus		6.90	0.47		99		
204451	**Low discrimination**						**204451**
					YEP		
Staphylococcus cohnii subsp. cohnii		41.60	0.38	URE 1	12		
Staphylococcus cohnii subsp. urealyticum		25.49	0.35	PHS 9	64		
Staphylococcus kloosii		25.39	0.46		99		
204452	**Very good identification**						**204452**
					OXI		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		99.33	0.88		1		
Staphylococcus saprophyticus subsp. saprophyticus		0.31	0.08		1		
Macrococcus equipersicus		0.13	0.11		99		
204453	**Good identification**						**204453**
Staphylococcus cohnii subsp. urealyticum		90.36	0.55	GLR 99			
Staphylococcus kloosii		9.64	0.47				
204460	**Good identification**						**204460**
Staphylococcus kloosii		100.0	0.34	MAN 99			
204462	**Good identification**						**204462**
Staphylococcus kloosii		100.0	0.35	MAN 99			
204470	**Very good identification**						**204470**
					OXI		
Staphylococcus kloosii		99.11	0.75		1		
Macrococcus equipersicus		0.55	0.11		99		
Macrococcus bovicus		0.34	0.05		99		
204471	**Good identification**						**204471**
Staphylococcus kloosii		100.0	0.34	MNS 1			
204472	**Very good identification**						**204472**
Staphylococcus kloosii		100.0	0.76				
204473	**Low discrimination**						**204473**
Staphylococcus kloosii		72.53	0.35	MNS 1			
Staphylococcus cohnii subsp. urealyticum		27.47	0.14				
204502	**Low discrimination**						**204502**
					NOV		
Staphylococcus kloosii		54.52	0.27	MAN 99	99		
Kocuria varians		45.49	0.10	MLT 95	1		
204510	**Low discrimination**						**204510**
					OXI		
Staphylococcus kloosii		72.91	0.67	MLT 95	1		
Macrococcus equipersicus		15.47	0.36		99		
Macrococcus bovicus		11.24	0.31		99		
204512	**Very good identification**						**204512**
Staphylococcus kloosii		100.0	0.68	MLT 95			
204513	**Low discrimination**						**204513**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		52.68	0.32	GLR 99	99	64	
Staphylococcus felis		39.73	0.15		1	1	
Staphylococcus kloosii		7.59	0.27		99	99	
204530	**Very good identification**						**204530**
Staphylococcus kloosii		100.0	0.55	MLT 95			
204532	**Very good identification**						**204532**
Staphylococcus kloosii		100.0	0.55	MLT 95			
204540	**Low discrimination**						**204540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	YEP	
Staphylococcus kloosii		78.04	0.52	MAN 99	1	99	
Macrococcus equipersicus		12.28	0.18		99	99	
Staphylococcus cohnii subsp. cohnii		5.31	0.16		1	12	
204542	**Very good identification**						**204542**
Staphylococcus kloosii		100.0	0.53	MAN 99			
204543	**Acceptable identification**						**204543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		90.39	0.27	GLR 99	99		
Staphylococcus chromogenes		9.61	-.02	MAN 91	1		
204550	**Good identification**						**204550**
					OXI		
Staphylococcus kloosii		94.68	0.93		1		
Macrococcus equipersicus		2.86	0.45		99		
Macrococcus bovicus		1.79	0.39		99		
204551	**Low discrimination**						**204551**
Staphylococcus kloosii		65.94	0.52	MNS 1			
Staphylococcus cohnii subsp. urealyticum		14.62	0.27				
Staphylococcus saprophyticus subsp. bovis		10.02	0.18				
204552	**Excellent identification**						**204552**
					OXI	PYR	
Staphylococcus kloosii		99.79	0.94		1	52	
Staphylococcus saprophyticus subsp. bovis		0.14	0.18		1	99	
Macrococcus equipersicus		0.03	0.04		99	1	
204553	**Low discrimination**						**204553**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		65.60	0.48	GLR 99	99	64	
Staphylococcus kloosii		31.70	0.53		99	99	
Staphylococcus caprae		1.59	-.02		1	1	
204560	**Good identification**						**204560**
Staphylococcus kloosii		100.0	0.40	MAN 99			
204562	**Good identification**						**204562**
Staphylococcus kloosii		100.0	0.41	MAN 99			
204570	**Very good identification**						**204570**
					OXI		
Staphylococcus kloosii		99.80	0.81		1		
Macrococcus equipersicus		0.12	0.04		99		
Macrococcus bovicus		0.08	-.02		99		
204571	**Acceptable identification**						**204571**
Staphylococcus kloosii		93.16	0.40	MNS 1			
Staphylococcus xylosus		6.84	-.04				
204572	**Very good identification**						**204572**
Staphylococcus kloosii		100.0	0.82				
204610	**Low discrimination**						**204610**
					OXI		
Macrococcus bovicus		46.21	0.27	PHS 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		26.18	0.41		1			
Macrococcus equipersicus		25.17	0.23		99			
204612	**Low discrimination**							**204612**
					NOV	YEP		
Staphylococcus kloosii		88.14	0.41	SUC 9	99	99		
Staphylococcus simulans		11.86	0.06	MLT 95	1	1		
204630	**Good identification**							**204630**
Staphylococcus kloosii		100.0	0.28	SUC 9				
204632	**Good identification**							**204632**
Staphylococcus kloosii		100.0	0.29	SUC 9				
204640	**Low discrimination**							**204640**
					COA			
Staphylococcus warneri		43.80	0.41	PHS 1	1			
Staphylococcus aureus subsp. anaerobius		33.11	0.16		99			
Staphylococcus hominis subsp. hominis		14.17	0.25		1			
204650	**Low discrimination**							**204650**
					OXI			
Staphylococcus kloosii		61.22	0.67	SUC 9	1			
Macrococcus bovicus		13.27	0.34		99			
Macrococcus equipersicus		8.37	0.32		99			
204652	**Low discrimination**							**204652**
					NOV			
Staphylococcus kloosii		72.63	0.68	SUC 9	99			
Staphylococcus saprophyticus subsp. saprophyticus		22.78	0.29		99			
Staphylococcus warneri		2.08	0.28		1			
204670	**Very good identification**							**204670**
Staphylococcus kloosii		100.0	0.54	SUC 9				
204672	**Good identification**							**204672**
Staphylococcus kloosii		93.95	0.55	SUC 9				
Staphylococcus arlettae		6.05	0.19					
204703	**Low discrimination**							**204703**
					YEP	PYR	COA	
Staphylococcus chromogenes		76.52	0.39	ARG 90	95	45	1	
Staphylococcus intermedius		10.81	0.23	NIT 99	1	90	99	
Staphylococcus hyicus		9.70	0.17		1	1	24	
204710	**Low discrimination**							**204710**
					OXI			
Staphylococcus kloosii		60.62	0.46	SUC 9	1			
Macrococcus bovicus		23.62	0.19	MLT 95	99			
Macrococcus equipersicus		12.87	0.15		99			
204712	**Good identification**							**204712**
Staphylococcus kloosii		100.0	0.47	SUC 9				
204713	**Low discrimination**							**204713**
					COA	YEP		
Staphylococcus intermedius		41.35	0.32	BGA 99	99	1		
Staphylococcus chromogenes		30.55	0.28	NIT 93	1	95		
Staphylococcus felis		28.10	0.22		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
204730	**Good identification**							**204730**
Staphylococcus kloosii		100.0	0.34	SUC 9				
204732	**Good identification**							**204732**
Staphylococcus kloosii		100.0	0.35	SUC 9				
204742	**Low discrimination**							**204742**
					NOV			
Staphylococcus hominis subsp. hominis		57.31	0.27	NIT 99	1			
Staphylococcus kloosii		22.97	0.32	PHS 3	99			
Staphylococcus warneri		14.66	0.21		1			
204743	**Low discrimination**							**204743**
					OXI	COA	YEP	
Staphylococcus chromogenes		81.74	0.39	ARG 90	1	1	95	
Staphylococcus intermedius		17.32	0.27	NIT 99	1	99	1	
Kocuria kristinae		0.95	-.03		99	1	15	
204750	**Low discrimination**							**204750**
					OXI	PYR		
Staphylococcus kloosii		74.20	0.73	SUC 9	1	52		
Staphylococcus saprophyticus subsp. bovis		18.42	0.43		1	99		
Macrococcus bovicus		3.55	0.27		99	1		
204751	**Low discrimination**							**204751**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		59.13	0.31	NIT 99	1	99		
Staphylococcus intermedius		21.18	0.20		99	1		
Staphylococcus kloosii		9.62	0.32		1	99		
204752	**Good identification**							**204752**
					PYR			
Staphylococcus kloosii		92.61	0.73	SUC 9	52			
Staphylococcus saprophyticus subsp. bovis		5.31	0.31		99			
Staphylococcus saprophyticus subsp. saprophyticus		1.30	0.07		7			
204753	**Low discrimination**							**204753**
					COA	NOV	YEP	
Staphylococcus intermedius		52.27	0.36	BGA 99	99	1	1	
Staphylococcus chromogenes		25.74	0.28	NIT 93	1	1	95	
Staphylococcus saprophyticus subsp. bovis		5.94	0.18		1	99	80	
204770	**Very good identification**							**204770**
Staphylococcus kloosii		99.01	0.60	SUC 9				
Staphylococcus saprophyticus subsp. bovis		0.99	0.02					
204772	**Low discrimination**							**204772**
Staphylococcus kloosii		84.18	0.61	SUC 9				
Staphylococcus arlettae		15.82	0.35					
204773	**Low discrimination**							**204773**
					COA	NOV		
Staphylococcus xylosus		42.21	0.31	GLR 95	1	99		
Staphylococcus arlettae		28.09	0.37	NIT 89	1	99		
Staphylococcus lutrae		15.47	0.18		99	1		
205000	**Low discrimination**							**205000**
					NOV	YEP		
Dermacoccus nishinomiyaensis		76.07	0.33	ESL 1	80	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		16.10	0.24	PHS 5	99	99	
Micrococcus sp.		7.83	0.03		1	41	
205010	**Good identification**						**205010**
					OXI		
Macrococcus equipersicus		97.06	0.50	PHS 1	99		
Macrococcus bovicus		2.46	0.16		99		
Staphylococcus kloosii		0.48	0.20		1		
205040	**Good identification**						**205040**
Macrococcus equipersicus		100.0	0.33	PHS 1			
205050	**Good identification**						**205050**
					OXI		
Macrococcus equipersicus		94.64	0.59	PHS 1	99		
Staphylococcus kloosii		3.29	0.46		1		
Macrococcus bovicus		2.07	0.23		99		
205052	**Low discrimination**						**205052**
					OXI		
Staphylococcus kloosii		78.85	0.47	TRE 99	1		
Macrococcus equipersicus		21.16	0.18		99		
205072	**Good identification**						**205072**
Staphylococcus kloosii		100.0	0.35	TRE 99			
205110	**Good identification**						**205110**
					OXI		
Macrococcus equipersicus		95.45	0.43	PHS 1	99		
Macrococcus bovicus		2.42	0.08		99		
Staphylococcus kloosii		2.14	0.26		1		
205112	**Low discrimination**						**205112**
					OXI		
Staphylococcus kloosii		70.61	0.27	TRE 99	1		
Macrococcus equipersicus		29.39	0.01	MLT 95	99		
205150	**Low discrimination**						**205150**
					OXI		
Macrococcus equipersicus		84.80	0.51	PHS 1	99		
Staphylococcus kloosii		13.35	0.52		1		
Macrococcus bovicus		1.85	0.16		99		
205152	**Good identification**						**205152**
					OXI		
Staphylococcus kloosii		94.41	0.53	TRE 99	1		
Macrococcus equipersicus		5.59	0.10		99		
205170	**Low discrimination**						**205170**
					OXI		
Staphylococcus kloosii		79.57	0.40	TRE 99	1		
Macrococcus equipersicus		20.43	0.10		99		
205172	**Good identification**						**205172**
Staphylococcus kloosii		100.0	0.41	TRE 99			
205210	**Acceptable identification**						**205210**
Macrococcus equipersicus		93.99	0.29	PHS 1			
Macrococcus bovicus		6.02	0.03	SUC 9			
205250	**Acceptable identification**						**205250**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					OXI	
Macrococcus equipersicus		91.75	0.38	PHS 1	99	
Macrococcus bovicus		5.07	0.11	SUC 9	99	
Staphylococcus kloosii		3.19	0.26		1	
205350	**Low discrimination**					**205350**
					OXI	
Macrococcus equipersicus		82.47	0.31	PHS 1	99	
Staphylococcus kloosii		12.98	0.32	SUC 9	1	
Macrococcus bovicus		4.56	0.03		99	
205352	**Good identification**					**205352**
Staphylococcus kloosii		100.0	0.32	SUC 9		
205410	**Low discrimination**					**205410**
					OXI	
Macrococcus equipersicus		64.80	0.50	PHS 1	99	
Staphylococcus kloosii		31.72	0.61		1	
Macrococcus bovicus		2.46	0.19		99	
205412	**Very good identification**					**205412**
					OXI	
Staphylococcus kloosii		98.13	0.62	MLT 95	1	
Macrococcus equipersicus		1.87	0.09		99	
205413	**Acceptable identification**					**205413**
Staphylococcus cohnii subsp. urealyticum		92.44	0.31	GLR 99		
Staphylococcus kloosii		7.56	0.21			
205430	**Acceptable identification**					**205430**
					OXI	
Staphylococcus kloosii		92.38	0.49	MLT 95	1	
Macrococcus equipersicus		7.62	0.09		99	
205432	**Good identification**					**205432**
Staphylococcus kloosii		100.0	0.50	MLT 95		
205440	**Low discrimination**					**205440**
					OXI	
Macrococcus equipersicus		60.23	0.33	PHS 1	99	
Staphylococcus kloosii		39.77	0.46	MAN 95	1	
205442	**Good identification**					**205442**
Staphylococcus kloosii		100.0	0.47	MAN 99		
205450	**Low discrimination**					**205450**
					OXI	
Staphylococcus kloosii		76.91	0.88		1	
Macrococcus equipersicus		22.36	0.59		99	
Macrococcus bovicus		0.73	0.27		99	
205451	**Low discrimination**					**205451**
					OXI	
Staphylococcus kloosii		59.48	0.46	MNS 1	1	
Staphylococcus cohnii subsp. urealyticum		23.23	0.26		1	
Macrococcus equipersicus		17.29	0.18		99	
205452	**Very good identification**					**205452**
					OXI	
Staphylococcus kloosii		99.69	0.88		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		0.27	0.18		99		
Staphylococcus cohnii subsp. urealyticum		0.04	0.06		1		
205453	**Low discrimination**						**205453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		77.79	0.47	GLR 99	99	64	
Staphylococcus kloosii		21.34	0.47		99	99	
Aerococcus viridans		0.87	0.18		1	1	
205460	**Good identification**						**205460**
Staphylococcus kloosii		100.0	0.34	MAN 99			
205462	**Good identification**						**205462**
Staphylococcus kloosii		100.0	0.35	MAN 99			
205470	**Very good identification**						**205470**
					OXI		
Staphylococcus kloosii		98.84	0.75		1		
Macrococcus equipersicus		1.16	0.18		99		
205471	**Good identification**						**205471**
Staphylococcus kloosii		100.0	0.34	MNS 1			
205472	**Very good identification**						**205472**
Staphylococcus kloosii		100.0	0.76				
205473	**Low discrimination**						**205473**
					NOV	YEP	
Staphylococcus kloosii		69.51	0.35	MNS 1	99	99	
Aerococcus viridans		11.36	0.18		1	1	
Staphylococcus cohnii subsp. urealyticum		10.24	0.06		99	64	
205502	**Low discrimination**						**205502**
					NOV	YEP	
Staphylococcus kloosii		74.89	0.27	MAN 99	99	99	
Staphylococcus piscifermentans		25.11	0.11	MLT 95	1	1	
205510	**Low discrimination**						**205510**
					OXI		
Staphylococcus kloosii		67.79	0.67	MLT 95	1		
Macrococcus equipersicus		30.57	0.43		99		
Macrococcus bovicus		1.16	0.12		99		
205512	**Very good identification**						**205512**
					OXI		
Staphylococcus kloosii		99.58	0.68	MLT 95	1		
Macrococcus equipersicus		0.42	0.01		99		
205530	**Very good identification**						**205530**
					OXI		
Staphylococcus kloosii		98.21	0.55	MLT 95	1		
Macrococcus equipersicus		1.79	0.01		99		
205532	**Very good identification**						**205532**
Staphylococcus kloosii		100.0	0.55	MLT 95			
205540	**Low discrimination**						**205540**
					OXI		
Staphylococcus kloosii		74.94	0.52	MAN 99	1		
Macrococcus equipersicus		25.06	0.25		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
205542	**Very good identification**						**205542**
					NOV	YEP	
Staphylococcus kloosii		98.27	0.53	MAN 99	99	99	
Staphylococcus piscifermentans		1.73	0.11		1	1	
205550	**Good identification**						**205550**
					OXI		
Staphylococcus kloosii		93.78	0.93		1		
Macrococcus equipersicus		6.02	0.51		99		
Macrococcus bovicus		0.20	0.19		99		
205551	**Low discrimination**						**205551**
					OXI		
Staphylococcus kloosii		86.93	0.52	MNS 1	1		
Staphylococcus cohnii subsp. urealyticum		7.50	0.19		1		
Macrococcus equipersicus		5.58	0.10		99		
205552	**Excellent identification**						**205552**
					OXI		
Staphylococcus kloosii		99.94	0.94		1		
Macrococcus equipersicus		0.06	0.10		99		
205553	**Low discrimination**						**205553**
					NOV	YEP	
Staphylococcus kloosii		54.77	0.53	MNS 1	99	99	
Staphylococcus cohnii subsp. urealyticum		44.08	0.39		99	64	
Aerococcus viridans		1.15	0.18		1	1	
205560	**Good identification**						**205560**
Staphylococcus kloosii		100.0	0.40	MAN 99			
205562	**Good identification**						**205562**
Staphylococcus kloosii		100.0	0.41	MAN 99			
205570	**Very good identification**						**205570**
					OXI		
Staphylococcus kloosii		99.74	0.81		1		
Macrococcus equipersicus		0.26	0.10		99		
205571	**Low discrimination**						**205571**
					YEP		
Staphylococcus kloosii		67.18	0.40	MNS 1	99		
Staphylococcus equorum		18.88	0.09		1		
Staphylococcus gallinarum		13.94	0.04		78		
205572	**Very good identification**						**205572**
Staphylococcus kloosii		100.0	0.82				
205573	**Low discrimination**						**205573**
					YEP		
Staphylococcus kloosii		45.04	0.41	MNS 1	99		
Staphylococcus equorum		30.04	0.17		1		
Staphylococcus gallinarum		21.13	0.12		78		
205610	**Low discrimination**						**205610**
					OXI		
Macrococcus equipersicus		52.44	0.29	PHS 1	99		
Staphylococcus kloosii		25.67	0.41	SUC 9	1		
Macrococcus carouselicus		16.86	0.14		99		
205612	**Good identification**						**205612**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.41	SUC 9		
205630	**Low discrimination**					**205630**
					OXI	
Staphylococcus kloosii		77.36	0.28	SUC 9	1	
Staphylococcus vitulinus		22.64	-.08	MLT 95	99	
205632	**Good identification**					**205632**
Staphylococcus kloosii		100.0	0.29	SUC 9		
205650	**Low discrimination**					**205650**
					OXI	
Staphylococcus kloosii		75.86	0.67	SUC 9	1	
Macrococcus equipersicus		22.05	0.38		99	
Macrococcus bovicus		1.83	0.15		99	
205652	**Very good identification**					**205652**
Staphylococcus kloosii		100.0	0.68	SUC 9		
205670	**Very good identification**					**205670**
Staphylococcus kloosii		100.0	0.54	SUC 9		
205671	**Low discrimination**					**205671**
					OXI	
Staphylococcus equorum		81.27	0.29	GLR 90	1	
Staphylococcus sciuri		12.60	0.16	NIT 92	99	
Staphylococcus gallinarum		6.13	0.04	GAL 91	1	
205672	**Good identification**					**205672**
					YEP	
Staphylococcus kloosii		92.81	0.55	SUC 9	99	
Staphylococcus arlettae		5.98	0.19		99	
Staphylococcus equorum		1.21	-.03		1	
205673	**Low discrimination**					**205673**
					YEP	
Staphylococcus equorum		85.88	0.38	GLR 90	1	
Staphylococcus gallinarum		6.17	0.12	NIT 92	78	
Staphylococcus arlettae		5.66	0.22	GAL 91	99	
205710	**Low discrimination**					**205710**
					OXI	
Staphylococcus kloosii		61.00	0.46	SUC 9	1	
Macrococcus equipersicus		27.51	0.22	MLT 95	99	
Macrococcus carouselicus		8.85	0.06		99	
205712	**Good identification**					**205712**
Staphylococcus kloosii		100.0	0.47	SUC 9		
205730	**Good identification**					**205730**
Staphylococcus kloosii		100.0	0.34	SUC 9		
205731	**Low discrimination**					**205731**
					OXI	
Staphylococcus equorum		86.55	0.29	GLR 90	1	
Staphylococcus sciuri		6.92	0.10	NIT 92	99	
Staphylococcus gallinarum		6.53	0.04	MLT 91	1	
205732	**Low discrimination**					**205732**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					YEP			
Staphylococcus kloosii		88.65	0.35	SUC 9	99			
Staphylococcus equorum		11.35	-.03	MLT 95	1			
205733	**Acceptable identification**							**205733**
Staphylococcus equorum		92.39	0.38	GLR 90				
Staphylococcus gallinarum		6.64	0.12	NIT 92				
Staphylococcus xylosus		0.98	-.04	MLT 91				
205740	**Low discrimination**							**205740**
					OXI	NOV	YEP	
Staphylococcus kloosii		65.46	0.32	SUC 9	1	99	99	
Macrococcus equipersicus		21.89	0.04	MAN 99	99	99	99	
Staphylococcus piscifermentans		12.66	0.11		1	1	1	
205742	**Low discrimination**							**205742**
					NOV	YEP		
Staphylococcus kloosii		78.88	0.32	SUC 9	99	99		
Staphylococcus piscifermentans		21.12	0.14	MAN 99	1	1		
205750	**Good identification**							**205750**
					OXI			
Staphylococcus kloosii		92.98	0.73	SUC 9	1			
Macrococcus equipersicus		5.97	0.31		99			
Macrococcus bovicus		0.49	0.07		99			
205752	**Very good identification**							**205752**
Staphylococcus kloosii		100.0	0.73	SUC 9				
205763	**Low discrimination**							**205763**
Staphylococcus gallinarum		87.32	0.32	NIT 99				
Staphylococcus equorum		12.68	0.17	MAN 91				
205770	**Good identification**							**205770**
					YEP			
Staphylococcus kloosii		94.31	0.60	SUC 9	99			
Staphylococcus equorum		2.71	0.09		1			
Staphylococcus gallinarum		2.00	0.04		78			
205771	**Low discrimination**							**205771**
					OXI			
Staphylococcus equorum		56.04	0.50	GLR 90	1			
Staphylococcus gallinarum		41.39	0.45	NIT 92	1			
Staphylococcus sciuri		1.67	0.22		99			
205772	**Low discrimination**							**205772**
					YEP			
Staphylococcus kloosii		76.69	0.61	SUC 9	99			
Staphylococcus arlettae		14.41	0.35		99			
Staphylococcus equorum		5.22	0.17		1			
205773	**Low discrimination**							**205773**
					YEP			
Staphylococcus equorum		56.81	0.59	GLR 90	1			
Staphylococcus gallinarum		39.95	0.53	NIT 92	78			
Staphylococcus arlettae		2.10	0.37		99			
206000	**Good identification**							**206000**
					OXI			
Dermacoccus nishinomiyaensis		90.12	0.69	PHS 5	93			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		8.66	0.55		13			
Micrococcus sp.		0.87	0.18		89			
206001	**Low discrimination**							**206001**
					OXI			
Dermacoccus nishinomiyaensis		53.56	0.28	PHS 5	93			
Kocuria varians		44.28	0.34	MNS 1	13			
Staphylococcus capitis subsp. capitis		2.16	0.00		1			
206002	**Low discrimination**							**206002**
					OXI			
Kocuria varians		72.84	0.44	PHS 1	13			
Dermacoccus nishinomiyaensis		27.16	0.28		93			
206010	**Low discrimination**							**206010**
					OXI	YEP	NOV	
Dermacoccus nishinomiyaensis		73.82	0.28	PHS 5	93	1	80	
Macrococcus bovicus		13.36	0.16	MAN 1	99	99	99	
Kocuria varians		7.09	0.14		13	80	1	
206020	**Low discrimination**							**206020**
					OXI			
Kocuria varians		74.09	0.46	PHS 1	13			
Dermacoccus nishinomiyaensis		22.18	0.28		93			
Kocuria rosea		3.73	0.10		35			
206022	**Good identification**							**206022**
Kocuria varians		100.0	0.35	PHS 1				
206040	**Low discrimination**							**206040**
					OXI			
Kocuria varians		64.31	0.42	PHS 1	13			
Dermacoccus nishinomiyaensis		30.81	0.28		93			
Staphylococcus hominis subsp. hominis		2.67	0.15		1			
206042	**Low discrimination**							**206042**
Kocuria varians		78.42	0.30	PHS 1				
Staphylococcus hominis subsp. hominis		13.53	0.16					
Staphylococcus epidermidis		8.05	0.08					
206060	**Good identification**							**206060**
Kocuria varians		100.0	0.32	PHS 1				
206100	**Low discrimination**							**206100**
					OXI			
Dermacoccus nishinomiyaensis		56.94	0.54	PHS 5	93			
Kocuria varians		43.07	0.59		13			
206101	**Low discrimination**							**206101**
					OXI	YEP		
Kocuria varians		84.94	0.37	PHS 1	13	80		
Dermacoccus nishinomiyaensis		13.05	0.13	MNS 8	93	1		
Staphylococcus chromogenes		2.01	-.02		1	95		
206102	**Good identification**							**206102**
					OXI			
Kocuria varians		95.48	0.48	PHS 1	13			
Dermacoccus nishinomiyaensis		4.52	0.13		93			
206120	**Good identification**							**206120**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		96.34	0.50	PHS 1	13		
Dermacoccus nishinomiyaensis		3.66	0.13		93		
206121	**Good identification**						**206121**
Kocuria varians		100.0	0.28	PHS 1			
206122	**Good identification**						**206122**
Kocuria varians		100.0	0.38	PHS 1			
206140	**Low discrimination**						**206140**
					OXI		
Kocuria varians		87.66	0.45	PHS 1	13		
Dermacoccus nishinomiyaensis		5.33	0.13		93		
Staphylococcus epidermidis		3.03	0.13		1		
206142	**Low discrimination**						**206142**
Kocuria varians		64.85	0.34	PHS 1			
Staphylococcus epidermidis		25.16	0.24				
Staphylococcus hominis subsp. hominis		8.08	0.17				
206143	**Low discrimination**						**206143**
					YEP		
Staphylococcus epidermidis		65.07	0.29	VPT 96	1		
Staphylococcus chromogenes		27.07	0.17	SUC 99	95		
Kocuria varians		7.85	0.12		80		
206150	**Low discrimination**						**206150**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		75.87	0.31	TRE 99	1	99	
Macrococcus bovicus		17.33	0.16		99	1	
Macrococcus equipersicus		3.77	0.04		99	1	
206160	**Good identification**						**206160**
Kocuria varians		100.0	0.36	PHS 1			
206200	**Low discrimination**						**206200**
					OXI		
Dermacoccus nishinomiyaensis		80.60	0.44	PHS 5	93		
Kocuria varians		16.54	0.38	SUC 6	13		
Micrococcus sp.		1.20	-.03		89		
206203	**Low discrimination**						**206203**
					YEP		
Staphylococcus chromogenes		84.81	0.39	ARG 90	95		
Staphylococcus hyicus		8.41	0.14	GAL 90	1		
Staphylococcus epidermidis		3.27	0.14	TRE 92	1		
206211	**Acceptable identification**						**206211**
					YEP		
Staphylococcus capitis subsp. capitis		91.61	0.29	URE 1	1		
Staphylococcus chromogenes		8.39	0.08	PHS 11	95		
206213	**Low discrimination**						**206213**
					COA	YEP	
Staphylococcus chromogenes		41.43	0.28	ARG 90	1	95	
Staphylococcus delphini		35.90	0.18	GAL 90	99	1	
Staphylococcus felis		11.90	0.11	TRE 92	1	1	
206240	**Low discrimination**						**206240**
					NOV		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		65.70	0.56	PHS 3	1			
Staphylococcus epidermidis		14.49	0.39		1			
Staphylococcus hominis subsp. novobiosepticus		12.23	0.30		99			
206241	**Low discrimination**							**206241**
					NOV	YEP		
Staphylococcus epidermidis		79.20	0.44	VPT 96	1	1		
Staphylococcus hominis subsp. novobiosepticus		8.99	0.17		99	1		
Staphylococcus chromogenes		7.30	0.19		1	95		
206242	**Low discrimination**							**206242**
					NOV			
Staphylococcus hominis subsp. hominis		59.12	0.57	PHS 3	1			
Staphylococcus epidermidis		35.19	0.49		1			
Staphylococcus hominis subsp. novobiosepticus		5.05	0.24		99			
206243	**Low discrimination**							**206243**
					COA	YEP		
Staphylococcus epidermidis		73.21	0.55	VPT 96	1	1		
Staphylococcus chromogenes		19.18	0.39		1	95		
Staphylococcus delphini		4.69	0.18		99	1		
206250	**Low discrimination**							**206250**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		58.77	0.39	PHS 3	1	1	1	
Staphylococcus saprophyticus subsp. bovis		36.38	0.31	MAN 13	1	99	99	
Macrococcus bovicus		4.85	0.11		99	1	99	
206252	**Low discrimination**							**206252**
					COA	NOV	PYR	
Staphylococcus hominis subsp. hominis		60.08	0.40	PHS 3	1	1	1	
Staphylococcus delphini		28.44	0.18	MAN 13	99	1	15	
Staphylococcus saprophyticus subsp. bovis		7.92	0.18		1	99	99	
206253	**Very good identification**							**206253**
					COA	YEP		
Staphylococcus delphini		98.23	0.59	ARG 99	99	1		
Staphylococcus chromogenes		1.15	0.28		1	95		
Staphylococcus capitis subsp. ureolyticus		0.22	0.03		1	77		
206300	**Low discrimination**							**206300**
					OXI			
Kocuria varians		56.73	0.41	PHS 1	13			
Dermacoccus nishinomiyaensis		35.11	0.29	SUC 12	93			
Staphylococcus epidermidis		3.16	0.13		1			
206301	**Low discrimination**							**206301**
					YEP			
Staphylococcus chromogenes		79.98	0.39	ARG 90	95			
Staphylococcus hyicus		7.94	0.14	TRE 92	1			
Staphylococcus epidermidis		5.52	0.19	LAC 90	1			
206302	**Low discrimination**							**206302**
					YEP			
Kocuria varians		41.39	0.30	PHS 1	80			
Staphylococcus epidermidis		25.85	0.24	SUC 12	1			
Staphylococcus chromogenes		19.97	0.17		95			
206303	**Low discrimination**							**206303**
					YEP			
Staphylococcus chromogenes		85.12	0.59	ARG 90	95			
Staphylococcus hyicus		10.79	0.36	TRE 92	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		2.07	0.29		1			
206311	**Low discrimination**							**206311**
					COA	YEP		
Staphylococcus chromogenes		81.27	0.28	ARG 90	1	95		
Staphylococcus intermedius		18.73	0.16	TRE 92	99	1		
206313	**Low discrimination**							**206313**
					COA	YEP		
Staphylococcus chromogenes		73.07	0.47	ARG 90	1	95		
Staphylococcus intermedius		11.49	0.32	TRE 92	99	1		
Staphylococcus felis		7.81	0.22		1	1		
206320	**Good identification**							**206320**
Kocuria varians		100.0	0.32	PHS 1				
206340	**Low discrimination**							**206340**
					NOV			
Staphylococcus epidermidis		42.40	0.54	VPT 96	1			
Staphylococcus hominis subsp. hominis		36.74	0.57		1			
Staphylococcus hominis subsp. novobiosepticus		18.94	0.39		99			
206341	**Low discrimination**							**206341**
					NOV	YEP		
Staphylococcus epidermidis		81.76	0.60	VPT 96	1	1		
Staphylococcus chromogenes		11.97	0.39		1	95		
Staphylococcus hominis subsp. novobiosepticus		4.92	0.27		99	1		
206342	**Low discrimination**							**206342**
					NOV			
Staphylococcus epidermidis		71.00	0.65	VPT 96	1			
Staphylococcus hominis subsp. hominis		22.81	0.58		1			
Staphylococcus hominis subsp. novobiosepticus		5.39	0.34		99			
206343	**Low discrimination**							**206343**
					COA	YEP		
Staphylococcus epidermidis		68.88	0.70	VPT 96	1	1		
Staphylococcus chromogenes		28.66	0.59		1	95		
Staphylococcus delphini		0.78	0.18		99	1		
206350	**Low discrimination**							**206350**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		66.62	0.43	TRE 99	99	99		
Staphylococcus hominis subsp. hominis		29.15	0.40		1	1		
Staphylococcus epidermidis		2.27	0.13		1	1		
206351	**Low discrimination**							**206351**
					COA	NOV	YEP	
Staphylococcus chromogenes		28.94	0.28	ARG 90	1	1	95	
Staphylococcus saprophyticus subsp. bovis		27.92	0.31	TRE 92	1	99	80	
Staphylococcus delphini		25.07	0.18	LAC 90	99	1	1	
206352	**Low discrimination**							**206352**
					COA	NOV	PYR	
Staphylococcus hominis subsp. hominis		45.54	0.41	PHS 3	1	1	1	
Staphylococcus saprophyticus subsp. bovis		22.17	0.31	MAN 13	1	99	99	
Staphylococcus delphini		19.90	0.18		99	1	15	
206353	**Low discrimination**							**206353**
					COA	YEP		
Staphylococcus delphini		87.32	0.59	ARG 99	99	1		
Staphylococcus chromogenes		9.16	0.47		1	95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus intermedius		2.16	0.36		99	1	
206363	**Low discrimination**						**206363**
					COA	YEP	
Staphylococcus epidermidis		54.24	0.29	VPT 96	1	1	
Staphylococcus chromogenes		22.56	0.17	XYL 1	1	95	
Staphylococcus lutrae		20.83	0.12		99	1	
206400	**Low discrimination**						**206400**
					OXI		
Dermacoccus nishinomiyaensis		53.31	0.28	PHS 5	93		
Kocuria varians		38.16	0.32	TRE 1	13		
Staphylococcus piscifermentans		6.50	0.25		1		
206403	**Low discrimination**						**206403**
					YEP		
Staphylococcus felis		80.59	0.29	ARG 92	1		
Staphylococcus chromogenes		15.86	0.20	MAN 87	95		
Staphylococcus simulans		1.98	0.04		1		
206413	**Good identification**						**206413**
					NOV		
Staphylococcus felis		96.02	0.46	ARG 92	1		
Staphylococcus cohnii subsp. urealyticum		1.77	0.24		99		
Staphylococcus simulans		1.41	0.17		1		
206450	**Good identification**						**206450**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		92.60	0.59		1	99	
Staphylococcus kloosii		3.17	0.46		1	52	
Macrococcus bovicus		2.99	0.27		99	1	
206451	**Good identification**						**206451**
					PYR		
Staphylococcus saprophyticus subsp. bovis		96.05	0.47		99		
Staphylococcus cohnii subsp. urealyticum		2.33	0.19		53		
Staphylococcus cohnii subsp. cohnii		1.62	0.15		1		
206452	**Low discrimination**						**206452**
					NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		85.17	0.47		99	80	
Staphylococcus kloosii		12.64	0.47		99	99	
Staphylococcus piscifermentans		1.36	0.21		1	1	
206453	**Low discrimination**						**206453**
					NOV		
Staphylococcus saprophyticus subsp. bovis		45.12	0.34		99		
Staphylococcus cohnii subsp. urealyticum		44.30	0.40		99		
Staphylococcus felis		4.28	0.04		1		
206472	**Low discrimination**						**206472**
Staphylococcus kloosii		78.60	0.35	NIT 1			
Staphylococcus saprophyticus subsp. bovis		21.40	0.06				
206500	**Low discrimination**						**206500**
					OXI		
Kocuria varians		74.20	0.36	PHS 1	13		
Dermacoccus nishinomiyaensis		13.17	0.13	TRE 7	93		
Staphylococcus piscifermentans		12.64	0.28		1		
206503	**Low discrimination**						**206503**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus felis		62.09	0.39	ARG 92	1			
Staphylococcus chromogenes		32.84	0.39	MAN 87	95			
Staphylococcus hyicus		4.16	0.17		1			
206510	**Low discrimination**							**206510**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		75.06	0.31	MLT 99	1	99	99	
Macrococcus bovicus		11.02	0.12		99	99	1	
Staphylococcus piscifermentans		7.42	0.21		1	1	15	
206513	**Very good identification**							**206513**
					YEP			
Staphylococcus felis		95.72	0.56	ARG 92	1			
Staphylococcus chromogenes		2.13	0.28		95			
Staphylococcus condimenti		1.42	0.14		1			
206520	**Good identification**							**206520**
Kocuria varians		100.0	0.26	PHS 1				
206540	**Low discrimination**							**206540**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		52.56	0.31	MAN 99	99	99		
Staphylococcus hominis subsp. hominis		19.69	0.26		1	1		
Kocuria varians		15.63	0.22		1	10		
206542	**Low discrimination**							**206542**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		37.96	0.27	PHS 3	1	1		
Staphylococcus piscifermentans		29.87	0.32	SUC 99	1	15		
Staphylococcus saprophyticus subsp. bovis		21.58	0.18		99	99		
206543	**Low discrimination**							**206543**
					COA	PYR	YEP	
Staphylococcus chromogenes		85.33	0.39	ARG 90	1	45	95	
Staphylococcus lutrae		6.85	0.12	SUC 99	99	10	1	
Staphylococcus intermedius		2.43	0.09		99	90	1	
206550	**Very good identification**							**206550**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		97.83	0.72		1	99		
Staphylococcus kloosii		1.63	0.52		1	52		
Macrococcus bovicus		0.34	0.19		99	1		
206551	**Very good identification**							**206551**
					NOV			
Staphylococcus saprophyticus subsp. bovis		98.90	0.59		99			
Staphylococcus chromogenes		0.48	0.08		1			
Staphylococcus cohnii subsp. urealyticum		0.26	0.12		99			
206552	**Good identification**							**206552**
					NOV	YEP		
Staphylococcus saprophyticus subsp. bovis		92.16	0.59		99	80		
Staphylococcus kloosii		6.64	0.53		99	99		
Staphylococcus piscifermentans		0.55	0.25		1	1		
206553	**Low discrimination**							**206553**
					NOV	YEP		
Staphylococcus saprophyticus subsp. bovis		59.59	0.47		99	80		
Staphylococcus chromogenes		10.44	0.28		1	95		
Staphylococcus caprae		7.76	0.18		1	1		
206563	**Very good identification**							**206563**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	NOV	YEP	
Staphylococcus lutrae		99.73	0.53	BGA 99	99	1	1	
Staphylococcus xylosus		0.15	-.01		1	99	40	
Staphylococcus chromogenes		0.13	-.02		1	1	95	
206570	**Low discrimination**							**206570**
Staphylococcus saprophyticus subsp. bovis		70.86	0.31	XYL 1				
Staphylococcus kloosii		29.15	0.40					
206573	**Very good identification**							**206573**
					COA	NOV		
Staphylococcus lutrae		98.34	0.59	BGA 99	99	1		
Staphylococcus xylosus		1.15	0.24		1	99		
Staphylococcus gallinarum		0.35	0.12		1	99		
206601	**Acceptable identification**							**206601**
					YEP			
Staphylococcus chromogenes		91.99	0.41	ARG 90	95			
Staphylococcus hyicus		7.14	0.14	GAL 90	1			
Staphylococcus felis		0.87	-.06	LAC 90	1			
206603	**Low discrimination**							**206603**
					YEP			
Staphylococcus chromogenes		80.94	0.61	ARG 90	95			
Staphylococcus felis		8.44	0.35	GAL 90	1			
Staphylococcus hyicus		8.03	0.36		1			
206610	**Low discrimination**							**206610**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		78.10	0.31	MLT 99	1	99	99	
Staphylococcus piscifermentans		7.72	0.21		1	1	15	
Micrococcus bovicus		6.69	0.07		99	99	1	
206611	**Low discrimination**							**206611**
					NOV	YEP		
Staphylococcus chromogenes		66.67	0.30	ARG 90	1	95		
Staphylococcus felis		14.99	0.11	GAL 90	1	1		
Staphylococcus saprophyticus subsp. bovis		12.59	0.18	LAC 90	99	80		
206612	**Low discrimination**							**206612**
					NOV			
Staphylococcus simulans		63.54	0.34	ARG 92	1			
Staphylococcus felis		12.38	0.11	BGA 90	1			
Staphylococcus saprophyticus subsp. bovis		10.39	0.18		99			
206613	**Low discrimination**							**206613**
					YEP			
Staphylococcus felis		62.97	0.52	ARG 92	1			
Staphylococcus chromogenes		25.46	0.49		95			
Staphylococcus simulans		10.93	0.45		1			
206620	**Very good identification**							**206620**
Staphylococcus muscae		99.97	0.59	URE 1				
Kocuria varians		0.03	0.05					
206640	**Good identification**							**206640**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		94.90	0.66	PHS 3	1	1		
Staphylococcus saprophyticus subsp. bovis		2.77	0.31		99	99		
Staphylococcus warneri		1.27	0.33		1	8		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
206641	**Low discrimination**							**206641**
					NOV	PYR		
Staphylococcus chromogenes		81.96	0.41	ARG 90	1	45		
Staphylococcus hominis subsp. hominis		12.20	0.31	GAL 90	1	1		
Staphylococcus saprophyticus subsp. bovis		4.36	0.18	LAC 90	99	99		
206642	**Very good identification**							**206642**
					YEP			
Staphylococcus hominis subsp. hominis		96.58	0.68	PHS 3	49			
Staphylococcus chromogenes		1.03	0.20		95			
Staphylococcus piscifermentans		0.83	0.32		1			
206643	**Very good identification**							**206643**
					YEP			
Staphylococcus chromogenes		95.93	0.61	ARG 90	95			
Staphylococcus hominis subsp. hominis		1.86	0.33	GAL 90	49			
Staphylococcus simulans		1.31	0.26		1			
206650	**Good identification**							**206650**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		94.44	0.72		99	99		
Staphylococcus hominis subsp. hominis		4.88	0.49		1	1		
Staphylococcus warneri		0.32	0.30		1	8		
206651	**Good identification**							**206651**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		93.82	0.59		1	99	99	
Staphylococcus chromogenes		5.02	0.30		1	1	45	
Staphylococcus sciuri		0.44	0.12		99	99	1	
206652	**Low discrimination**							**206652**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		77.56	0.59		99	99		
Staphylococcus hominis subsp. hominis		18.82	0.51		1	1		
Staphylococcus simulans		2.15	0.27		1	70		
206653	**Low discrimination**							**206653**
					NOV	YEP		
Staphylococcus chromogenes		52.73	0.49	ARG 90	1	95		
Staphylococcus saprophyticus subsp. bovis		27.38	0.47	GAL 90	99	80		
Staphylococcus simulans		10.17	0.38		1	1		
206670	**Good identification**							**206670**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		95.09	0.31	XYL 1	99	99		
Staphylococcus hominis subsp. hominis		4.91	0.08		1	1		
206673	**Low discrimination**							**206673**
					COA	NOV		
Staphylococcus xylosus		49.59	0.29	GLR 95	1	99		
Staphylococcus lutrae		22.72	0.18	GAL 91	99	1		
Staphylococcus equorum		12.83	0.19		1	99		
206701	**Low discrimination**							**206701**
					YEP	PYR	COA	
Staphylococcus chromogenes		88.46	0.61	ARG 90	95	45	1	
Staphylococcus hyicus		8.78	0.36	LAC 90	1	1	24	
Staphylococcus intermedius		2.46	0.30		1	90	99	
206702	**Low discrimination**							**206702**
					YEP			
Staphylococcus chromogenes		68.67	0.39	ARG 90	95			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		9.28	0.36	MNS 99	1			
Staphylococcus hyicus		8.71	0.17		1			
206703	**Low discrimination**							**206703**
					YEP			
Staphylococcus chromogenes		84.40	0.80	ARG 90	95			
Staphylococcus hyicus		10.70	0.58		1			
Staphylococcus felis		3.27	0.46		1			
206710	**Acceptable identification**							**206710**
					NOV	YEP		
Staphylococcus saprophyticus subsp. bovis		91.05	0.43	MLT 99	99	80		
Staphylococcus piscifermentans		3.38	0.25		1	1		
Staphylococcus chromogenes		2.74	0.08		1	95		
206711	**Low discrimination**							**206711**
					COA	NOV	YEP	
Staphylococcus chromogenes		69.21	0.49	ARG 90	1	1	95	
Staphylococcus intermedius		18.42	0.39	LAC 90	99	1	1	
Staphylococcus saprophyticus subsp. bovis		5.81	0.31		1	99	80	
206712	**Low discrimination**							**206712**
					NOV	YEP		
Staphylococcus chromogenes		29.30	0.28	ARG 90	1	95		
Staphylococcus saprophyticus subsp. bovis		27.04	0.31	MNS 99	99	80		
Staphylococcus felis		26.95	0.22		1	1		
206713	**Low discrimination**							**206713**
					COA	YEP		
Staphylococcus chromogenes		47.05	0.69	ARG 90	1	95		
Staphylococcus felis		43.28	0.63		1	1		
Staphylococcus intermedius		8.55	0.55		99	1		
206723	**Low discrimination**							**206723**
					YEP	COA		
Staphylococcus chromogenes		78.05	0.39	ARG 90	95	1		
Staphylococcus hyicus		9.89	0.17	XYL 1	1	24		
Staphylococcus lutrae		6.27	0.12		1	99		
206733	**Low discrimination**							**206733**
					NOV	YEP		
Staphylococcus chromogenes		24.53	0.28	ARG 90	1	95		
Staphylococcus felis		22.57	0.22	XYL 1	1	1		
Staphylococcus xylosus		22.51	0.27		99	40		
206740	**Low discrimination**							**206740**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		87.80	0.67	PHS 3	1	1		
Staphylococcus saprophyticus subsp. bovis		9.47	0.43		99	99		
Staphylococcus chromogenes		1.01	0.20		1	45		
206741	**Good identification**							**206741**
					COA	NOV	YEP	
Staphylococcus chromogenes		91.72	0.61	ARG 90	1	1	95	
Staphylococcus intermedius		3.82	0.34	LAC 90	99	1	1	
Staphylococcus saprophyticus subsp. bovis		2.17	0.31		1	99	80	
206742	**Low discrimination**							**206742**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		87.92	0.68	PHS 3	1	1		
Staphylococcus chromogenes		7.76	0.39		1	45		
Staphylococcus saprophyticus subsp. bovis		2.02	0.31		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
206743	**Very good identification**							**206743**
					COA	YEP	PYR	
Staphylococcus chromogenes		96.44	0.80	ARG 90	1	95	45	
Staphylococcus intermedius		2.74	0.50		99	1	90	
Staphylococcus hominis subsp. hominis		0.23	0.33		1	49	1	
206750	**Very good identification**							**206750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		98.46	0.84		99	99		
Staphylococcus hominis subsp. hominis		1.38	0.50		1	1		
Staphylococcus kloosii		0.04	0.32		99	52		
206751	**Low discrimination**							**206751**
					COA	NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		84.82	0.72		1	99	80	
Staphylococcus chromogenes		10.21	0.49		1	1	95	
Staphylococcus intermedius		4.08	0.43		99	1	1	
206752	**Good identification**							**206752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		91.98	0.72		99	99		
Staphylococcus hominis subsp. hominis		6.05	0.51		1	1		
Staphylococcus chromogenes		1.01	0.28		1	45		
206753	**Low discrimination**							**206753**
					COA	NOV	YEP	
Staphylococcus chromogenes		57.47	0.69	ARG 90	1	1	95	
Staphylococcus intermedius		15.66	0.59		99	1	1	
Staphylococcus saprophyticus subsp. bovis		13.26	0.59		1	99	80	
206762	**Low discrimination**							**206762**
					COA	YEP		
Staphylococcus hominis subsp. hominis		55.88	0.27	PHS 3	1	49		
Staphylococcus lutrae		39.19	0.12	XYL 1	99	1		
Staphylococcus chromogenes		4.93	-.02		1	95		
206763	**Low discrimination**							**206763**
					COA	NOV	YEP	
Staphylococcus lutrae		81.62	0.53	BGA 99	99	1	1	
Staphylococcus chromogenes		10.27	0.39		1	1	95	
Staphylococcus gallinarum		5.47	0.32		1	99	78	
206770	**Acceptable identification**							**206770**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		94.59	0.43	XYL 1	99	99		
Staphylococcus gallinarum		2.02	0.04		99	12		
Staphylococcus hominis subsp. hominis		1.32	0.09		1	1		
206771	**Low discrimination**							**206771**
Staphylococcus gallinarum		51.67	0.45	ESL 99				
Staphylococcus xylosus		28.38	0.41					
Staphylococcus equorum		8.13	0.31					
206772	**Low discrimination**							**206772**
					COA	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		47.95	0.31	XYL 1	1	99	99	
Staphylococcus lutrae		28.70	0.18		99	10	1	
Staphylococcus gallinarum		10.02	0.12		1	12	99	
206773	**Low discrimination**							**206773**
					COA	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lutrae		60.18	0.59	BGA 99	99	1		
Staphylococcus gallinarum		21.02	0.53		1	99		
Staphylococcus xylosus		13.42	0.50		1	99		
207000	**Acceptable identification**							**207000**
					OXI			
Dermacoccus nishinomiyaensis		91.24	0.28	ESL 1	93			
Kocuria varians		8.76	0.14	PHS 5	13			
207343	**Low discrimination**							**207343**
					NOV	YEP		
Staphylococcus epidermidis		66.49	0.29	VPT 96	1	1		
Staphylococcus chromogenes		27.66	0.17	ESL 1	1	95		
Staphylococcus hominis subsp. novobiosepticus		5.85	-.01		99	1		
207351	**Low discrimination**							**207351**
					OXI			
Staphylococcus sciuri		88.90	0.28	URE 1	99			
Staphylococcus gallinarum		11.10	0.04	TRE 97	1			
207363	**Good identification**							**207363**
Staphylococcus gallinarum		98.34	0.32	TRE 99				
Staphylococcus equorum		1.66	-.02	MAN 91				
207371	**Low discrimination**							**207371**
					OXI			
Staphylococcus gallinarum		81.70	0.45	TRE 99	1			
Staphylococcus equorum		12.85	0.31		1			
Staphylococcus sciuri		4.99	0.25		99			
207373	**Low discrimination**							**207373**
Staphylococcus gallinarum		84.88	0.53	TRE 99				
Staphylococcus equorum		14.02	0.39					
Staphylococcus xylosus		0.57	0.09					
207400	**Good identification**							**207400**
Staphylococcus piscifermentans		100.0	0.44	ARG 99				
207402	**Good identification**							**207402**
Staphylococcus piscifermentans		100.0	0.48	ARG 99				
207410	**Low discrimination**							**207410**
					OXI	NOV	YEP	
Staphylococcus piscifermentans		77.34	0.37	ARG 99	1	1	1	
Macrococcus equipersicus		11.88	0.09		99	99	99	
Staphylococcus kloosii		5.82	0.20		1	99	99	
207412	**Acceptable identification**							**207412**
					NOV	YEP		
Staphylococcus piscifermentans		94.85	0.41	ARG 99	1	1		
Staphylococcus kloosii		5.15	0.21		99	99		
207440	**Good identification**							**207440**
Staphylococcus piscifermentans		100.0	0.44	ARG 99				
207442	**Good identification**							**207442**
Staphylococcus piscifermentans		100.0	0.48	ARG 99				
207450	**Low discrimination**							**207450**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus kloosii		41.83	0.46	NIT 1	99	99		
Staphylococcus piscifermentans		29.28	0.37		1	1		
Staphylococcus saprophyticus subsp. bovis		12.34	0.18		99	80		
207452	**Low discrimination**							**207452**
					NOV	YEP		
Staphylococcus kloosii		49.09	0.47	NIT 1	99	99		
Staphylococcus piscifermentans		47.57	0.41		1	1		
Staphylococcus saprophyticus subsp. bovis		3.34	0.06		99	80		
207453	**Good identification**							**207453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	GLR 99				
207470	**Good identification**							**207470**
Staphylococcus kloosii		100.0	0.34	NIT 1				
207472	**Good identification**							**207472**
Staphylococcus kloosii		100.0	0.35	NIT 1				
207500	**Good identification**							**207500**
Staphylococcus piscifermentans		100.0	0.48	ARG 99				
207502	**Good identification**							**207502**
Staphylococcus piscifermentans		100.0	0.52	ARG 99				
207510	**Low discrimination**							**207510**
					OXI	NOV	YEP	
Staphylococcus piscifermentans		87.63	0.41	ARG 99	1	1	1	
Staphylococcus kloosii		8.53	0.26		1	99	99	
Micrococcus equipersicus		3.85	0.01		99	99	99	
207512	**Acceptable identification**							**207512**
					NOV	YEP		
Staphylococcus piscifermentans		93.43	0.44	ARG 99	1	1		
Staphylococcus kloosii		6.57	0.27		99	99		
207540	**Good identification**							**207540**
Staphylococcus piscifermentans		100.0	0.48	ARG 99				
207542	**Good identification**							**207542**
Staphylococcus piscifermentans		100.0	0.52	ARG 99				
207550	**Low discrimination**							**207550**
					NOV	YEP		
Staphylococcus kloosii		44.72	0.52	NIT 1	99	99		
Staphylococcus saprophyticus subsp. bovis		27.18	0.31		99	80		
Staphylococcus piscifermentans		24.19	0.41		1	1		
207552	**Low discrimination**							**207552**
					NOV	YEP		
Staphylococcus kloosii		52.93	0.53	NIT 1	99	99		
Staphylococcus piscifermentans		39.64	0.44		1	1		
Staphylococcus saprophyticus subsp. bovis		7.42	0.18		99	80		
207563	**Low discrimination**							**207563**
					COA	NOV		
Staphylococcus gallinarum		85.65	0.32	SUC 99	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lutrae		12.91	0.12	MAN 91	99	1	
Staphylococcus equorum		1.45	-.02		1	99	
207570	**Low discrimination**						**207570**
Staphylococcus kloosii		82.81	0.40	NIT 1			
Staphylococcus gallinarum		17.19	0.04				
207571	**Low discrimination**						**207571**
					OXI		
Staphylococcus gallinarum		84.43	0.45	SUC 99	1		
Staphylococcus equorum		13.28	0.31		1		
Staphylococcus sciuri		1.68	0.15		99		
207572	**Low discrimination**						**207572**
					YEP		
Staphylococcus kloosii		64.66	0.41	NIT 1	99		
Staphylococcus gallinarum		30.33	0.12		78		
Staphylococcus equorum		5.01	-.02		1		
207573	**Low discrimination**						**207573**
					COA	NOV	
Staphylococcus gallinarum		83.01	0.53	SUC 99	1	99	
Staphylococcus equorum		13.71	0.39		1	99	
Staphylococcus lutrae		2.40	0.18		99	1	
207600	**Good identification**						**207600**
Staphylococcus piscifermentans		100.0	0.48	ARG 99			
207602	**Good identification**						**207602**
Staphylococcus piscifermentans		100.0	0.52	ARG 99			
207610	**Low discrimination**						**207610**
					OXI	NOV	
Staphylococcus piscifermentans		64.31	0.41	ARG 99	1	1	
Staphylococcus sciuri		27.05	0.23		99	99	
Staphylococcus vitulinus		8.64	0.04		99	99	
207611	**Good identification**						**207611**
Staphylococcus sciuri		100.0	0.41	URE 1			
207612	**Good identification**						**207612**
					OXI	NOV	
Staphylococcus piscifermentans		95.63	0.44	ARG 99	1	1	
Staphylococcus sciuri		4.37	0.06		99	99	
207631	**Low discrimination**						**207631**
					OXI		
Staphylococcus sciuri		62.47	0.39	URE 1	99		
Staphylococcus equorum		35.20	0.31		1		
Staphylococcus gallinarum		2.33	0.04		1		
207633	**Low discrimination**						**207633**
					OXI		
Staphylococcus equorum		85.07	0.39	GLR 90	1		
Staphylococcus sciuri		9.56	0.22	GAL 91	99		
Staphylococcus gallinarum		5.37	0.12	MLT 91	1		
207640	**Low discrimination**						**207640**
					OXI	NOV	
Staphylococcus piscifermentans		84.21	0.48	ARG 99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		14.02	0.25		1	1		
Staphylococcus sciuri		1.77	0.03		99	99		
207642	**Low discrimination**							**207642**
Staphylococcus piscifermentans		88.47	0.52	ARG 99				
Staphylococcus hominis subsp. hominis		11.53	0.26					
207650	**Low discrimination**							**207650**
					OXI	NOV	PYR	
Staphylococcus sciuri		41.01	0.34	URE 1	99	99	1	
Staphylococcus saprophyticus subsp. bovis		29.13	0.31	MNS 89	1	99	99	
Staphylococcus piscifermentans		25.92	0.41		1	1	15	
207651	**Very good identification**							**207651**
					OXI	PYR		
Staphylococcus sciuri		96.06	0.53	URE 1	99	1		
Staphylococcus saprophyticus subsp. bovis		2.11	0.18		1	99		
Staphylococcus equorum		1.11	0.10		1	5		
207652	**Low discrimination**							**207652**
					OXI	NOV	PYR	
Staphylococcus piscifermentans		67.91	0.44	ARG 99	1	1	15	
Staphylococcus saprophyticus subsp. bovis		12.72	0.18		1	99	99	
Staphylococcus sciuri		11.66	0.18		99	99	1	
207653	**Low discrimination**							**207653**
					OXI			
Staphylococcus sciuri		70.51	0.37	URE 1	99			
Staphylococcus equorum		12.85	0.19	LAC 14	1			
Staphylococcus gallinarum		7.95	0.12		1			
207663	**Low discrimination**							**207663**
					OXI			
Staphylococcus gallinarum		84.29	0.32	GAL 99	1			
Staphylococcus equorum		13.92	0.19	MAN 91	1			
Staphylococcus sciuri		1.78	0.03		99			
207670	**Low discrimination**							**207670**
					OXI			
Staphylococcus sciuri		83.05	0.32	URE 1	99			
Staphylococcus equorum		10.28	0.10	MNS 89	1			
Staphylococcus gallinarum		6.67	0.04		1			
207671	**Low discrimination**							**207671**
					OXI			
Staphylococcus equorum		43.13	0.51	GLR 90	1			
Staphylococcus sciuri		28.48	0.51	GAL 91	99			
Staphylococcus gallinarum		28.01	0.45		1			
207673	**Low discrimination**							**207673**
					OXI			
Staphylococcus equorum		59.84	0.60	GLR 90	1			
Staphylococcus gallinarum		37.00	0.53	GAL 91	1			
Staphylococcus sciuri		2.50	0.34		99			
207700	**Good identification**							**207700**
Staphylococcus piscifermentans		100.0	0.52	ARG 99				
207702	**Very good identification**							**207702**
					YEP			
Staphylococcus piscifermentans		99.18	0.55	ARG 99	1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus chromogenes		0.82	-.02		95			
207703	**Low discrimination**							**207703**
					YEP			
Staphylococcus chromogenes		83.02	0.39	ARG 90	95			
Staphylococcus hyicus		10.53	0.17	ESL 1	1			
Staphylococcus felis		3.22	0.05		1			
207710	**Low discrimination**							**207710**
					OXI	NOV	PYR	
Staphylococcus piscifermentans		63.51	0.44	ARG 99	1	1	15	
Staphylococcus sciuri		34.57	0.28		99	99	1	
Staphylococcus saprophyticus subsp. bovis		1.92	0.02		1	99	99	
207711	**Low discrimination**							**207711**
					OXI			
Staphylococcus sciuri		85.46	0.47	URE 1	99			
Staphylococcus lentus		9.17	0.29		99			
Staphylococcus equorum		1.91	0.10		1			
207712	**Acceptable identification**							**207712**
					OXI	NOV		
Staphylococcus piscifermentans		94.42	0.48	ARG 99	1	1		
Staphylococcus sciuri		5.58	0.12		99	99		
207713	**Low discrimination**							**207713**
					OXI	YEP	NOV	
Staphylococcus lentus		37.18	0.40	URE 1	99	1	99	
Staphylococcus chromogenes		18.00	0.28		1	95	1	
Staphylococcus felis		16.56	0.22		1	1	1	
207723	**Low discrimination**							**207723**
					NOV	YEP		
Staphylococcus gallinarum		84.45	0.32	MAN 91	99	78		
Staphylococcus equorum		13.95	0.19	MLT 99	99	1		
Staphylococcus chromogenes		1.60	-.02		1	95		
207731	**Low discrimination**							**207731**
					OXI			
Staphylococcus equorum		48.89	0.51	GLR 90	1			
Staphylococcus gallinarum		31.75	0.45	MLT 91	1			
Staphylococcus sciuri		16.66	0.45		99			
207733	**Low discrimination**							**207733**
					OXI			
Staphylococcus equorum		58.34	0.60	GLR 90	1			
Staphylococcus gallinarum		36.07	0.53	MLT 91	1			
Staphylococcus lentus		3.90	0.40		99			
207740	**Low discrimination**							**207740**
					OXI	NOV		
Staphylococcus piscifermentans		86.18	0.52	ARG 99	1	1		
Staphylococcus hominis subsp. hominis		10.37	0.26		1	1		
Staphylococcus sciuri		2.34	0.09		99	99		
207742	**Good identification**							**207742**
					YEP			
Staphylococcus piscifermentans		90.71	0.55	ARG 99	1			
Staphylococcus hominis subsp. hominis		8.54	0.27		49			
Staphylococcus chromogenes		0.75	-.02		95			
207743	**Low discrimination**							**207743**
					OXI	NOV		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus chromogenes		61.60	0.39	ARG 90	1	1		
Staphylococcus gallinarum		32.82	0.32	ESL 1	1	99		
Staphylococcus sciuri		1.79	0.12		99	99		
207750	**Low discrimination**							**207750**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		47.78	0.43	ESL 1	1	99	99	
Staphylococcus sciuri		32.65	0.40		99	99	1	
Staphylococcus piscifermentans		15.94	0.44		1	1	15	
207751	**Low discrimination**							**207751**
					OXI			
Staphylococcus sciuri		64.20	0.59	URE 1	99			
Staphylococcus gallinarum		24.54	0.45		1			
Staphylococcus equorum		3.86	0.31		1			
207752	**Low discrimination**							**207752**
					OXI	NOV	PYR	
Staphylococcus piscifermentans		50.33	0.48	ARG 99	1	1	15	
Staphylococcus saprophyticus subsp. bovis		25.13	0.31		1	99	99	
Staphylococcus sciuri		11.18	0.24		99	99	1	
207753	**Low discrimination**							**207753**
					OXI			
Staphylococcus gallinarum		60.19	0.53	XYL 99	1			
Staphylococcus lentus		13.21	0.47		99			
Staphylococcus sciuri		10.47	0.43		99			
207761	**Very good identification**							**207761**
					OXI			
Staphylococcus gallinarum		97.73	0.65	MAN 91	1			
Staphylococcus equorum		1.57	0.31		1			
Staphylococcus sciuri		0.61	0.25		99			
207762	**Good identification**							**207762**
					NOV			
Staphylococcus gallinarum		96.16	0.32	MAN 91	99			
Staphylococcus piscifermentans		2.22	0.14	MNS 99	1			
Staphylococcus equorum		1.62	-.02		99			
207763	**Very good identification**							**207763**
					COA	NOV		
Staphylococcus gallinarum		98.01	0.73	MAN 91	1	99		
Staphylococcus equorum		1.65	0.39		1	99		
Staphylococcus lutrae		0.15	0.12		99	1		
207770	**Low discrimination**							**207770**
					OXI			
Staphylococcus gallinarum		70.88	0.45	MNS 99	1			
Staphylococcus sciuri		17.29	0.38		99			
Staphylococcus equorum		11.15	0.31		1			
207771	**Low discrimination**							**207771**
					OXI			
Staphylococcus gallinarum		84.74	0.86		1			
Staphylococcus equorum		13.33	0.72		1			
Staphylococcus sciuri		1.69	0.56		99			
207772	**Low discrimination**							**207772**
					OXI			
Staphylococcus gallinarum		84.22	0.53	MNS 99	1			
Staphylococcus equorum		13.91	0.39		1			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus sciuri		1.37	0.22		99			
207773	**Low discrimination**							**207773**
					OXI			
Staphylococcus gallinarum		85.42	0.94		1			
Staphylococcus equorum		14.11	0.80		1			
Staphylococcus lentus		0.19	0.47		99			
210000	**Low discrimination**							**210000**
					OXI			
Dermacoccus nishinomiyaensis		62.48	0.59	ORN 1	93			
Micrococcus sp.		33.51	0.44		89			
Kocuria varians		3.60	0.41		13			
210002	**Low discrimination**							**210002**
					OXI			
Kocuria varians		51.17	0.30	ORN 1	13			
Dermacoccus nishinomiyaensis		31.78	0.18		93			
Micrococcus sp.		17.04	0.03		89			
210010	**Low discrimination**							**210010**
					YEP			
Macrococcus equipersicus		60.97	0.43	ORN 1	99			
Macrococcus bovicus		29.53	0.35		99			
Dermacoccus nishinomiyaensis		6.19	0.18		1			
210020	**Low discrimination**							**210020**
					OXI			
Kocuria varians		56.63	0.32	ORN 1	13			
Dermacoccus nishinomiyaensis		28.23	0.18		93			
Micrococcus sp.		15.14	0.03		89			
210050	**Low discrimination**							**210050**
Macrococcus equipersicus		70.53	0.52	ORN 1				
Macrococcus bovicus		29.47	0.43					
210100	**Low discrimination**							**210100**
					OXI			
Dermacoccus nishinomiyaensis		66.92	0.44	ORN 1	93			
Kocuria varians		30.39	0.45		13			
Micrococcus sp.		1.90	0.03		89			
210102	**Acceptable identification**							**210102**
					OXI			
Kocuria varians		92.69	0.33	ORN 1	13			
Dermacoccus nishinomiyaensis		7.31	0.03		93			
210110	**Low discrimination**							**210110**
					YEP			
Macrococcus equipersicus		64.53	0.36	ORN 1	99			
Macrococcus bovicus		31.25	0.28		99			
Dermacoccus nishinomiyaensis		2.91	0.03		1			
210120	**Acceptable identification**							**210120**
					OXI			
Kocuria varians		94.05	0.35	ORN 1	13			
Dermacoccus nishinomiyaensis		5.95	0.03		93			
210140	**Low discrimination**							**210140**
					OXI	YEP	NOV	
Kocuria varians		70.36	0.31	ORN 1	13	80	1	
Macrococcus equipersicus		22.51	0.18		99	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		7.13	0.03		93	1	80	
210150	**Low discrimination**							**210150**
Macrococcus equipersicus		70.53	0.45	ORN 1				
Macrococcus bovicus		29.47	0.35					
210200	**Low discrimination**							**210200**
					OXI			
Dermacoccus nishinomiyaensis		51.18	0.34	ORN 1	93			
Micrococcus sp.		42.52	0.23	SUC 6	89			
Kocuria varians		6.30	0.23		13			
210250	**Low discrimination**							**210250**
					OXI	NOV		
Macrococcus bovicus		50.49	0.31	ORN 1	99	99		
Macrococcus equipersicus		47.80	0.32		99	99		
Staphylococcus warneri		1.72	0.12		1	1		
210410	**Low discrimination**							**210410**
					OXI	YEP		
Macrococcus equipersicus		57.60	0.43	ORN 1	99	99		
Macrococcus bovicus		41.84	0.39		99	99		
Staphylococcus cohnii subsp. cohnii		0.56	0.07		1	12		
210450	**Low discrimination**							**210450**
					OXI	YEP		
Macrococcus equipersicus		60.75	0.52	ORN 1	99	99		
Macrococcus bovicus		38.08	0.47		99	99		
Staphylococcus cohnii subsp. cohnii		0.69	0.17		1	12		
210510	**Low discrimination**							**210510**
Macrococcus equipersicus		57.93	0.36	ORN 1				
Macrococcus bovicus		42.07	0.31					
210550	**Low discrimination**							**210550**
					OXI	PYR		
Macrococcus equipersicus		61.16	0.45	ORN 1	99	1		
Macrococcus bovicus		38.34	0.39		99	1		
Staphylococcus saprophyticus subsp. bovis		0.50	0.02		1	99		
210610	**Low discrimination**							**210610**
					OXI	NOV		
Macrococcus bovicus		62.60	0.27	ORN 1	99	99		
Macrococcus equipersicus		34.10	0.23		99	99		
Staphylococcus warneri		3.31	0.12		1	1		
210640	**Low discrimination**							**210640**
					PYR			
Staphylococcus warneri		81.24	0.41	ORN 1	8			
Staphylococcus hominis subsp. hominis		8.58	0.15		1			
Staphylococcus petrasii subsp. pragensis		5.64	-.09		99			
210642	**Low discrimination**							**210642**
					NOV			
Staphylococcus warneri		58.76	0.31	ORN 1	1			
Staphylococcus hominis subsp. hominis		21.87	0.16		1			
Staphylococcus saprophyticus subsp. saprophyticus		19.38	0.00		99			
210650	**Low discrimination**							**210650**
					OXI	NOV		
Macrococcus bovicus		42.36	0.34	ORN 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Micrococcus equipersicus		26.74	0.32		99	99	
Staphylococcus warneri		18.23	0.38		1	1	
210652	**Acceptable identification**						**210652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		91.62	0.29	ORN 1	99		
Staphylococcus warneri		8.38	0.28		1		
210740	**Low discrimination**						**210740**
Staphylococcus warneri		73.40	0.30	ORN 1			
Staphylococcus hominis subsp. hominis		26.60	0.16				
210750	**Low discrimination**						**210750**
					OXI	NOV	
Micrococcus bovicus		45.17	0.27	ORN 1	99	99	
Micrococcus equipersicus		28.51	0.24		99	99	
Staphylococcus warneri		14.32	0.28		1	1	
211010	**Good identification**						**211010**
Micrococcus equipersicus		97.53	0.50	ORN 1			
Micrococcus bovicus		2.47	0.16				
211040	**Good identification**						**211040**
Micrococcus equipersicus		100.0	0.33	ORN 1			
211050	**Very good identification**						**211050**
Micrococcus equipersicus		97.86	0.59	ORN 1			
Micrococcus bovicus		2.14	0.23				
211110	**Good identification**						**211110**
Micrococcus equipersicus		97.53	0.43	ORN 1			
Micrococcus bovicus		2.47	0.08				
211150	**Good identification**						**211150**
Micrococcus equipersicus		97.86	0.51	ORN 1			
Micrococcus bovicus		2.14	0.16				
211210	**Acceptable identification**						**211210**
Micrococcus equipersicus		93.99	0.29	ORN 1			
Micrococcus bovicus		6.02	0.03	SUC 9			
211250	**Acceptable identification**						**211250**
Micrococcus equipersicus		94.77	0.38	ORN 1			
Micrococcus bovicus		5.23	0.11	SUC 9			
211350	**Acceptable identification**						**211350**
Micrococcus equipersicus		94.77	0.31	ORN 1			
Micrococcus bovicus		5.23	0.03	SUC 9			
211410	**Acceptable identification**						**211410**
Micrococcus equipersicus		94.91	0.50	ORN 1			
Micrococcus bovicus		3.60	0.19				
Micrococcus carouselicus		1.49	0.08				
211440	**Good identification**						**211440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Micrococcus equipersicus		100.0	0.33	ORN 1		
211450	**Very good identification**					**211450**
Micrococcus equipersicus		96.83	0.59	ORN 1		
Micrococcus bovicus		3.17	0.27			
211510	**Acceptable identification**					**211510**
Micrococcus equipersicus		94.91	0.43	ORN 1		
Micrococcus bovicus		3.60	0.12			
Micrococcus carouselicus		1.49	0.00			
211550	**Good identification**					**211550**
Micrococcus equipersicus		96.83	0.51	ORN 1		
Micrococcus bovicus		3.17	0.19			
211610	**Low discrimination**					**211610**
Micrococcus equipersicus		70.55	0.29	ORN 1		
Micrococcus carouselicus		22.68	0.14	SUC 9		
Micrococcus bovicus		6.77	0.07			
211650	**Acceptable identification**					**211650**
Micrococcus equipersicus		92.35	0.38	ORN 1		
Micrococcus bovicus		7.65	0.15	SUC 9		
211750	**Acceptable identification**					**211750**
Micrococcus equipersicus		92.35	0.31	ORN 1		
Micrococcus bovicus		7.65	0.07	SUC 9		
212000	**Low discrimination**					**212000**
					OXI	
Dermacoccus nishinomiyaensis		63.64	0.54	ORN 1	93	
Kocuria varians		31.85	0.55		13	
Micrococcus sp.		3.19	0.18		89	
212001	**Low discrimination**					**212001**
					OXI	
Kocuria varians		81.16	0.34	ORN 1	13	
Dermacoccus nishinomiyaensis		18.84	0.13	MNS 8	93	
212002	**Acceptable identification**					**212002**
					OXI	
Kocuria varians		93.32	0.44	ORN 1	13	
Dermacoccus nishinomiyaensis		6.68	0.13		93	
212020	**Acceptable identification**					**212020**
					OXI	
Kocuria varians		90.27	0.46	ORN 1	13	
Dermacoccus nishinomiyaensis		5.19	0.13		93	
Kocuria rosea		4.55	0.10		35	
212022	**Good identification**					**212022**
Kocuria varians		100.0	0.35	ORN 1		
212040	**Low discrimination**					**212040**
					OXI	
Kocuria varians		87.73	0.42	ORN 1	13	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Dermacoccus nishinomiyaensis		8.07	0.13		93		
Micrococcus sp.		3.02	-.05		89		
212042	**Acceptable identification**						**212042**
Kocuria varians		94.67	0.30	ORN 1			
Staphylococcus hominis subsp. hominis		5.33	0.06				
212060	**Good identification**						**212060**
Kocuria varians		100.0	0.32	ORN 1			
212100	**Low discrimination**						**212100**
					OXI		
Kocuria varians		79.76	0.59	ORN 1	13		
Dermacoccus nishinomiyaensis		20.24	0.39		93		
212101	**Good identification**						**212101**
					OXI		
Kocuria varians		97.14	0.37	ORN 1	13		
Dermacoccus nishinomiyaensis		2.86	-.02	MNS 8	93		
212102	**Good identification**						**212102**
					OXI		
Kocuria varians		99.10	0.48	ORN 1	13		
Dermacoccus nishinomiyaensis		0.90	-.02		93		
212120	**Good identification**						**212120**
					OXI		
Kocuria varians		99.28	0.50	ORN 1	13		
Dermacoccus nishinomiyaensis		0.72	-.02		93		
212121	**Good identification**						**212121**
Kocuria varians		100.0	0.28	ORN 1			
212122	**Good identification**						**212122**
Kocuria varians		100.0	0.38	ORN 1			
212140	**Good identification**						**212140**
					OXI	NOV	
Kocuria varians		96.44	0.45	ORN 1	13	1	
Staphylococcus hominis subsp. novobiosepticus		1.49	-.02		1	99	
Dermacoccus nishinomiyaensis		1.13	-.02		93	80	
212142	**Acceptable identification**						**212142**
					NOV		
Kocuria varians		93.44	0.34	ORN 1	1		
Staphylococcus hominis subsp. hominis		3.80	0.07		1		
Staphylococcus hominis subsp. novobiosepticus		2.75	-.07		99		
212160	**Good identification**						**212160**
Kocuria varians		100.0	0.36	ORN 1			
212200	**Low discrimination**						**212200**
					OXI		
Kocuria varians		48.46	0.38	ORN 1	13		
Dermacoccus nishinomiyaensis		45.33	0.29	SUC 12	93		
Micrococcus sp.		3.52	-.03		89		
212220	**Good identification**						**212220**
Kocuria varians		100.0	0.28	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
212240	**Low discrimination**							**212240**
					NOV			
Staphylococcus hominis subsp. hominis		59.25	0.46	ORN 1	1			
Staphylococcus hominis subsp. novobiosepticus		33.76	0.30		99			
Kocuria varians		6.03	0.24		1			
212242	**Low discrimination**							**212242**
					NOV			
Staphylococcus hominis subsp. hominis		75.21	0.47	ORN 1	1			
Staphylococcus hominis subsp. novobiosepticus		19.66	0.24		99			
Staphylococcus epidermidis		3.30	0.16		1			
212250	**Low discrimination**							**212250**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		75.24	0.29	ORN 1	1	1	1	
Macrococcus bovicus		19.00	0.11	MAN 13	99	1	99	
Staphylococcus saprophyticus subsp. bovis		5.76	0.02		1	99	99	
212252	**Good identification**							**212252**
Staphylococcus hominis subsp. hominis		100.0	0.30	ORN 1				
212300	**Low discrimination**							**212300**
					OXI	NOV		
Kocuria varians		86.26	0.41	ORN 1	13	1		
Dermacoccus nishinomiyaensis		10.24	0.14	SUC 12	93	80		
Staphylococcus hominis subsp. novobiosepticus		2.14	-.02		1	99		
212302	**Low discrimination**							**212302**
					NOV			
Kocuria varians		89.85	0.30	ORN 1	1			
Staphylococcus hominis subsp. hominis		5.89	0.07	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		4.26	-.07		99			
212320	**Good identification**							**212320**
Kocuria varians		100.0	0.32	ORN 1				
212340	**Low discrimination**							**212340**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		56.28	0.39	VPT 90	99			
Staphylococcus hominis subsp. hominis		35.66	0.47	ORN 1	1			
Kocuria varians		5.02	0.28		1			
212341	**Low discrimination**							**212341**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		66.18	0.27	VPT 90	99			
Staphylococcus epidermidis		26.49	0.27	ORN 1	1			
Staphylococcus hominis subsp. hominis		3.42	0.12		1			
212342	**Low discrimination**							**212342**
					NOV			
Staphylococcus hominis subsp. hominis		50.32	0.48	ORN 1	1			
Staphylococcus hominis subsp. novobiosepticus		36.43	0.34		99			
Staphylococcus epidermidis		11.54	0.31		1			
212343	**Low discrimination**							**212343**
					NOV	PYR		
Staphylococcus epidermidis		59.84	0.37	VPT 96	1	1		
Staphylococcus hominis subsp. novobiosepticus		25.42	0.21	ORN 12	99	1		
Staphylococcus lugdunensis		11.88	0.18		1	96		
212350	**Low discrimination**							**212350**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR	NOV	
Staphylococcus hominis subsp. hominis		67.14	0.30	ORN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		18.99	0.14	MAN 13	99	99	
Staphylococcus hominis subsp. novobiosepticus		7.16	-.02		1	99	
212352	**Acceptable identification**						**212352**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		90.16	0.31	ORN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		5.43	0.02	MAN 13	99	99	
Staphylococcus hominis subsp. novobiosepticus		4.41	-.07		1	99	
212400	**Low discrimination**						**212400**
					OXI		
Kocuria varians		75.68	0.32	ORN 1	13		
Dermacoccus nishinomiyaensis		20.29	0.13	TRE 7	93		
Staphylococcus auricularis		4.03	0.05		1		
212450	**Low discrimination**						**212450**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		52.21	0.31	ORN 1	1	99	
Macrococcus bovicus		41.74	0.27		99	1	
Macrococcus equipersicus		6.05	0.11		99	1	
212500	**Good identification**						**212500**
					OXI		
Kocuria varians		96.71	0.36	ORN 1	13		
Dermacoccus nishinomiyaensis		3.29	-.02	TRE 7	93		
212520	**Good identification**						**212520**
Kocuria varians		100.0	0.26	ORN 1			
212550	**Acceptable identification**						**212550**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		91.07	0.43	ORN 1	1	99	
Macrococcus bovicus		7.80	0.19		99	1	
Macrococcus equipersicus		1.13	0.04		99	1	
212551	**Good identification**						**212551**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	ORN 1			
212552	**Good identification**						**212552**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	ORN 1			
212640	**Good identification**						**212640**
Staphylococcus hominis subsp. hominis		94.55	0.56	ORN 1			
Staphylococcus warneri		3.87	0.33				
Staphylococcus auricularis		0.88	0.12				
212641	**Acceptable identification**						**212641**
					OXI	PYR	
Staphylococcus lugdunensis		93.96	0.39	VPT 99	1	96	
Staphylococcus hominis subsp. hominis		5.13	0.21	LAC 90	1	1	
Kocuria kristinae		0.91	-.04		99	93	
212642	**Very good identification**						**212642**
					PYR		
Staphylococcus hominis subsp. hominis		96.10	0.58	ORN 1	1		
Staphylococcus lugdunensis		2.78	0.18		96		
Staphylococcus warneri		1.12	0.23		8		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
212643	**Very good identification**						**212643**
					PYR		
Staphylococcus lugdunensis		99.29	0.59	VPT 99	96		
Staphylococcus hominis subsp. hominis		0.71	0.23		1		
212650	**Low discrimination**						**212650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		64.49	0.43	ORN 1	99	99	
Staphylococcus hominis subsp. hominis		26.94	0.39		1	1	
Staphylococcus warneri		5.35	0.30		1	8	
212651	**Acceptable identification**						**212651**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		92.91	0.31	ORN 1	99	99	
Staphylococcus lugdunensis		3.92	-.02		1	96	
Staphylococcus hominis subsp. hominis		3.17	0.04		1	1	
212652	**Low discrimination**						**212652**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		63.85	0.41	ORN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		32.55	0.31	MAN 13	99	99	
Staphylococcus warneri		3.60	0.20		1	8	
212740	**Very good identification**						**212740**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		96.42	0.57	ORN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		1.29	0.14		99	99	
Staphylococcus warneri		1.15	0.23		1	8	
212741	**Acceptable identification**						**212741**
					NOV	PYR	
Staphylococcus lugdunensis		93.56	0.39	VPT 99	1	96	
Staphylococcus hominis subsp. hominis		5.54	0.22	LAC 90	1	1	
Staphylococcus saprophyticus subsp. bovis		0.91	0.02		99	99	
212742	**Very good identification**						**212742**
					PYR		
Staphylococcus hominis subsp. hominis		96.60	0.58	ORN 1	1		
Staphylococcus lugdunensis		2.58	0.18		96		
Staphylococcus warneri		0.33	0.13		8		
212743	**Very good identification**						**212743**
					PYR		
Staphylococcus lugdunensis		99.13	0.59	VPT 99	96		
Staphylococcus hominis subsp. hominis		0.77	0.23		1		
Staphylococcus chromogenes		0.10	-.02		45		
212750	**Low discrimination**						**212750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		88.89	0.55	ORN 1	99	99	
Staphylococcus hominis subsp. hominis		10.06	0.40		1	1	
Staphylococcus warneri		0.58	0.20		1	8	
212751	**Good identification**						**212751**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.06	0.43	ORN 1	99	99	
Staphylococcus lugdunensis		1.03	-.02		1	96	
Staphylococcus hominis subsp. hominis		0.91	0.05		1	1	
212752	**Low discrimination**						**212752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		64.94	0.43	ORN 1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		34.49	0.41		1	1	
Staphylococcus warneri		0.57	0.10		1	8	
212753	**Low discrimination**						**212753**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		68.30	0.31	ORN 1	99	99	
Staphylococcus lugdunensis		25.93	0.18		1	96	
Staphylococcus hominis subsp. hominis		2.96	0.06		1	1	
213771	**Low discrimination**						**213771**
					OXI		
Staphylococcus equorum		76.14	0.27	ORN 1	1		
Staphylococcus sciuri		8.50	0.11	GLR 90	99		
Staphylococcus gallinarum		7.34	0.04		1		
213773	**Low discrimination**						**213773**
					OXI		
Staphylococcus equorum		81.69	0.36	ORN 1	1		
Staphylococcus gallinarum		7.49	0.12	GLR 90	1		
Staphylococcus lentus		6.58	0.18		99		
214000	**Acceptable identification**						**214000**
Dermacoccus nishinomiyaensis		90.67	0.33	ORN 1			
Micrococcus sp.		9.33	0.03	PHS 5			
214450	**Low discrimination**						**214450**
					OXI		
Staphylococcus kloosii		81.80	0.46	ORN 1	1		
Macrococcus equipersicus		11.19	0.11		99		
Macrococcus bovicus		7.01	0.05		99		
214452	**Good identification**						**214452**
Staphylococcus kloosii		100.0	0.47	ORN 1			
214470	**Good identification**						**214470**
Staphylococcus kloosii		100.0	0.34	ORN 1			
214472	**Good identification**						**214472**
Staphylococcus kloosii		100.0	0.35	ORN 1			
214512	**Good identification**						**214512**
Staphylococcus kloosii		100.0	0.27	ORN 1			
214550	**Good identification**						**214550**
					OXI		
Staphylococcus kloosii		95.32	0.52	ORN 1	1		
Macrococcus equipersicus		2.88	0.04		99		
Macrococcus bovicus		1.80	-.02		99		
214552	**Very good identification**						**214552**
Staphylococcus kloosii		100.0	0.53	ORN 1			
214570	**Good identification**						**214570**
Staphylococcus kloosii		100.0	0.40	ORN 1			
214572	**Good identification**						**214572**
Staphylococcus kloosii		100.0	0.41	ORN 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
214750	**Low discrimination**					**214750**
Staphylococcus kloosii		80.11	0.32	ORN 1		
Staphylococcus saprophyticus subsp. bovis		19.89	0.02	SUC 9		
214752	**Good identification**					**214752**
Staphylococcus kloosii		100.0	0.32	ORN 1		
215450	**Low discrimination**					**215450**
					OXI	
Staphylococcus kloosii		77.48	0.46	ORN 1	1	
Macrococcus equipersicus		22.52	0.18		99	
215452	**Good identification**					**215452**
Staphylococcus kloosii		100.0	0.47	ORN 1		
215470	**Good identification**					**215470**
Staphylococcus kloosii		100.0	0.34	ORN 1		
215472	**Good identification**					**215472**
Staphylococcus kloosii		100.0	0.35	ORN 1		
215512	**Good identification**					**215512**
Staphylococcus kloosii		100.0	0.27	ORN 1		
215550	**Acceptable identification**					**215550**
					OXI	
Staphylococcus kloosii		93.97	0.52	ORN 1	1	
Macrococcus equipersicus		6.03	0.10		99	
215552	**Very good identification**					**215552**
Staphylococcus kloosii		100.0	0.53	ORN 1		
215570	**Good identification**					**215570**
Staphylococcus kloosii		100.0	0.40	ORN 1		
215572	**Good identification**					**215572**
Staphylococcus kloosii		100.0	0.41	ORN 1		
215750	**Good identification**					**215750**
Staphylococcus kloosii		100.0	0.32	ORN 1		
215752	**Good identification**					**215752**
Staphylococcus kloosii		100.0	0.32	ORN 1		
216000	**Acceptable identification**					**216000**
					OXI	
Dermacoccus nishinomiyaensis		91.24	0.28	ORN 1	93	
Kocuria varians		8.76	0.14	PHS 5	13	
216241	**Good identification**					**216241**
Staphylococcus epidermidis		100.0	0.27	VPT 96		
216242	**Low discrimination**					**216242**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		88.93	0.31	VPT 96			
Staphylococcus hominis subsp. hominis		11.07	0.16	ORN 12			
216243	**Good identification**						**216243**
					YEP		
Staphylococcus epidermidis		98.10	0.37	VPT 96	1		
Staphylococcus chromogenes		1.90	-.02	ORN 12	95		
216340	**Acceptable identification**						**216340**
					NOV		
Staphylococcus epidermidis		91.14	0.37	VPT 96	1		
Staphylococcus hominis subsp. hominis		5.85	0.15	ORN 12	1		
Staphylococcus hominis subsp. novobiosepticus		3.02	-.02		99		
216341	**Good identification**						**216341**
					YEP		
Staphylococcus epidermidis		98.93	0.42	VPT 96	1		
Staphylococcus chromogenes		1.07	-.02	ORN 12	95		
216342	**Good identification**						**216342**
					NOV		
Staphylococcus epidermidis		97.14	0.47	VPT 96	1		
Staphylococcus hominis subsp. hominis		2.31	0.17	ORN 12	1		
Staphylococcus hominis subsp. novobiosepticus		0.55	-.07		99		
216343	**Good identification**						**216343**
					YEP		
Staphylococcus epidermidis		97.01	0.52	VPT 96	1		
Staphylococcus chromogenes		2.99	0.17	ORN 12	95		
216550	**Good identification**						**216550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	ORN 1			
216650	**Good identification**						**216650**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		95.09	0.31	ORN 1	99	99	
Staphylococcus hominis subsp. hominis		4.91	0.08		1	1	
216703	**Low discrimination**						**216703**
					YEP		
Staphylococcus chromogenes		84.42	0.39	ARG 90	95		
Staphylococcus hyicus		10.70	0.17	ORN 1	1		
Staphylococcus felis		3.27	0.05		1		
216713	**Low discrimination**						**216713**
					COA	YEP	
Staphylococcus chromogenes		47.58	0.28	ARG 90	1	95	
Staphylococcus felis		43.77	0.22	ORN 1	1	1	
Staphylococcus intermedius		8.65	0.14		99	1	
216742	**Low discrimination**						**216742**
					YEP		
Staphylococcus hominis subsp. hominis		81.81	0.27	ORN 1	49		
Staphylococcus epidermidis		10.97	0.06	PHS 3	1		
Staphylococcus chromogenes		7.22	-.02		95		
216743	**Low discrimination**						**216743**
					PYR	YEP	
Staphylococcus chromogenes		86.52	0.39	ARG 90	45	95	
Staphylococcus lugdunensis		8.55	0.18	ORN 1	96	50	
Staphylococcus epidermidis		2.47	0.11		1	1	
216750	**Good identification**						**216750**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		98.62	0.43	ORN 1	99	99		
Staphylococcus hominis subsp. hominis		1.38	0.09		1	1		
216751	**Low discrimination**							**216751**
					COA	NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		85.58	0.31	ORN 1	1	99	80	
Staphylococcus chromogenes		10.30	0.08		1	1	95	
Staphylococcus intermedius		4.11	0.02		99	1	1	
216752	**Acceptable identification**							**216752**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		93.83	0.31	ORN 1	99	99		
Staphylococcus hominis subsp. hominis		6.17	0.10		1	1		
216753	**Low discrimination**							**216753**
					COA	NOV	YEP	
Staphylococcus chromogenes		58.92	0.28	ARG 90	1	1	95	
Staphylococcus intermedius		16.06	0.18	ORN 1	99	1	1	
Staphylococcus saprophyticus subsp. bovis		13.60	0.18		1	99	80	
217763	**Good identification**							**217763**
Staphylococcus gallinarum		98.34	0.32	ORN 1				
Staphylococcus equorum		1.66	-.02	MAN 91				
217771	**Low discrimination**							**217771**
					OXI			
Staphylococcus gallinarum		84.95	0.45	ORN 1	1			
Staphylococcus equorum		13.36	0.31		1			
Staphylococcus sciuri		1.69	0.15		99			
217773	**Low discrimination**							**217773**
					OXI			
Staphylococcus gallinarum		85.54	0.53	ORN 1	1			
Staphylococcus equorum		14.13	0.39		1			
Staphylococcus lentus		0.19	0.05		99			
220000	**Low discrimination**							**220000**
					OXI			
Kocuria varians		55.22	0.72		13			
Dermacoccus nishinomiyaensis		29.02	0.59		93			
Micrococcus sp.		15.56	0.44		89			
220001	**Low discrimination**							**220001**
					OXI			
Kocuria varians		88.72	0.51	MNS 8	13			
Micrococcus sp.		5.87	0.09		89			
Dermacoccus nishinomiyaensis		5.42	0.18		93			
220002	**Very good identification**							**220002**
					OXI			
Kocuria varians		97.19	0.61		13			
Dermacoccus nishinomiyaensis		1.83	0.18		93			
Micrococcus sp.		0.98	0.03		89			
220003	**Good identification**							**220003**
Kocuria varians		100.0	0.39	MNS 8				
220010	**Low discrimination**							**220010**
					NOV	OXI		
Macrococcus equipersicus		54.56	0.43	BGA 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Micrococcus bovicus		26.42	0.35		99	99	
Kocuria varians		10.53	0.31		1	13	
220020	**Very good identification**						**220020**
					OXI		
Kocuria varians		97.73	0.63		13		
Dermacoccus nishinomiyaensis		1.48	0.18		93		
Micrococcus sp.		0.79	0.03		89		
220021	**Good identification**						**220021**
Kocuria varians		100.0	0.41	MNS 8			
220022	**Good identification**						**220022**
Kocuria varians		100.0	0.52				
220023	**Good identification**						**220023**
Kocuria varians		100.0	0.30	MNS 8			
220040	**Low discrimination**						**220040**
					NOV	OXI	
Kocuria varians		86.09	0.59		1	13	
Micrococcus sp.		8.32	0.21		1	89	
Macrococcus equipersicus		2.92	0.26		99	99	
220041	**Good identification**						**220041**
					OXI		
Kocuria varians		97.78	0.37	MNS 8	13		
Micrococcus sp.		2.22	-.14		89		
220042	**Good identification**						**220042**
					NOV		
Kocuria varians		98.71	0.48		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.29	-.05		99		
220050	**Low discrimination**						**220050**
					OXI		
Macrococcus equipersicus		68.38	0.52	BGA 1	99		
Macrococcus bovicus		28.57	0.43		99		
Staphylococcus saprophyticus subsp. saprophyticus		1.66	0.08		1		
220060	**Good identification**						**220060**
Kocuria varians		100.0	0.50				
220061	**Good identification**						**220061**
Kocuria varians		100.0	0.28	MNS 8			
220062	**Good identification**						**220062**
Kocuria varians		100.0	0.38				
220100	**Good identification**						**220100**
					OXI		
Kocuria varians		93.51	0.76		13		
Dermacoccus nishinomiyaensis		6.24	0.44		93		
Micrococcus sp.		0.18	0.03		89		
220101	**Very good identification**						**220101**
					OXI		
Kocuria varians		99.23	0.54	MNS 8	13		
Dermacoccus nishinomiyaensis		0.77	0.03		93		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
220102	**Very good identification**							**220102**
					OXI			
Kocuria varians		99.76	0.65		13			
Dermacoccus nishinomiyaensis		0.24	0.03		93			
220103	**Good identification**							**220103**
Kocuria varians		100.0	0.43	MNS 8				
220110	**Low discrimination**							**220110**
					NOV	OXI		
Macrococcus equipersicus		45.36	0.36	BGA 1	99	99		
Kocuria varians		30.64	0.35		1	13		
Macrococcus bovicus		21.96	0.28		99	99		
220120	**Very good identification**							**220120**
					OXI			
Kocuria varians		99.81	0.67		13			
Dermacoccus nishinomiyaensis		0.19	0.03		93			
220121	**Good identification**							**220121**
Kocuria varians		100.0	0.45	MNS 8				
220122	**Very good identification**							**220122**
Kocuria varians		100.0	0.55					
220123	**Good identification**							**220123**
Kocuria varians		100.0	0.34	MNS 8				
220140	**Very good identification**							**220140**
					OXI	YEP	NOV	
Kocuria varians		98.74	0.62		13	80	1	
Macrococcus equipersicus		0.96	0.18		99	99	99	
Dermacoccus nishinomiyaensis		0.30	0.03		93	1	80	
220141	**Good identification**							**220141**
Kocuria varians		100.0	0.41	MNS 8				
220142	**Good identification**							**220142**
Kocuria varians		100.0	0.51					
220143	**Good identification**							**220143**
Kocuria varians		100.0	0.29	MNS 8				
220150	**Low discrimination**							**220150**
					NOV	OXI		
Macrococcus equipersicus		67.34	0.45	BGA 1	99	99		
Macrococcus bovicus		28.14	0.35		99	99		
Kocuria varians		3.69	0.21		1	13		
220160	**Very good identification**							**220160**
Kocuria varians		100.0	0.53					
220161	**Good identification**							**220161**
Kocuria varians		100.0	0.31	MNS 8				
220162	**Good identification**							**220162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		100.0	0.42				
220200	**Low discrimination**						**220200**
					OXI		
Kocuria varians		68.63	0.55	SUC 12	13		
Dermacoccus nishinomiyaensis		16.88	0.34		93		
Micrococcus sp.		14.03	0.23		89		
220201	**Low discrimination**						**220201**
					OXI	PYR	
Kocuria varians		86.56	0.33	SUC 12	13	10	
Kocuria kristinae		9.29	0.05	MNS 8	99	93	
Micrococcus sp.		4.15	-.11		89	63	
220202	**Good identification**						**220202**
Kocuria varians		100.0	0.43	SUC 12			
220220	**Good identification**						**220220**
Kocuria varians		100.0	0.45	SUC 12			
220222	**Good identification**						**220222**
Kocuria varians		100.0	0.34	SUC 12			
220240	**Low discrimination**						**220240**
					NOV		
Kocuria varians		54.85	0.41	SUC 12	1		
Staphylococcus warneri		31.45	0.43		1		
Staphylococcus hominis subsp. novobiosepticus		5.05	0.10		99		
220242	**Low discrimination**						**220242**
					NOV		
Kocuria varians		34.64	0.30	SUC 12	1		
Staphylococcus saprophyticus subsp. saprophyticus		33.67	0.16		99		
Staphylococcus warneri		23.48	0.33		1		
220250	**Low discrimination**						**220250**
					OXI		
Staphylococcus saprophyticus subsp. saprophyticus		45.31	0.29	TRE 99	1		
Macrococcus bovicus		19.27	0.31		99		
Macrococcus equipersicus		18.24	0.32		99		
220252	**Good identification**						**220252**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		97.70	0.44	TRE 99	99		
Staphylococcus warneri		2.06	0.30		1		
Staphylococcus simulans		0.25	0.02		1		
220260	**Good identification**						**220260**
Kocuria varians		100.0	0.32	SUC 12			
220300	**Very good identification**						**220300**
					OXI		
Kocuria varians		96.97	0.58	SUC 12	13		
Dermacoccus nishinomiyaensis		3.03	0.20		93		
220301	**Good identification**						**220301**
					OXI	PYR	
Kocuria varians		97.67	0.36	SUC 12	13	10	
Kocuria kristinae		2.33	-.05	MNS 8	99	93	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
220302	**Good identification**					**220302**
	Kocuria varians	100.0	0.47	SUC 12		
220320	**Good identification**					**220320**
	Kocuria varians	100.0	0.49	SUC 12		
220321	**Good identification**					**220321**
	Kocuria varians	100.0	0.27	SUC 12		
220322	**Good identification**					**220322**
	Kocuria varians	100.0	0.38	SUC 12		
220340	**Low discrimination**					**220340**
					NOV	
	Kocuria varians	76.02	0.45	SUC 12	1	
	Staphylococcus hominis subsp. novobiosepticus	14.00	0.20		99	
	Staphylococcus warneri	9.18	0.33		1	
220342	**Low discrimination**					**220342**
					NOV	
	Kocuria varians	60.55	0.33	SUC 12	1	
	Staphylococcus hominis subsp. novobiosepticus	21.29	0.14		99	
	Staphylococcus warneri	8.64	0.23		1	
220360	**Good identification**					**220360**
	Kocuria varians	100.0	0.35	SUC 12		
220373	**Good identification**					**220373**
	Staphylococcus xylosus	100.0	0.27	GLR 95		
220400	**Low discrimination**					**220400**
					OXI	
	Kocuria varians	84.64	0.49	TRE 7	13	
	Dermacoccus nishinomiyaensis	5.97	0.18		93	
	Micrococcus sp.	3.20	0.03		89	
220401	**Acceptable identification**					**220401**
					NOV	
	Kocuria varians	93.78	0.27	TRE 7	1	
	Staphylococcus cohnii subsp. cohnii	6.22	0.05	MNS 8	99	
220402	**Good identification**					**220402**
					NOV	
	Kocuria varians	96.32	0.38	TRE 7	1	
	Staphylococcus saprophyticus subsp. saprophyticus	3.68	-.05		99	
220410	**Low discrimination**					**220410**
					OXI	
	Macrococcus equipersicus	54.37	0.43	BGA 1	99	
	Macrococcus bovicus	39.49	0.39		99	
	Staphylococcus saprophyticus subsp. saprophyticus	3.57	0.08		1	
220413	**Low discrimination**					**220413**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	82.41	0.40	GLR 99	99	
	Staphylococcus simulans	13.01	0.18		1	
	Aerococcus viridans	4.58	0.25		1	
220420	**Good identification**					**220420**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.40	TRE 7				
220422	**Good identification**							**220422**
Kocuria varians		100.0	0.29	TRE 7				
220440	**Low discrimination**							**220440**
					NOV			
Staphylococcus warneri		28.63	0.43	SUC 95	1			
Kocuria varians		27.56	0.36		1			
Staphylococcus saprophyticus subsp. saprophyticus		23.65	0.21		99			
220442	**Low discrimination**							**220442**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		87.98	0.36	SUC 91	99			
Staphylococcus warneri		6.27	0.33	MAN 96	1			
Kocuria varians		5.10	0.24		1			
220443	**Low discrimination**							**220443**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		76.89	0.35	GLR 99	99	53		
Aerococcus viridans		10.16	0.28	MAN 91	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		5.95	-.05		99	7		
220450	**Low discrimination**							**220450**
					OXI			
Staphylococcus saprophyticus subsp. saprophyticus		55.47	0.49	SUC 91	1			
Macrococcus equipersicus		23.06	0.52		99			
Macrococcus bovicus		14.46	0.47		99			
220451	**Low discrimination**							**220451**
					PYR			
Staphylococcus cohnii subsp. urealyticum		45.45	0.35	GLR 99	53			
Staphylococcus saprophyticus subsp. bovis		20.02	0.22	LAC 91	99			
Staphylococcus saprophyticus subsp. saprophyticus		14.89	0.08		7			
220452	**Very good identification**							**220452**
					PYR			
Staphylococcus saprophyticus subsp. saprophyticus		98.69	0.65	SUC 91	7			
Staphylococcus kloosii		0.58	0.47		52			
Staphylococcus saprophyticus subsp. bovis		0.24	0.22		99			
220453	**Low discrimination**							**220453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		80.24	0.55	GLR 99	99	53		
Staphylococcus saprophyticus subsp. saprophyticus		14.73	0.23		99	7		
Aerococcus viridans		3.15	0.38		1	95		
220463	**Good identification**							**220463**
Aerococcus viridans		100.0	0.28	URE 1				
220473	**Low discrimination**							**220473**
					NOV			
Aerococcus viridans		61.93	0.38	URE 1	1			
Staphylococcus xylosus		22.12	0.08	ESL 90	99			
Staphylococcus cohnii subsp. urealyticum		15.95	0.14		99			
220500	**Very good identification**							**220500**
					OXI	YEP	NOV	
Kocuria varians		98.10	0.53	TRE 7	13	80	1	
Macrococcus equipersicus		1.03	0.09		99	99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		0.88	0.03		93	1	80	
220501	**Good identification**							**220501**
Kocuria varians		100.0	0.31	TRE 7				
220502	**Good identification**							**220502**
Kocuria varians		100.0	0.42	TRE 7				
220510	**Low discrimination**							**220510**
					NOV	OXI		
Macrococcus equipersicus		53.82	0.36	BGA 1	99	99		
Macrococcus bovicus		39.09	0.31		99	99		
Kocuria varians		2.74	0.12		1	13		
220513	**Low discrimination**							**220513**
					COA	NOV		
Staphylococcus cohnii subsp. urealyticum		83.38	0.32	GLR 99	1	99		
Aerococcus viridans		10.81	0.25		1	1		
Staphylococcus intermedius		5.81	0.01		99	1		
220520	**Good identification**							**220520**
Kocuria varians		100.0	0.44	TRE 7				
220522	**Good identification**							**220522**
Kocuria varians		100.0	0.32	TRE 7				
220540	**Low discrimination**							**220540**
					NOV	OXI		
Kocuria varians		69.30	0.39	TRE 7	1	13		
Staphylococcus warneri		15.16	0.33		1	1		
Macrococcus equipersicus		8.93	0.18		99	99		
220542	**Low discrimination**							**220542**
					NOV			
Kocuria varians		42.68	0.28	TRE 7	1			
Staphylococcus saprophyticus subsp. saprophyticus		42.66	0.14		99			
Staphylococcus warneri		11.03	0.23		1			
220543	**Low discrimination**							**220543**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		69.27	0.27	GLR 99	99	53		
Aerococcus viridans		21.36	0.28	MAN 91	1	95		
Kocuria varians		9.36	0.06		1	10		
220550	**Low discrimination**							**220550**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		32.67	0.47	NIT 99	1	99		
Macrococcus equipersicus		26.78	0.45		99	1		
Macrococcus bovicus		16.79	0.39		99	1		
220551	**Low discrimination**							**220551**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		74.24	0.34	NIT 99	1	99		
Staphylococcus cohnii subsp. urealyticum		18.06	0.27		1	53		
Macrococcus equipersicus		2.46	0.04		99	1		
220552	**Low discrimination**							**220552**
					PYR			
Staphylococcus saprophyticus subsp. saprophyticus		78.84	0.43	GAL 8	7			
Staphylococcus kloosii		10.33	0.53	SUC 91	52			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. bovis		8.69	0.34		99			
220553	**Low discrimination**							**220553**
					NOV			
Staphylococcus cohnii subsp. urealyticum		78.27	0.48	GLR 99	99			
Staphylococcus saprophyticus subsp. bovis		7.96	0.22		99			
Aerococcus viridans		7.16	0.38		1			
220560	**Good identification**							**220560**
Kocuria varians		100.0	0.30	TRE 7				
220570	**Low discrimination**							**220570**
					OXI	PYR		
Staphylococcus kloosii		74.41	0.40	PHS 99	1	52		
Staphylococcus saprophyticus subsp. bovis		10.96	0.05		1	99		
Macrococcus equipersicus		8.99	0.04		99	1		
220572	**Low discrimination**							**220572**
					NOV	YEP	PYR	
Staphylococcus kloosii		61.24	0.41	PHS 99	99	99	52	
Aerococcus viridans		19.88	0.30		1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		18.88	0.02		99	16	7	
220573	**Low discrimination**							**220573**
					COA	NOV	PYR	
Staphylococcus xylosus		55.60	0.29	GLR 95	1	99	65	
Staphylococcus lutrae		27.31	0.18	NIT 89	99	1	10	
Aerococcus viridans		15.40	0.38	SUC 95	1	1	95	
220600	**Low discrimination**							**220600**
					NOV			
Staphylococcus warneri		57.49	0.43	MLT 95	1			
Kocuria varians		34.38	0.31		1			
Staphylococcus saprophyticus subsp. saprophyticus		4.85	0.00		99			
220601	**Good identification**							**220601**
					OXI	PYR		
Kocuria kristinae		96.44	0.31	VPT 90	99	93		
Kocuria varians		3.56	0.10	URE 10	13	10		
220603	**Low discrimination**							**220603**
					OXI			
Staphylococcus simulans		74.59	0.34	ARG 92	1			
Kocuria kristinae		25.41	0.19	NIT 96	99			
220610	**Low discrimination**							**220610**
					OXI	NOV		
Staphylococcus saprophyticus subsp. saprophyticus		55.51	0.29	MLT 99	1	99		
Staphylococcus warneri		19.85	0.40		1	1		
Macrococcus bovicus		15.18	0.27		99	99		
220612	**Low discrimination**							**220612**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		88.43	0.44	MLT 99	99			
Staphylococcus simulans		9.52	0.35		1			
Staphylococcus warneri		1.86	0.30		1			
220613	**Acceptable identification**							**220613**
					NOV	PYR		
Staphylococcus simulans		94.45	0.46	ARG 92	1	70		
Staphylococcus saprophyticus subsp. saprophyticus		2.65	0.03	NIT 96	99	7		
Aerococcus viridans		1.39	0.25		1	95		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
220640	**Low discrimination**							**220640**
					NOV			
Staphylococcus warneri		67.94	0.69		1			
Staphylococcus saprophyticus subsp. saprophyticus		29.87	0.41		99			
Staphylococcus auricularis		0.48	0.17		1			
220641	**Good identification**							**220641**
					OXI	NOV	PYR	
Kocuria kristinae		96.41	0.51	VPT 90	99	1	93	
Staphylococcus warneri		2.95	0.35	URE 10	1	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		0.64	0.00		1	99	7	
220642	**Low discrimination**							**220642**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		87.65	0.57	MAN 96	99			
Staphylococcus warneri		11.73	0.60		1			
Staphylococcus simulans		0.26	0.16		1			
220643	**Low discrimination**							**220643**
					OXI	NOV		
Kocuria kristinae		73.33	0.39	VPT 90	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		11.07	0.16	URE 10	1	99		
Staphylococcus simulans		10.75	0.27		1	1		
220650	**Good identification**							**220650**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		90.83	0.70		99	7		
Staphylococcus warneri		6.23	0.67		1	8		
Staphylococcus saprophyticus subsp. bovis		1.95	0.47		99	99		
220651	**Low discrimination**							**220651**
					OXI	PYR	NOV	
Staphylococcus saprophyticus subsp. saprophyticus		50.77	0.29	MNS 1	1	7	99	
Staphylococcus saprophyticus subsp. bovis		27.01	0.34		1	99	99	
Kocuria kristinae		13.24	0.23		99	93	1	
220652	**Very good identification**							**220652**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		99.44	0.85		99	7		
Staphylococcus warneri		0.40	0.57		1	8		
Staphylococcus saprophyticus subsp. bovis		0.09	0.34		99	99		
220653	**Low discrimination**							**220653**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		79.40	0.44	MNS 1	99	7		
Staphylococcus simulans		12.85	0.39		1	70		
Staphylococcus saprophyticus subsp. bovis		1.86	0.22		99	99		
220660	**Low discrimination**							**220660**
					NOV			
Staphylococcus warneri		59.53	0.28	XYL 1	1			
Staphylococcus saprophyticus subsp. saprophyticus		26.17	0.00		99			
Kocuria varians		14.30	0.09		1			
220670	**Acceptable identification**							**220670**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		91.73	0.29	XYL 1	99	7		
Staphylococcus warneri		6.30	0.25		1	8		
Staphylococcus saprophyticus subsp. bovis		1.97	0.05		99	99		
220672	**Good identification**							**220672**
					NOV	PYR		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. saprophyticus		98.12	0.44	XYL 1	99	7		
Aerococcus viridans		0.89	0.30		1	95		
Staphylococcus arlettae		0.59	0.15		99	1		
220673	**Low discrimination**							**220673**
					NOV	PYR		
Staphylococcus xylosus		78.51	0.34	GLR 95	99	65		
Aerococcus viridans		11.57	0.38	NIT 89	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		5.53	0.03	GAL 91	99	7		
220700	**Low discrimination**							**220700**
Kocuria varians		73.96	0.35	SUC 12				
Staphylococcus warneri		26.04	0.33	TRE 7				
220703	**Low discrimination**							**220703**
					OXI	COA		
Staphylococcus intermedius		83.88	0.33	NIT 93	1	99		
Kocuria kristinae		8.25	0.09	PHS 97	99	1		
Staphylococcus simulans		6.32	0.12		1	1		
220713	**Low discrimination**							**220713**
					COA	NOV		
Staphylococcus intermedius		84.99	0.42	NIT 93	99	1		
Staphylococcus simulans		9.47	0.24	PHS 97	1	1		
Staphylococcus xylosus		2.87	0.10		1	99		
220733	**Acceptable identification**							**220733**
					COA	NOV		
Staphylococcus xylosus		93.07	0.32	GLR 95	1	99		
Aerococcus viridans		4.51	0.25	NIT 89	1	1		
Staphylococcus intermedius		2.42	0.01	MLT 93	99	1		
220740	**Low discrimination**							**220740**
					NOV			
Staphylococcus warneri		83.01	0.59		1			
Staphylococcus saprophyticus subsp. saprophyticus		10.05	0.19		99			
Kocuria varians		2.72	0.22		1			
220741	**Low discrimination**							**220741**
					OXI	COA	PYR	
Kocuria kristinae		89.61	0.41	VPT 90	99	1	93	
Staphylococcus intermedius		6.18	0.21	URE 10	1	99	90	
Staphylococcus warneri		2.60	0.24		1	1	8	
220742	**Low discrimination**							**220742**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		64.79	0.35	GAL 8	99			
Staphylococcus warneri		31.48	0.49	MAN 96	1			
Staphylococcus hominis subsp. hominis		1.62	0.17		1			
220743	**Low discrimination**							**220743**
					OXI	COA	NOV	
Staphylococcus intermedius		56.96	0.37	NIT 93	1	99	1	
Kocuria kristinae		33.62	0.29	PHS 97	99	1	1	
Staphylococcus xylosus		2.94	0.09		1	1	99	
220750	**Low discrimination**							**220750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		43.54	0.48	GAL 8	99	7		
Staphylococcus saprophyticus subsp. bovis		43.06	0.59		99	99		
Staphylococcus warneri		10.85	0.56		1	8		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
220751	**Low discrimination**							**220751**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		70.96	0.47	NIT 99	1	99		
Staphylococcus intermedius		12.97	0.30		99	1		
Staphylococcus xylosus		8.37	0.24		1	99		
220752	**Good identification**							**220752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		94.20	0.63	GAL 8	99	7		
Staphylococcus saprophyticus subsp. bovis		4.11	0.47		99	99		
Staphylococcus warneri		1.38	0.46		1	8		
220753	**Low discrimination**							**220753**
					COA	NOV		
Staphylococcus intermedius		54.25	0.46	NIT 93	99	1		
Staphylococcus xylosus		16.21	0.33	PHS 97	1	99		
Staphylococcus saprophyticus subsp. bovis		12.08	0.34		1	99		
220763	**Low discrimination**							**220763**
					COA	NOV	PYR	
Staphylococcus xylosus		77.84	0.30	GLR 95	1	99	65	
Staphylococcus lutrae		16.24	0.12	NIT 89	99	1	10	
Aerococcus viridans		5.92	0.28	MAN 94	1	1	95	
220771	**Good identification**							**220771**
					PYR			
Staphylococcus xylosus		98.06	0.46	GLR 95	65			
Staphylococcus equorum		0.76	0.04	NIT 89	5			
Staphylococcus saprophyticus subsp. bovis		0.73	0.05		99			
220773	**Good identification**							**220773**
					COA	NOV		
Staphylococcus xylosus		92.45	0.55	GLR 95	1	99		
Staphylococcus arlettae		2.89	0.34	NIT 89	1	99		
Staphylococcus lutrae		2.39	0.18		99	1		
221000	**Low discrimination**							**221000**
					OXI	YEP	NOV	
Kocuria varians		41.90	0.31	ESL 1	13	80	1	
Macrococcus equipersicus		24.28	0.24		99	99	99	
Dermacoccus nishinomiyaensis		22.02	0.18		93	1	80	
221010	**Good identification**							**221010**
Macrococcus equipersicus		97.53	0.50	BGA 1				
Macrococcus bovicus		2.47	0.16					
221040	**Low discrimination**							**221040**
					NOV	OXI		
Macrococcus equipersicus		87.71	0.33	BGA 1	99	99		
Kocuria varians		12.29	0.18	MAN 95	1	13		
221050	**Very good identification**							**221050**
Macrococcus equipersicus		97.86	0.59	BGA 1				
Macrococcus bovicus		2.14	0.23					
221053	**Low discrimination**							**221053**
					NOV			
Aerococcus viridans		82.54	0.31	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		17.46	0.06	TRE 95	99			
221073	**Good identification**							**221073**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		100.0	0.31	URE 1				
221100	**Low discrimination**							**221100**
					OXI	YEP	NOV	
Kocuria varians		81.15	0.35	ESL 1	13	80	1	
Macrococcus equipersicus		13.44	0.16		99	99	99	
Dermacoccus nishinomiyaensis		5.42	0.03		93	1	80	
221110	**Good identification**							**221110**
Macrococcus equipersicus		97.53	0.43	BGA 1				
Macrococcus bovicus		2.47	0.08					
221150	**Good identification**							**221150**
Macrococcus equipersicus		97.86	0.51	BGA 1				
Macrococcus bovicus		2.14	0.16					
221153	**Good identification**							**221153**
Aerococcus viridans		100.0	0.31	URE 1				
221173	**Good identification**							**221173**
Aerococcus viridans		100.0	0.31	URE 1				
221210	**Acceptable identification**							**221210**
Macrococcus equipersicus		93.99	0.29	BGA 1				
Macrococcus bovicus		6.02	0.03	SUC 9				
221250	**Acceptable identification**							**221250**
Macrococcus equipersicus		94.77	0.38	BGA 1				
Macrococcus bovicus		5.23	0.11	SUC 9				
221253	**Good identification**							**221253**
Aerococcus viridans		100.0	0.31	URE 1				
221273	**Good identification**							**221273**
Aerococcus viridans		100.0	0.31	URE 1				
221350	**Acceptable identification**							**221350**
Macrococcus equipersicus		94.77	0.31	BGA 1				
Macrococcus bovicus		5.23	0.03	SUC 9				
221353	**Good identification**							**221353**
Aerococcus viridans		100.0	0.31	URE 1				
221402	**Good identification**							**221402**
Aerococcus viridans		100.0	0.27	URE 1				
221403	**Low discrimination**							**221403**
					NOV			
Aerococcus viridans		81.25	0.35	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		18.75	0.11		99			
221410	**Acceptable identification**							**221410**
Macrococcus equipersicus		94.11	0.50	BGA 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		3.57	0.19					
Macrococcus carouselicus		1.47	0.08					
221412	**Low discrimination**							**221412**
					OXI	NOV	YEP	
Aerococcus viridans		70.00	0.37	URE 1	1	1	1	
Macrococcus equipersicus		19.60	0.09		99	99	99	
Staphylococcus kloosii		10.40	0.21		1	99	99	
221413	**Low discrimination**							**221413**
					NOV			
Aerococcus viridans		56.25	0.45	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		43.75	0.31		99			
221422	**Good identification**							**221422**
Aerococcus viridans		100.0	0.27	URE 1				
221423	**Good identification**							**221423**
Aerococcus viridans		100.0	0.35	URE 1				
221432	**Good identification**							**221432**
Aerococcus viridans		100.0	0.37	URE 1				
221433	**Good identification**							**221433**
Aerococcus viridans		100.0	0.45	URE 1				
221440	**Good identification**							**221440**
					OXI	NOV		
Macrococcus equipersicus		96.38	0.33	BGA 1	99	99		
Aerococcus viridans		3.62	0.20	MAN 95	1	1		
221441	**Low discrimination**							**221441**
					OXI	NOV		
Aerococcus viridans		59.27	0.28	URE 1	1	1		
Kocuria kristinae		23.48	-.03	LAC 90	99	1		
Staphylococcus cohnii subsp. urealyticum		17.25	0.06		1	99		
221442	**Low discrimination**							**221442**
					NOV	PYR		
Aerococcus viridans		86.83	0.40	URE 1	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		13.17	-.05		99	7		
221443	**Low discrimination**							**221443**
					NOV			
Aerococcus viridans		75.36	0.47	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		24.64	0.26		99			
221450	**Good identification**							**221450**
					OXI			
Macrococcus equipersicus		92.18	0.59	BGA 1	99			
Staphylococcus kloosii		3.20	0.46		1			
Macrococcus bovicus		3.02	0.27		99			
221451	**Low discrimination**							**221451**
					OXI	NOV		
Aerococcus viridans		36.89	0.38	URE 1	1	1		
Staphylococcus cohnii subsp. urealyticum		36.18	0.26	LAC 90	1	99		
Macrococcus equipersicus		26.93	0.18		99	99		
221453	**Low discrimination**							**221453**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		52.42	0.47	GLR 99	99			
Aerococcus viridans		47.58	0.57		1			
221461	**Good identification**							**221461**
Aerococcus viridans		100.0	0.28	URE 1				
221462	**Good identification**							**221462**
Aerococcus viridans		100.0	0.40	URE 1				
221463	**Good identification**							**221463**
Aerococcus viridans		100.0	0.47	URE 1				
221471	**Good identification**							**221471**
Aerococcus viridans		100.0	0.38	URE 1				
221472	**Low discrimination**							**221472**
					NOV	YEP		
Aerococcus viridans		85.01	0.50	URE 1	1	1		
Staphylococcus kloosii		14.99	0.35		99	99		
221473	**Very good identification**							**221473**
					NOV			
Aerococcus viridans		97.94	0.57	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		1.09	0.06		99			
Staphylococcus xylosus		0.97	-.04		99			
221502	**Good identification**							**221502**
Aerococcus viridans		100.0	0.27	URE 1				
221503	**Good identification**							**221503**
Aerococcus viridans		100.0	0.35	URE 1				
221510	**Acceptable identification**							**221510**
					OXI			
Macrococcus equipersicus		92.13	0.43	BGA 1	99			
Macrococcus bovicus		3.50	0.12		99			
Staphylococcus kloosii		2.06	0.26		1			
221512	**Low discrimination**							**221512**
					OXI	NOV	YEP	
Aerococcus viridans		71.01	0.37	URE 1	1	1	1	
Staphylococcus kloosii		20.47	0.27		1	99	99	
Macrococcus equipersicus		8.52	0.01		99	99	99	
221513	**Low discrimination**							**221513**
					NOV			
Aerococcus viridans		75.00	0.45	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		25.00	0.24		99			
221522	**Good identification**							**221522**
Aerococcus viridans		100.0	0.27	URE 1				
221523	**Good identification**							**221523**
Aerococcus viridans		100.0	0.35	URE 1				
221532	**Good identification**							**221532**
Aerococcus viridans		100.0	0.37	URE 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
221533	**Good identification**							**221533**
Aerococcus viridans		100.0	0.45	URE 1				
221541	**Good identification**							**221541**
Aerococcus viridans		100.0	0.28	URE 1				
221542	**Good identification**							**221542**
Aerococcus viridans		100.0	0.40	URE 1				
221543	**Low discrimination**							**221543**
					NOV			
Aerococcus viridans		87.71	0.47	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		12.29	0.19		99			
221550	**Low discrimination**							**221550**
					OXI			
Macrococcus equipersicus		82.66	0.51	BGA 1	99			
Staphylococcus kloosii		13.01	0.52		1			
Macrococcus bovicus		2.71	0.19		99			
221551	**Low discrimination**							**221551**
					OXI	NOV		
Aerococcus viridans		57.70	0.38	URE 1	1	1		
Staphylococcus cohnii subsp. urealyticum		24.25	0.19	LAC 90	1	99		
Macrococcus equipersicus		18.05	0.10		99	99		
221552	**Low discrimination**							**221552**
					NOV	YEP	PYR	
Staphylococcus kloosii		53.57	0.53	PHS 99	99	99	52	
Aerococcus viridans		39.13	0.50		1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		4.13	0.02		99	16	7	
221553	**Low discrimination**							**221553**
					NOV			
Aerococcus viridans		67.92	0.57	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		32.08	0.39		99			
221561	**Good identification**							**221561**
Aerococcus viridans		100.0	0.28	URE 1				
221562	**Good identification**							**221562**
Aerococcus viridans		100.0	0.40	URE 1				
221563	**Good identification**							**221563**
Aerococcus viridans		100.0	0.47	URE 1				
221570	**Low discrimination**							**221570**
					OXI	NOV	YEP	
Staphylococcus kloosii		62.18	0.40	PHS 99	1	99	99	
Aerococcus viridans		21.86	0.30		1	1	1	
Macrococcus equipersicus		15.96	0.10		99	99	99	
221571	**Low discrimination**							**221571**
					NOV	PYR		
Aerococcus viridans		66.49	0.38	URE 1	1	95		
Staphylococcus xylosus		21.09	0.07	LAC 90	99	65		
Staphylococcus equorum		12.42	0.04		99	5		
221572	**Low discrimination**							**221572**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Aerococcus viridans		74.50	0.50	URE 1	1	1	
Staphylococcus kloosii		25.50	0.41		99	99	
221573	**Low discrimination**						**221573**
					NOV	PYR	
Aerococcus viridans		86.68	0.57	URE 1	1	95	
Staphylococcus xylosus		8.70	0.16		99	65	
Staphylococcus equorum		4.63	0.12		99	5	
221602	**Good identification**						**221602**
Aerococcus viridans		100.0	0.27	URE 1			
221603	**Low discrimination**						**221603**
					OXI		
Aerococcus viridans		67.38	0.35	URE 1	1		
Kocuria kristinae		32.62	0.07		99		
221610	**Low discrimination**						**221610**
Macrococcus equipersicus		68.59	0.29	BGA 1			
Macrococcus carouselicus		22.05	0.14	SUC 9			
Macrococcus bovicus		6.58	0.07				
221612	**Low discrimination**						**221612**
					NOV	PYR	
Aerococcus viridans		66.85	0.37	URE 1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		33.15	0.03		99	7	
221613	**Acceptable identification**						**221613**
Aerococcus viridans		92.89	0.45	URE 1			
Staphylococcus simulans		7.11	0.05				
221622	**Good identification**						**221622**
Aerococcus viridans		100.0	0.27	URE 1			
221623	**Good identification**						**221623**
Aerococcus viridans		100.0	0.35	URE 1			
221632	**Good identification**						**221632**
Aerococcus viridans		100.0	0.37	URE 1			
221633	**Low discrimination**						**221633**
					NOV	PYR	
Aerococcus viridans		79.15	0.45	URE 1	1	95	
Staphylococcus equorum		16.36	0.12		99	5	
Staphylococcus xylosus		4.49	-.01		99	65	
221640	**Low discrimination**						**221640**
					OXI	NOV	
Staphylococcus warneri		44.21	0.28	ESL 1	1	1	
Macrococcus equipersicus		21.01	0.12		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		19.44	0.00		1	99	
221641	**Good identification**						**221641**
					OXI		
Kocuria kristinae		97.51	0.39	VPT 90	99		
Aerococcus viridans		2.49	0.28	URE 10	1		
221650	**Low discrimination**						**221650**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI			
Staphylococcus saprophyticus subsp. saprophyticus		47.67	0.29	ESL 1	1			
Macrococcus equipersicus		40.79	0.38		99			
Macrococcus bovicus		3.38	0.15		99			
221651	**Low discrimination**							**221651**
					OXI			
Aerococcus viridans		64.74	0.38	URE 1	1			
Kocuria kristinae		35.26	0.10	LAC 90	99			
221652	**Acceptable identification**							**221652**
					NOV	YEP	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		91.64	0.44	ESL 1	99	16	7	
Aerococcus viridans		7.47	0.50		1	1	95	
Staphylococcus kloosii		0.52	0.26		99	99	52	
221653	**Low discrimination**							**221653**
					NOV	PYR		
Aerococcus viridans		89.39	0.57	URE 1	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		4.75	0.03		99	7		
Staphylococcus cohnii subsp. urealyticum		3.05	0.16		99	53		
221661	**Low discrimination**							**221661**
					OXI			
Aerococcus viridans		71.63	0.28	URE 1	1			
Kocuria kristinae		28.37	-.03	LAC 90	99			
221662	**Good identification**							**221662**
Aerococcus viridans		100.0	0.40	URE 1				
221663	**Good identification**							**221663**
					NOV			
Aerococcus viridans		96.52	0.47	URE 1	1			
Staphylococcus xylosus		3.48	-.03		99			
221670	**Good identification**							**221670**
Aerococcus viridans		100.0	0.30	URE 1				
221672	**Low discrimination**							**221672**
					NOV	PYR		
Aerococcus viridans		83.47	0.50	URE 1	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		10.35	0.03		99	7		
Staphylococcus arlettae		6.19	0.15		99	1		
221673	**Low discrimination**							**221673**
					NOV	PYR		
Aerococcus viridans		57.04	0.57	URE 1	1	95		
Staphylococcus equorum		29.81	0.32		99	5		
Staphylococcus xylosus		10.75	0.22		99	65		
221702	**Good identification**							**221702**
Aerococcus viridans		100.0	0.27	URE 1				
221703	**Low discrimination**							**221703**
					OXI			
Aerococcus viridans		86.10	0.35	URE 1	1			
Kocuria kristinae		13.90	-.03		99			
221712	**Good identification**							**221712**
Aerococcus viridans		100.0	0.37	URE 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
221713	**Low discrimination**							**221713**
					OXI	COA	NOV	
Aerococcus viridans		81.39	0.45	URE 1	1	1	1	
Staphylococcus lentus		9.69	0.12		99	1	99	
Staphylococcus intermedius		4.86	0.01		1	99	1	
221722	**Good identification**							**221722**
Aerococcus viridans		100.0	0.27	URE 1				
221723	**Good identification**							**221723**
Aerococcus viridans		100.0	0.35	URE 1				
221732	**Good identification**							**221732**
Aerococcus viridans		100.0	0.37	URE 1				
221733	**Low discrimination**							**221733**
					NOV	PYR		
Staphylococcus equorum		55.25	0.32	GLR 90	99	5		
Aerococcus viridans		26.43	0.45	NIT 92	1	95		
Staphylococcus xylosus		15.17	0.20	MLT 91	99	65		
221741	**Acceptable identification**							**221741**
					OXI			
Kocuria kristinae		92.89	0.29	VPT 90	99			
Aerococcus viridans		7.11	0.28	URE 10	1			
221742	**Low discrimination**							**221742**
					NOV	PYR		
Aerococcus viridans		88.23	0.40	URE 1	1	95		
Staphylococcus saprophyticus subsp. saprophyticus		11.77	-.06		99	7		
221743	**Low discrimination**							**221743**
					OXI	NOV		
Aerococcus viridans		71.69	0.47	URE 1	1	1		
Kocuria kristinae		26.03	0.16		99	1		
Staphylococcus xylosus		2.28	-.04		1	99		
221750	**Low discrimination**							**221750**
					OXI	PYR		
Macrococcus equipersicus		52.33	0.31	BGA 1	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		12.41	0.07	SUC 9	1	7		
Staphylococcus saprophyticus subsp. bovis		12.27	0.18		1	99		
221751	**Low discrimination**							**221751**
					NOV	PYR		
Aerococcus viridans		44.94	0.38	URE 1	1	95		
Staphylococcus xylosus		23.55	0.11	LAC 90	99	65		
Staphylococcus equorum		8.39	0.04		99	5		
221753	**Low discrimination**							**221753**
					OXI	NOV		
Aerococcus viridans		75.96	0.57	URE 1	1	1		
Staphylococcus xylosus		12.59	0.21		1	99		
Staphylococcus lentus		4.59	0.18		99	99		
221762	**Good identification**							**221762**
Aerococcus viridans		100.0	0.40	URE 1				
221763	**Low discrimination**							**221763**
					NOV	PYR		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Aerococcus viridans		65.57	0.47	URE 1	1	95		
Staphylococcus xylosus		23.93	0.18		99	65		
Staphylococcus equorum		10.50	0.12		99	5		
221770	**Low discrimination**							**221770**
					NOV	YEP	PYR	
Aerococcus viridans		58.24	0.30	URE 1	1	1	95	
Staphylococcus equorum		25.38	0.04	LAC 90	99	1	5	
Staphylococcus kloosii		16.38	0.19		99	99	52	
221771	**Low discrimination**							**221771**
					NOV	PYR		
Staphylococcus equorum		71.50	0.45	GLR 90	99	5		
Staphylococcus xylosus		23.30	0.33	NIT 92	99	65		
Aerococcus viridans		3.87	0.38		1	95		
221772	**Low discrimination**							**221772**
					NOV	YEP	PYR	
Aerococcus viridans		61.61	0.50	URE 1	1	1	95	
Staphylococcus arlettae		25.87	0.31		99	99	1	
Staphylococcus equorum		7.67	0.12		99	1	5	
221773	**Low discrimination**							**221773**
					NOV	PYR		
Staphylococcus equorum		61.87	0.53	GLR 90	99	5		
Staphylococcus xylosus		22.32	0.43	NIT 92	99	65		
Aerococcus viridans		11.71	0.57		1	95		
222000	**Good identification**							**222000**
					OXI			
Kocuria varians		93.90	0.87		13			
Dermacoccus nishinomiyaensis		5.69	0.54		93			
Micrococcus sp.		0.29	0.18		89			
222001	**Very good identification**							**222001**
					OXI			
Kocuria varians		99.30	0.65	MNS 8	13			
Dermacoccus nishinomiyaensis		0.70	0.13		93			
222002	**Very good identification**							**222002**
					OXI			
Kocuria varians		99.78	0.75		13			
Dermacoccus nishinomiyaensis		0.22	0.13		93			
222003	**Very good identification**							**222003**
Kocuria varians		99.23	0.53	MNS 8				
Staphylococcus simulans		0.77	0.07					
222010	**Low discrimination**							**222010**
					OXI	YEP	NOV	
Kocuria varians		87.56	0.46	MAN 1	13	80	1	
Dermacoccus nishinomiyaensis		5.30	0.13		93	1	80	
Macrococcus bovicus		5.00	0.16		99	99	99	
222012	**Acceptable identification**							**222012**
Kocuria varians		92.58	0.34	MAN 1				
Staphylococcus simulans		7.43	0.09					
222020	**Very good identification**							**222020**
					OXI			
Kocuria varians		99.67	0.77		13			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		0.17	0.13		93			
Kocuria rosea		0.15	0.10		35			
222021	**Very good identification**							**222021**
Kocuria varians		100.0	0.55	MNS 8				
222022	**Very good identification**							**222022**
Kocuria varians		100.0	0.66					
222023	**Good identification**							**222023**
Kocuria varians		100.0	0.44	MNS 8				
222030	**Good identification**							**222030**
Kocuria varians		100.0	0.36	MAN 1				
222040	**Very good identification**							**222040**
					OXI			
Kocuria varians		99.36	0.73		13			
Dermacoccus nishinomiyaensis		0.28	0.13		93			
Staphylococcus auricularis		0.18	0.15		1			
222041	**Good identification**							**222041**
Kocuria varians		100.0	0.51	MNS 8				
222042	**Very good identification**							**222042**
Kocuria varians		99.83	0.62					
Staphylococcus hominis subsp. hominis		0.17	0.06					
222043	**Good identification**							**222043**
Kocuria varians		98.43	0.40	MNS 8				
Staphylococcus simulans		1.57	0.00					
222050	**Low discrimination**							**222050**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		49.55	0.34	TRE 99	99	99	1	
Kocuria varians		29.01	0.32		1	10	13	
Macrococcus bovicus		17.61	0.23		99	1	99	
222060	**Very good identification**							**222060**
Kocuria varians		100.0	0.64					
222061	**Good identification**							**222061**
Kocuria varians		100.0	0.42	MNS 8				
222062	**Good identification**							**222062**
Kocuria varians		100.0	0.52					
222063	**Good identification**							**222063**
Kocuria varians		100.0	0.31	MNS 8				
222100	**Very good identification**							**222100**
					OXI			
Kocuria varians		99.24	0.90		13			
Dermacoccus nishinomiyaensis		0.76	0.39		93			

Code		ID	T	tests	Differentiation tests			Code
Taxon		score	index	against				
222101	**Very good identification**							**222101**
					OXI			
Kocuria varians		99.91	0.68	MNS 8	13			
Dermacoccus nishinomiyaensis		0.09	-.02		93			
222102	**Very good identification**							**222102**
					OXI			
Kocuria varians		99.97	0.79		13			
Dermacoccus nishinomiyaensis		0.03	-.02		93			
222103	**Very good identification**							**222103**
Kocuria varians		100.0	0.57	MNS 8				
222110	**Good identification**							**222110**
					NOV	PYR	OXI	
Kocuria varians		96.70	0.49	MAN 1	1	10	13	
Macrococcus bovicus		1.58	0.08		99	1	99	
Staphylococcus saprophyticus subsp. bovis		0.98	0.05		99	99	1	
222111	**Good identification**							**222111**
Kocuria varians		100.0	0.27	MAN 1				
222112	**Good identification**							**222112**
Kocuria varians		100.0	0.38	MAN 1				
222120	**Very good identification**							**222120**
					OXI			
Kocuria varians		99.98	0.81		13			
Dermacoccus nishinomiyaensis		0.02	-.02		93			
222121	**Very good identification**							**222121**
Kocuria varians		100.0	0.59	MNS 8				
222122	**Very good identification**							**222122**
Kocuria varians		100.0	0.70					
222123	**Good identification**							**222123**
Kocuria varians		100.0	0.48	MNS 8				
222130	**Good identification**							**222130**
Kocuria varians		100.0	0.40	MAN 1				
222132	**Good identification**							**222132**
Kocuria varians		100.0	0.28	MAN 1				
222140	**Very good identification**							**222140**
					NOV	PYR		
Kocuria varians		99.84	0.77		1	10		
Staphylococcus hominis subsp. novobiosepticus		0.05	-.02		99	1		
Staphylococcus saprophyticus subsp. bovis		0.05	0.05		99	99		
222141	**Very good identification**							**222141**
Kocuria varians		100.0	0.55	MNS 8				
222142	**Very good identification**							**222142**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		99.79	0.65		1			
Staphylococcus hominis subsp. hominis		0.12	0.07		1			
Staphylococcus hominis subsp. novobiosepticus		0.09	-.07		99			
222143	**Good identification**							**222143**
Kocuria varians		98.91	0.44	MNS 8				
Staphylococcus epidermidis		1.10	0.01					
222150	**Low discrimination**							**222150**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		79.00	0.47	TRE 99	99	99	1	
Kocuria varians		17.34	0.36		1	10	13	
Macrococcus bovicus		3.01	0.16		99	1	99	
222151	**Acceptable identification**							**222151**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		92.91	0.34	TRE 99	99	99		
Kocuria varians		7.09	0.14		1	10		
222152	**Low discrimination**							**222152**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		80.15	0.34	TRE 99	99	99		
Kocuria varians		19.85	0.24		1	10		
222160	**Very good identification**							**222160**
Kocuria varians		100.0	0.67					
222161	**Good identification**							**222161**
Kocuria varians		100.0	0.45	MNS 8				
222162	**Very good identification**							**222162**
Kocuria varians		100.0	0.56					
222163	**Low discrimination**							**222163**
					COA			
Kocuria varians		78.47	0.34	MNS 8	1			
Staphylococcus lutrae		21.53	0.12		99			
222200	**Very good identification**							**222200**
					OXI			
Kocuria varians		96.72	0.69	SUC 12	13			
Dermacoccus nishinomiyaensis		2.74	0.29		93			
Micrococcus sp.		0.21	-.03		89			
222201	**Good identification**							**222201**
Kocuria varians		100.0	0.47	SUC 12				
222202	**Very good identification**							**222202**
Kocuria varians		96.30	0.57	SUC 12				
Staphylococcus simulans		3.43	0.25					
Staphylococcus hominis subsp. hominis		0.27	0.06					
222203	**Low discrimination**							**222203**
Staphylococcus simulans		57.85	0.36	ARG 92				
Kocuria varians		42.15	0.36	TRE 95				
222210	**Low discrimination**							**222210**
					NOV	PYR	OXI	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		73.54	0.28	SUC 12	1	10	13	
Staphylococcus vitulinus		10.59	0.00	MAN 1	99	1	99	
Staphylococcus saprophyticus subsp. bovis		8.17	0.05		99	99	1	
222212	**Acceptable identification**							**222212**
Staphylococcus simulans		93.38	0.37	ARG 92				
Kocuria varians		6.62	0.16	TRE 95				
222213	**Good identification**							**222213**
					COA	PYR		
Staphylococcus simulans		97.89	0.48	ARG 92	1	70		
Staphylococcus capitis subsp. ureolyticus		1.50	-.02	TRE 95	1	1		
Staphylococcus intermedius		0.61	0.01		99	90		
222220	**Very good identification**							**222220**
Kocuria varians		100.0	0.59	SUC 12				
222221	**Good identification**							**222221**
Kocuria varians		100.0	0.38	SUC 12				
222222	**Good identification**							**222222**
Kocuria varians		100.0	0.48	SUC 12				
222240	**Low discrimination**							**222240**
					NOV			
Kocuria varians		64.76	0.55	SUC 12	1			
Staphylococcus hominis subsp. hominis		19.29	0.46		1			
Staphylococcus hominis subsp. novobiosepticus		10.99	0.30		99			
222241	**Low discrimination**							**222241**
					NOV			
Kocuria varians		55.09	0.33	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		26.88	0.17	MNS 8	99			
Staphylococcus epidermidis		10.44	0.16		1			
222242	**Low discrimination**							**222242**
					NOV			
Staphylococcus hominis subsp. hominis		44.34	0.47	BGA 1	1			
Kocuria varians		35.77	0.44		1			
Staphylococcus hominis subsp. novobiosepticus		11.59	0.24		99			
222243	**Low discrimination**							**222243**
Staphylococcus simulans		37.23	0.29	ARG 92				
Staphylococcus epidermidis		28.21	0.27	TRE 95				
Kocuria varians		13.26	0.22					
222250	**Low discrimination**							**222250**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		83.19	0.47	TRE 99	99	99		
Staphylococcus hominis subsp. hominis		7.32	0.29		1	1		
Staphylococcus warneri		5.99	0.33		1	8		
222251	**Low discrimination**							**222251**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		79.00	0.34	TRE 99	99	99		
Staphylococcus capitis subsp. ureolyticus		19.26	0.09		1	1		
Staphylococcus simulans		1.74	0.00		1	70		
222252	**Low discrimination**							**222252**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		43.68	0.34	TRE 99	99	99	
Staphylococcus simulans		28.41	0.30		1	70	
Staphylococcus hominis subsp. hominis		18.04	0.30		1	1	
222253	**Low discrimination**						**222253**
					NOV	PYR	
Staphylococcus simulans		50.58	0.41	ARG 92	1	70	
Staphylococcus capitis subsp. ureolyticus		41.52	0.26	TRE 95	1	1	
Staphylococcus saprophyticus subsp. bovis		5.81	0.22		99	99	
222260	**Good identification**						**222260**
Kocuria varians		99.15	0.46	SUC 12			
Staphylococcus hominis subsp. hominis		0.85	0.05				
222262	**Good identification**						**222262**
Kocuria varians		96.56	0.35	SUC 12			
Staphylococcus hominis subsp. hominis		3.44	0.06				
222300	**Very good identification**						**222300**
					OXI	NOV	
Kocuria varians		99.52	0.72	SUC 12	13	1	
Dermacoccus nishinomiyaensis		0.36	0.14		93	80	
Staphylococcus hominis subsp. novobiosepticus		0.08	-.02		1	99	
222301	**Good identification**						**222301**
					COA	PYR	
Kocuria varians		96.30	0.51	SUC 12	1	10	
Staphylococcus intermedius		3.70	0.17	MNS 8	99	90	
222302	**Very good identification**						**222302**
Kocuria varians		99.45	0.61	SUC 12			
Staphylococcus simulans		0.21	0.03				
Staphylococcus hominis subsp. hominis		0.20	0.07				
222303	**Low discrimination**						**222303**
					COA	PYR	
Kocuria varians		49.36	0.39	SUC 12	1	10	
Staphylococcus intermedius		41.27	0.33	MNS 8	99	90	
Staphylococcus simulans		3.93	0.14		1	70	
222310	**Low discrimination**						**222310**
					NOV	PYR	
Kocuria varians		77.14	0.31	SUC 12	1	10	
Staphylococcus saprophyticus subsp. bovis		22.86	0.18	MAN 1	99	99	
222313	**Low discrimination**						**222313**
					COA		
Staphylococcus intermedius		85.19	0.42	PHS 97	99		
Staphylococcus simulans		11.99	0.26	TRE 93	1		
Staphylococcus felis		1.08	-.04		1		
222320	**Very good identification**						**222320**
Kocuria varians		100.0	0.63	SUC 12			
222321	**Good identification**						**222321**
Kocuria varians		100.0	0.41	SUC 12			
222322	**Good identification**						**222322**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.52	SUC 12				
222323	**Acceptable identification**							**222323**
					NOV			
Kocuria varians		94.90	0.30	SUC 12	1			
Staphylococcus xylosus		5.10	-.03	MNS 8	99			
222340	**Low discrimination**							**222340**
					NOV			
Kocuria varians		62.02	0.59	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		21.06	0.39		99			
Staphylococcus hominis subsp. hominis		13.34	0.47		1			
222341	**Low discrimination**							**222341**
					NOV			
Kocuria varians		32.85	0.37	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		32.06	0.27	MNS 8	99			
Staphylococcus epidermidis		23.53	0.32		1			
222342	**Low discrimination**							**222342**
					NOV			
Kocuria varians		33.96	0.48	SUC 12	1			
Staphylococcus hominis subsp. hominis		30.41	0.48		1			
Staphylococcus hominis subsp. novobiosepticus		22.01	0.34		99			
222343	**Low discrimination**							**222343**
					COA	NOV	PYR	
Staphylococcus epidermidis		46.16	0.42	VPT 96	1	1	1	
Staphylococcus intermedius		32.80	0.37		99	1	90	
Staphylococcus hominis subsp. novobiosepticus		10.70	0.21		1	99	1	
222350	**Very good identification**							**222350**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		95.98	0.59	TRE 99	99	99		
Staphylococcus hominis subsp. hominis		2.29	0.30		1	1		
Kocuria varians		0.72	0.18		1	10		
222351	**Low discrimination**							**222351**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		81.80	0.47	TRE 99	1	99		
Staphylococcus intermedius		14.95	0.30		99	1		
Staphylococcus xylosus		3.25	0.14		1	99		
222352	**Low discrimination**							**222352**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		86.30	0.47	TRE 99	99	99		
Staphylococcus hominis subsp. hominis		9.65	0.31		1	1		
Staphylococcus simulans		1.22	0.08		1	70		
222353	**Low discrimination**							**222353**
					COA	NOV		
Staphylococcus intermedius		72.63	0.46	PHS 97	99	1		
Staphylococcus saprophyticus subsp. bovis		16.17	0.34	TRE 93	1	99		
Staphylococcus xylosus		7.32	0.23		1	99		
222360	**Good identification**							**222360**
					NOV			
Kocuria varians		98.43	0.50	SUC 12	1			
Staphylococcus hominis subsp. novobiosepticus		0.96	-.02		99			
Staphylococcus hominis subsp. hominis		0.61	0.05		1			
222361	**Low discrimination**							**222361**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV			
Kocuria varians		75.63	0.28	SUC 12	1			
Staphylococcus xylosus		24.37	0.11	MNS 8	99			
222362	**Good identification**							**222362**
					NOV			
Kocuria varians		95.75	0.38	SUC 12	1			
Staphylococcus hominis subsp. hominis		2.47	0.07		1			
Staphylococcus hominis subsp. novobiosepticus		1.78	-.07		99			
222371	**Good identification**							**222371**
Staphylococcus xylosus		97.84	0.36	GLR 95				
Staphylococcus saprophyticus subsp. bovis		2.16	0.05	TRE 96				
222373	**Acceptable identification**							**222373**
					COA	PYR	NOV	
Staphylococcus xylosus		92.13	0.45	GLR 95	1	65	99	
Staphylococcus lutrae		7.06	0.18	TRE 96	99	10	1	
Staphylococcus intermedius		0.80	0.05		99	90	1	
222400	**Very good identification**							**222400**
					OXI			
Kocuria varians		96.60	0.63	TRE 7	13			
Staphylococcus auricularis		2.51	0.30		1			
Dermacoccus nishinomiyaensis		0.79	0.13		93			
222401	**Good identification**							**222401**
Kocuria varians		96.79	0.42	TRE 7				
Staphylococcus auricularis		3.21	0.10	MNS 8				
222402	**Acceptable identification**							**222402**
Kocuria varians		94.22	0.52	TRE 7				
Staphylococcus simulans		4.82	0.23					
Staphylococcus auricularis		0.96	0.10					
222403	**Low discrimination**							**222403**
Staphylococcus simulans		62.72	0.34	ARG 92				
Kocuria varians		31.87	0.30	SUC 96				
Staphylococcus felis		5.41	0.03					
222410	**Low discrimination**							**222410**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		68.60	0.34	MLT 99	99	99	1	
Macrococcus bovicus		15.67	0.19		99	1	99	
Kocuria varians		13.77	0.22		1	10	13	
222412	**Low discrimination**							**222412**
					NOV	PYR		
Staphylococcus simulans		78.90	0.35	ARG 92	1	70		
Staphylococcus saprophyticus subsp. bovis		17.21	0.22	SUC 96	99	99		
Kocuria varians		3.90	0.11		1	10		
222413	**Low discrimination**							**222413**
					NOV			
Staphylococcus simulans		82.83	0.46	ARG 92	1			
Staphylococcus felis		11.96	0.20	SUC 96	1			
Staphylococcus cohnii subsp. urealyticum		3.86	0.24		99			
222420	**Very good identification**							**222420**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.54	TRE 7				
222421	**Good identification**							**222421**
Kocuria varians		100.0	0.32	TRE 7				
222422	**Good identification**							**222422**
Kocuria varians		100.0	0.43	TRE 7				
222440	**Low discrimination**							**222440**
					NOV	PYR		
Kocuria varians		61.97	0.50	TRE 7	1	10		
Staphylococcus auricularis		17.11	0.37		1	40		
Staphylococcus saprophyticus subsp. bovis		14.21	0.34		99	99		
222441	**Low discrimination**							**222441**
					NOV	PYR		
Kocuria varians		49.71	0.28	TRE 7	1	10		
Staphylococcus saprophyticus subsp. bovis		32.76	0.22	MNS 8	99	99		
Staphylococcus auricularis		17.54	0.18		1	40		
222442	**Low discrimination**							**222442**
					NOV	PYR		
Kocuria varians		61.28	0.39	TRE 7	1	10		
Staphylococcus saprophyticus subsp. bovis		12.45	0.22		99	99		
Staphylococcus auricularis		6.67	0.18		1	40		
222443	**Low discrimination**							**222443**
					NOV			
Staphylococcus simulans		61.52	0.27	ARG 92	1			
Kocuria varians		15.27	0.17	SUC 96	1			
Staphylococcus cohnii subsp. urealyticum		14.29	0.19		99			
222450	**Very good identification**							**222450**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.87	0.75		1	99	99	
Macrococcus bovicus		0.53	0.27		99	99	1	
Staphylococcus warneri		0.29	0.33		1	1	8	
222451	**Very good identification**							**222451**
Staphylococcus saprophyticus subsp. bovis		99.52	0.63					
Staphylococcus cohnii subsp. urealyticum		0.40	0.19					
Staphylococcus xylosus		0.08	-.05					
222452	**Good identification**							**222452**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		93.70	0.63		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		3.99	0.23		99	7		
Staphylococcus simulans		1.95	0.28		1	70		
222453	**Low discrimination**							**222453**
					NOV			
Staphylococcus saprophyticus subsp. bovis		68.90	0.50		99			
Staphylococcus simulans		19.20	0.39		1			
Staphylococcus cohnii subsp. urealyticum		11.28	0.40		99			
222460	**Good identification**							**222460**
Kocuria varians		100.0	0.41	TRE 7				
222462	**Good identification**							**222462**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.29	TRE 7				
222470	**Good identification**							**222470**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	XYL 1				
222473	**Low discrimination**							**222473**
					COA	NOV		
Staphylococcus xylosus		58.48	0.27	GLR 95	1	99		
Staphylococcus lutrae		35.89	0.18	GAL 91	99	1		
Staphylococcus saprophyticus subsp. bovis		5.62	0.09	SUC 95	1	99		
222500	**Very good identification**							**222500**
					NOV	PYR		
Kocuria varians		99.57	0.67	TRE 7	1	10		
Staphylococcus auricularis		0.19	0.10		1	40		
Staphylococcus saprophyticus subsp. bovis		0.14	0.05		99	99		
222501	**Good identification**							**222501**
					COA	PYR		
Kocuria varians		99.08	0.45	TRE 7	1	10		
Staphylococcus intermedius		0.93	-.01	MNS 8	99	90		
222502	**Very good identification**							**222502**
Kocuria varians		99.71	0.56	TRE 7				
Staphylococcus simulans		0.30	0.01					
222503	**Low discrimination**							**222503**
					COA	PYR		
Kocuria varians		58.95	0.34	TRE 7	1	10		
Staphylococcus felis		22.35	0.14	MNS 8	1	20		
Staphylococcus intermedius		11.98	0.15		99	90		
222510	**Acceptable identification**							**222510**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		90.93	0.47	MLT 99	99	99	1	
Kocuria varians		6.85	0.26		1	10	13	
Macrococcus bovicus		2.23	0.12		99	1	99	
222511	**Acceptable identification**							**222511**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		91.61	0.34	MLT 99	1	99	99	
Staphylococcus intermedius		5.99	0.08		99	1	90	
Kocuria varians		2.40	0.04		1	1	10	
222512	**Low discrimination**							**222512**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		84.42	0.34	MLT 99	99	99		
Staphylococcus simulans		8.41	0.14		1	70		
Kocuria varians		7.17	0.15		1	10		
222513	**Low discrimination**							**222513**
					COA			
Staphylococcus felis		53.94	0.31	ARG 92	1			
Staphylococcus condimenti		13.93	0.14	PHS 99	1			
Staphylococcus intermedius		11.69	0.24		99			
222520	**Very good identification**							**222520**
Kocuria varians		100.0	0.58	TRE 7				
222521	**Good identification**							**222521**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.36	TRE 7				
222522	**Good identification**							**222522**
Kocuria varians		100.0	0.46	TRE 7				
222540	**Low discrimination**							**222540**
					NOV	PYR		
Kocuria varians		60.12	0.54	TRE 7	1	10		
Staphylococcus saprophyticus subsp. bovis		36.75	0.47		99	99		
Staphylococcus auricularis		1.23	0.18		1	40		
222541	**Low discrimination**							**222541**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		62.30	0.34	MAN 99	1	99	99	
Kocuria varians		35.44	0.32		1	1	10	
Staphylococcus intermedius		2.26	0.03		99	1	90	
222542	**Low discrimination**							**222542**
					NOV	PYR		
Kocuria varians		61.87	0.42	TRE 7	1	10		
Staphylococcus saprophyticus subsp. bovis		33.52	0.34		99	99		
Staphylococcus hominis subsp. hominis		3.21	0.17		1	1		
222550	**Very good identification**							**222550**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		99.89	0.88		1	99	99	
Micrococcus bovicus		0.06	0.19		99	99	1	
Staphylococcus warneri		0.02	0.22		1	1	8	
222551	**Very good identification**							**222551**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		99.66	0.75		1	99		
Staphylococcus xylosus		0.20	0.16		1	99		
Staphylococcus intermedius		0.10	0.12		99	1		
222552	**Very good identification**							**222552**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		99.83	0.75		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		0.09	0.02		99	7		
Staphylococcus simulans		0.05	0.06		1	70		
222553	**Good identification**							**222553**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		91.98	0.63		1	99		
Staphylococcus intermedius		2.24	0.28		99	1		
Staphylococcus xylosus		2.13	0.26		1	99		
222560	**Good identification**							**222560**
					NOV	PYR		
Kocuria varians		98.27	0.44	TRE 7	1	10		
Staphylococcus saprophyticus subsp. bovis		1.73	0.05		99	99		
222562	**Low discrimination**							**222562**
					COA			
Kocuria varians		75.93	0.33	TRE 7	1			
Staphylococcus lutrae		24.07	0.12		99			
222563	**Very good identification**							**222563**
					COA	NOV		
Staphylococcus lutrae		97.72	0.53	PHS 99	99	1		
Staphylococcus xylosus		2.02	0.23		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		0.27	0.11		1	1	
222570	**Good identification**						**222570**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.85	0.47	XYL 1	99	99	
Staphylococcus xylosus		0.58	-.03		99	65	
Kocuria varians		0.57	0.03		1	10	
222571	**Low discrimination**						**222571**
					COA	NOV	
Staphylococcus xylosus		62.28	0.38	GLR 95	1	99	
Staphylococcus saprophyticus subsp. bovis		26.96	0.34	SUC 95	1	99	
Staphylococcus lutrae		10.76	0.18		99	1	
222572	**Low discrimination**						**222572**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		68.24	0.34	XYL 1	1	99	
Staphylococcus lutrae		27.23	0.18		99	1	
Staphylococcus xylosus		4.53	0.06		1	99	
222573	**Low discrimination**						**222573**
					COA	NOV	
Staphylococcus lutrae		85.27	0.59	PHS 99	99	1	
Staphylococcus xylosus		14.19	0.47		1	99	
Staphylococcus saprophyticus subsp. bovis		0.54	0.22		1	99	
222600	**Low discrimination**						**222600**
Kocuria varians		71.89	0.46	SUC 12			
Staphylococcus auricularis		12.64	0.29	TRE 7			
Staphylococcus warneri		10.56	0.36				
222602	**Low discrimination**						**222602**
Staphylococcus simulans		86.88	0.51	ARG 92			
Kocuria varians		9.66	0.34				
Staphylococcus warneri		1.68	0.26				
222603	**Very good identification**						**222603**
					COA		
Staphylococcus simulans		97.64	0.62	ARG 92	1		
Staphylococcus felis		0.71	0.09		1		
Staphylococcus intermedius		0.63	0.15		99		
222610	**Low discrimination**						**222610**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		85.47	0.47	MLT 99	99	99	
Staphylococcus warneri		6.16	0.33		1	8	
Staphylococcus simulans		5.94	0.23		1	70	
222611	**Low discrimination**						**222611**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		49.41	0.34	MLT 99	1	99	
Staphylococcus simulans		45.96	0.34		1	1	
Staphylococcus intermedius		3.23	0.08		99	1	
222612	**Very good identification**						**222612**
					NOV	PYR	
Staphylococcus simulans		95.81	0.64	ARG 92	1	70	
Staphylococcus saprophyticus subsp. bovis		3.48	0.34		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		0.38	0.03		99	7	
222613	**Very good identification**						**222613**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA			
Staphylococcus simulans		97.99	0.75	ARG 92	1			
Staphylococcus felis		1.20	0.26		1			
Staphylococcus intermedius		0.43	0.24		99			
222620	**Good identification**							**222620**
Kocuria varians		100.0	0.36	SUC 12				
222633	**Low discrimination**							**222633**
					NOV			
Staphylococcus simulans		50.01	0.34	ARG 92	1			
Staphylococcus xylosus		49.99	0.30	XYL 1	99			
222640	**Low discrimination**							**222640**
					NOV	PYR		
Staphylococcus warneri		36.22	0.62		1	8		
Staphylococcus hominis subsp. hominis		35.72	0.56		1	1		
Staphylococcus saprophyticus subsp. bovis		19.17	0.47		99	99		
222641	**Low discrimination**							**222641**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		38.41	0.34	MAN 99	1	99	99	
Kocuria kristinae		34.01	0.28		99	1	93	
Staphylococcus warneri		5.93	0.27		1	1	8	
222642	**Low discrimination**							**222642**
Staphylococcus hominis subsp. hominis		54.30	0.58	BGA 1				
Staphylococcus simulans		19.17	0.44					
Staphylococcus warneri		15.63	0.52					
222643	**Low discrimination**							**222643**
					COA	NOV		
Staphylococcus simulans		89.48	0.55	ARG 92	1	1		
Staphylococcus saprophyticus subsp. bovis		2.16	0.22		1	99		
Staphylococcus intermedius		1.93	0.19		99	1		
222650	**Very good identification**							**222650**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		97.84	0.88		99	99		
Staphylococcus warneri		1.35	0.59		1	8		
Staphylococcus saprophyticus subsp. saprophyticus		0.46	0.29		99	7		
222651	**Very good identification**							**222651**
					NOV			
Staphylococcus saprophyticus subsp. bovis		98.94	0.75		99			
Staphylococcus simulans		0.42	0.26		1			
Staphylococcus xylosus		0.38	0.22		99			
222652	**Low discrimination**							**222652**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		79.11	0.75		99	99		
Staphylococcus simulans		9.88	0.57		1	70		
Staphylococcus saprophyticus subsp. saprophyticus		8.51	0.44		99	7		
222653	**Low discrimination**							**222653**
					NOV			
Staphylococcus simulans		60.81	0.67	ARG 92	1			
Staphylococcus saprophyticus subsp. bovis		36.38	0.63		99			
Staphylococcus xylosus		1.58	0.31		99			
222663	**Low discrimination**							**222663**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	NOV		
Staphylococcus xylosus		71.78	0.28	GLR 95	1	99		
Staphylococcus lutrae		18.71	0.12	GAL 91	99	1		
Staphylococcus simulans		9.51	0.14	MAN 94	1	1		
222670	**Good identification**							**222670**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		97.58	0.47	XYL 1	99	99		
Staphylococcus warneri		1.35	0.18		1	8		
Staphylococcus xylosus		1.07	0.03		99	65		
222671	**Low discrimination**							**222671**
					PYR			
Staphylococcus xylosus		80.55	0.44	GLR 95	65			
Staphylococcus saprophyticus subsp. bovis		18.56	0.34	GAL 91	99			
Staphylococcus equorum		0.89	0.05		5			
222672	**Low discrimination**							**222672**
					NOV			
Staphylococcus saprophyticus subsp. bovis		73.68	0.34	XYL 1	99			
Staphylococcus simulans		9.20	0.16		1			
Staphylococcus xylosus		9.19	0.12		99			
222673	**Good identification**							**222673**
					COA	NOV		
Staphylococcus xylosus		91.21	0.53	GLR 95	1	99		
Staphylococcus simulans		3.09	0.26	GAL 91	1	1		
Staphylococcus lutrae		2.95	0.18		99	1		
222700	**Acceptable identification**							**222700**
					NOV	PYR		
Kocuria varians		90.11	0.49	SUC 12	1	10		
Staphylococcus saprophyticus subsp. bovis		3.58	0.18	TRE 7	99	99		
Staphylococcus warneri		2.79	0.25		1	8		
222701	**Low discrimination**							**222701**
					COA	YEP	PYR	
Staphylococcus intermedius		79.34	0.40	PHS 97	99	1	90	
Kocuria varians		11.71	0.27	LAC 86	1	80	10	
Staphylococcus chromogenes		4.04	0.12		1	95	45	
222702	**Low discrimination**							**222702**
					COA	PYR		
Kocuria varians		55.35	0.38	SUC 12	1	10		
Staphylococcus simulans		28.87	0.30	TRE 7	1	70		
Staphylococcus intermedius		7.18	0.15		99	90		
222703	**Low discrimination**							**222703**
					COA	YEP		
Staphylococcus intermedius		78.11	0.56	PHS 97	99	1		
Staphylococcus simulans		10.63	0.40		1	1		
Staphylococcus chromogenes		5.83	0.32		1	95		
222710	**Very good identification**							**222710**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.49	0.59	MLT 99	1	99	99	
Staphylococcus warneri		0.56	0.22		1	1	8	
Staphylococcus intermedius		0.40	0.08		99	1	90	
222711	**Low discrimination**							**222711**
					COA	NOV		
Staphylococcus intermedius		59.32	0.49	PHS 97	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. bovis		36.64	0.47	LAC 86	1	99		
Staphylococcus xylosus		2.63	0.19		1	99		
222712	**Low discrimination**							**222712**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		58.29	0.47	MLT 99	1	99		
Staphylococcus simulans		34.86	0.42		1	1		
Staphylococcus intermedius		5.86	0.24		99	1		
222713	**Low discrimination**							**222713**
					COA			
Staphylococcus intermedius		73.05	0.65	PHS 97	99			
Staphylococcus simulans		14.70	0.53		1			
Staphylococcus felis		6.91	0.37		1			
222720	**Good identification**							**222720**
Kocuria varians		100.0	0.40	SUC 12				
222722	**Good identification**							**222722**
Kocuria varians		100.0	0.29	SUC 12				
222731	**Good identification**							**222731**
					COA	NOV		
Staphylococcus xylosus		95.56	0.41	GLR 95	1	99		
Staphylococcus intermedius		1.89	0.08	MLT 93	99	1		
Staphylococcus equorum		1.38	0.05		1	99		
222733	**Low discrimination**							**222733**
					COA	PYR	NOV	
Staphylococcus xylosus		89.50	0.51	GLR 95	1	65	99	
Staphylococcus intermedius		3.83	0.24	MLT 93	99	90	1	
Staphylococcus lutrae		3.80	0.18		99	10	1	
222740	**Low discrimination**							**222740**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		57.93	0.59	MAN 99	99	99		
Staphylococcus hominis subsp. hominis		29.24	0.57		1	1		
Staphylococcus warneri		8.64	0.52		1	8		
222741	**Low discrimination**							**222741**
					OXI	COA	NOV	
Staphylococcus saprophyticus subsp. bovis		46.20	0.47	MAN 99	1	1	99	
Staphylococcus intermedius		41.50	0.44		1	99	1	
Kocuria kristinae		3.41	0.18		99	1	1	
222742	**Low discrimination**							**222742**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		62.77	0.58	BGA 1	1	1		
Staphylococcus saprophyticus subsp. bovis		26.48	0.47		99	99		
Staphylococcus warneri		5.27	0.42		1	8		
222743	**Low discrimination**							**222743**
					COA	NOV	YEP	
Staphylococcus intermedius		83.70	0.60	PHS 97	99	1	1	
Staphylococcus chromogenes		4.16	0.32		1	1	95	
Staphylococcus saprophyticus subsp. bovis		3.79	0.34		1	99	80	
222750	**Excellent identification**							**222750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		99.79	1.00		99	99		
Staphylococcus warneri		0.11	0.49		1	8		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		0.08	0.40		1	1		
222751	**Very good identification**							**222751**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		96.66	0.88		1	99		
Staphylococcus intermedius		2.37	0.53		99	1		
Staphylococcus xylosus		0.93	0.43		1	99		
222752	**Very good identification**							**222752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		98.83	0.88		99	99		
Staphylococcus hominis subsp. hominis		0.35	0.41		1	1		
Staphylococcus simulans		0.27	0.35		1	70		
222753	**Low discrimination**							**222753**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		56.89	0.75		1	99		
Staphylococcus intermedius		34.29	0.69		99	1		
Staphylococcus xylosus		6.24	0.52		1	99		
222761	**Acceptable identification**							**222761**
					COA	NOV		
Staphylococcus xylosus		90.05	0.40	GLR 95	1	99		
Staphylococcus lutrae		6.61	0.12	MAN 94	99	1		
Staphylococcus saprophyticus subsp. bovis		1.30	0.05		1	99		
222763	**Low discrimination**							**222763**
					COA	PYR	NOV	
Staphylococcus lutrae		71.29	0.53	PHS 99	99	10	1	
Staphylococcus xylosus		27.93	0.49		1	65	99	
Staphylococcus intermedius		0.78	0.19		99	90	1	
222770	**Very good identification**							**222770**
Staphylococcus saprophyticus subsp. bovis		97.30	0.59	XYL 1				
Staphylococcus xylosus		2.70	0.23					
222771	**Low discrimination**							**222771**
					PYR			
Staphylococcus xylosus		89.67	0.64	GLR 95	65			
Staphylococcus saprophyticus subsp. bovis		8.17	0.47		99			
Staphylococcus equorum		0.99	0.25		5			
222772	**Low discrimination**							**222772**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		70.66	0.47	XYL 1	1	99		
Staphylococcus xylosus		22.29	0.33		1	99		
Staphylococcus lutrae		7.05	0.18		99	1		
222773	**Low discrimination**							**222773**
					COA	NOV		
Staphylococcus xylosus		74.56	0.74	GLR 95	1	99		
Staphylococcus lutrae		23.58	0.59		99	1		
Staphylococcus equorum		0.74	0.34		1	99		
223000	**Acceptable identification**							**223000**
					OXI			
Kocuria varians		94.29	0.46	ESL 1	13			
Dermacoccus nishinomiyaensis		5.71	0.13		93			
223002	**Good identification**							**223002**
Kocuria varians		100.0	0.34	ESL 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
223020	**Good identification**						**223020**
Kocuria varians		100.0	0.36	ESL 1			
223040	**Good identification**						**223040**
Kocuria varians		100.0	0.32	ESL 1			
223100	**Good identification**						**223100**
					OXI		
Kocuria varians		99.24	0.49	ESL 1	13		
Dermacoccus nishinomiyaensis		0.76	-.02		93		
223101	**Good identification**						**223101**
Kocuria varians		100.0	0.27	ESL 1			
223102	**Good identification**						**223102**
Kocuria varians		100.0	0.38	ESL 1			
223120	**Good identification**						**223120**
Kocuria varians		100.0	0.40	ESL 1			
223122	**Good identification**						**223122**
Kocuria varians		100.0	0.28	ESL 1			
223140	**Good identification**						**223140**
Kocuria varians		100.0	0.36	ESL 1			
223200	**Good identification**						**223200**
Kocuria varians		100.0	0.28	ESL 1			
223300	**Good identification**						**223300**
Kocuria varians		100.0	0.31	ESL 1			
223373	**Low discrimination**						**223373**
					OXI		
Staphylococcus xylosus		46.95	0.33	GLR 95	1		
Staphylococcus equorum		44.85	0.34	TRE 96	1		
Staphylococcus lentus		4.42	0.18		99		
223450	**Low discrimination**						**223450**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		81.74	0.34	ESL 1	1	99	
Macrococcus equipersicus		13.43	0.18		99	1	
Macrococcus bovicus		4.84	0.07		99	1	
223453	**Low discrimination**						**223453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		82.60	0.31	GLR 99	99		
Staphylococcus saprophyticus subsp. bovis		13.11	0.09		99		
Aerococcus viridans		4.29	0.16		1		
223550	**Good identification**						**223550**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		97.66	0.47	ESL 1	1	99	
Macrococcus equipersicus		1.72	0.10		99	1	
Macrococcus bovicus		0.62	0.00		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
223551	**Acceptable identification**						**223551**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		92.19	0.34	ESL 1	1	99	
Staphylococcus xylosus		4.63	0.04		1	65	
Staphylococcus lentus		3.18	0.07		99	6	
223552	**Good identification**						**223552**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ESL 1			
223573	**Low discrimination**						**223573**
					COA	NOV	
Staphylococcus xylosus		46.51	0.35	GLR 95	1	99	
Staphylococcus equorum		35.18	0.34	SUC 95	1	99	
Staphylococcus lutrae		11.29	0.18		99	1	
223613	**Acceptable identification**						**223613**
					OXI	NOV	
Staphylococcus simulans		92.04	0.34	ARG 92	1	1	
Staphylococcus lentus		5.96	0.12	ESL 1	99	99	
Staphylococcus xylosus		2.00	-.04		1	99	
223633	**Low discrimination**						**223633**
					OXI		
Staphylococcus equorum		80.44	0.34	GLR 90	1		
Staphylococcus xylosus		15.54	0.18	GAL 91	1		
Staphylococcus lentus		4.03	0.12	MLT 91	99		
223650	**Good identification**						**223650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.64	0.47	ESL 1	99	99	
Staphylococcus warneri		1.36	0.18		1	8	
223651	**Low discrimination**						**223651**
					PYR		
Staphylococcus saprophyticus subsp. bovis		81.85	0.34	ESL 1	99		
Staphylococcus xylosus		7.72	0.09		65		
Staphylococcus equorum		3.91	0.05		5		
223652	**Low discrimination**						**223652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		81.14	0.34	ESL 1	99	99	
Staphylococcus simulans		10.13	0.16		1	70	
Staphylococcus saprophyticus subsp. saprophyticus		8.73	0.03		99	7	
223671	**Low discrimination**						**223671**
Staphylococcus equorum		79.82	0.46	GLR 90			
Staphylococcus xylosus		18.30	0.31	GAL 91			
Staphylococcus gallinarum		0.72	0.01				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
223673	**Low discrimination**							**223673**
					OXI			
Staphylococcus equorum		78.48	0.54	GLR 90	1			
Staphylococcus xylosus		19.92	0.41	GAL 91	1			
Staphylococcus lentus		0.79	0.18		99			
223711	**Low discrimination**							**223711**
					OXI	COA	NOV	
Staphylococcus lentus		87.13	0.41	URE 1	99	1	99	
Staphylococcus xylosus		3.72	0.07		1	1	99	
Staphylococcus intermedius		3.39	0.08		1	99	1	
223713	**Low discrimination**							**223713**
					OXI	COA	NOV	
Staphylococcus lentus		87.56	0.53	URE 1	99	1	99	
Staphylococcus intermedius		5.90	0.24		1	99	1	
Staphylococcus xylosus		3.00	0.16		1	1	99	
223731	**Low discrimination**							**223731**
					OXI			
Staphylococcus equorum		64.97	0.46	GLR 90	1			
Staphylococcus lentus		23.10	0.41	MLT 91	99			
Staphylococcus xylosus		11.34	0.29		1			
223733	**Low discrimination**							**223733**
					OXI			
Staphylococcus equorum		59.10	0.54	GLR 90	1			
Staphylococcus lentus		28.98	0.53	MLT 91	99			
Staphylococcus xylosus		11.41	0.38		1			
223750	**Very good identification**							**223750**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		99.79	0.59	ESL 1	1	99		
Staphylococcus lentus		0.22	0.07		99	6		
223751	**Low discrimination**							**223751**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		44.04	0.47	ESL 1	1	99		
Staphylococcus lentus		37.62	0.48		99	6		
Staphylococcus xylosus		10.50	0.30		1	65		
223752	**Good identification**							**223752**
					OXI	PYR		
Staphylococcus saprophyticus subsp. bovis		95.92	0.47	ESL 1	1	99		
Staphylococcus lentus		2.93	0.18		99	6		
Staphylococcus xylosus		0.66	-.02		1	65		
223753	**Low discrimination**							**223753**
					OXI			
Staphylococcus lentus		67.92	0.59	URE 1	99			
Staphylococcus xylosus		15.22	0.40		1			
Staphylococcus equorum		6.97	0.34		1			
223761	**Low discrimination**							**223761**
Staphylococcus xylosus		42.74	0.27	GLR 95				
Staphylococcus equorum		29.50	0.25	MAN 94				
Staphylococcus gallinarum		25.66	0.21					
223763	**Low discrimination**							**223763**
Staphylococcus xylosus		43.39	0.37	GLR 95				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus equorum		27.06	0.34	MAN 94		
Staphylococcus gallinarum		22.41	0.29			
223771	**Low discrimination**					**223771**
Staphylococcus equorum		71.85	0.67	GLR 90		
Staphylococcus xylosus		16.48	0.52			
Staphylococcus gallinarum		6.38	0.42			
223772	**Low discrimination**					**223772**
					OXI	
Staphylococcus equorum		68.06	0.34	GLR 90	1	
Staphylococcus xylosus		17.27	0.20	MNS 99	1	
Staphylococcus lentus		6.70	0.18		99	
223773	**Low discrimination**					**223773**
					OXI	
Staphylococcus equorum		69.43	0.75	GLR 90	1	
Staphylococcus xylosus		17.62	0.61		1	
Staphylococcus lentus		6.84	0.59		99	
224000	**Low discrimination**					**224000**
					OXI	
Dermacoccus nishinomiyaensis		68.12	0.33	BGA 1	93	
Kocuria varians		24.87	0.31	PHS 5	13	
Micrococcus sp.		7.01	0.03		89	
224050	**Low discrimination**					**224050**
					OXI	
Staphylococcus kloosii		83.76	0.46	TRE 99	1	
Macrococcus equipersicus		11.46	0.11		99	
Macrococcus bovicus		4.79	0.02		99	
224052	**Good identification**					**224052**
Staphylococcus kloosii		100.0	0.47	TRE 99		
224070	**Good identification**					**224070**
Staphylococcus kloosii		100.0	0.34	TRE 99		
224072	**Good identification**					**224072**
Staphylococcus kloosii		100.0	0.35	TRE 99		
224100	**Low discrimination**					**224100**
					OXI	
Kocuria varians		74.20	0.35	PHS 1	13	
Dermacoccus nishinomiyaensis		25.80	0.18		93	
224112	**Good identification**					**224112**
Staphylococcus kloosii		100.0	0.27	TRE 99		
224150	**Good identification**					**224150**
					OXI	
Staphylococcus kloosii		97.07	0.52	TRE 99	1	
Macrococcus equipersicus		2.93	0.04		99	
224152	**Very good identification**					**224152**
Staphylococcus kloosii		100.0	0.53	TRE 99		
224170	**Good identification**					**224170**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.40	TRE 99				
224172	**Good identification**							**224172**
Staphylococcus kloosii		100.0	0.41	TRE 99				
224303	**Good identification**							**224303**
					COA	YEP		
Staphylococcus intermedius		96.58	0.41	NIT 93	99	1		
Staphylococcus chromogenes		3.42	0.10	TRE 93	1	95		
224311	**Good identification**							**224311**
Staphylococcus intermedius		100.0	0.34	NIT 93				
224313	**Good identification**							**224313**
					COA	YEP		
Staphylococcus intermedius		99.12	0.50	NIT 93	99	1		
Staphylococcus felis		0.52	-.04	TRE 93	1	1		
Staphylococcus chromogenes		0.37	-.01		1	95		
224341	**Low discrimination**							**224341**
					COA	PYR		
Staphylococcus intermedius		83.57	0.29	NIT 93	99	90		
Staphylococcus epidermidis		16.43	0.16	TRE 93	1	1		
224343	**Low discrimination**							**224343**
					COA	YEP	PYR	
Staphylococcus intermedius		88.89	0.45	NIT 93	99	1	90	
Staphylococcus epidermidis		9.01	0.27	TRE 93	1	1	1	
Staphylococcus chromogenes		2.10	0.10		1	95	45	
224350	**Low discrimination**							**224350**
Staphylococcus kloosii		72.86	0.32	SUC 9				
Staphylococcus saprophyticus subsp. bovis		27.14	0.05	TRE 99				
224351	**Good identification**							**224351**
Staphylococcus intermedius		100.0	0.38	NIT 93				
224352	**Low discrimination**							**224352**
					COA	NOV		
Staphylococcus kloosii		51.56	0.32	SUC 9	1	99		
Staphylococcus intermedius		48.44	0.13	TRE 99	99	1		
224353	**Very good identification**							**224353**
					COA	YEP		
Staphylococcus intermedius		99.75	0.54	NIT 93	99	1		
Staphylococcus chromogenes		0.25	-.01	TRE 93	1	95		
224410	**Very good identification**							**224410**
					OXI			
Staphylococcus kloosii		98.35	0.61	MLT 95	1			
Micrococcus equipersicus		0.96	0.02		99			
Micrococcus bovicus		0.69	-.02		99			
224412	**Very good identification**							**224412**
Staphylococcus kloosii		100.0	0.62	MLT 95				
224413	**Low discrimination**							**224413**
					NOV			
Staphylococcus cohnii subsp. urealyticum		70.37	0.43	GLR 99	99			
Staphylococcus felis		25.67	0.20		1			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		2.47	0.08		1		
224430	**Good identification**						**224430**
Staphylococcus kloosii		100.0	0.49	MLT 95			
224432	**Good identification**						**224432**
Staphylococcus kloosii		100.0	0.50	MLT 95			
224440	**Good identification**						**224440**
Staphylococcus kloosii		100.0	0.46	MAN 99			
224442	**Acceptable identification**						**224442**
Staphylococcus kloosii		93.31	0.47	MAN 99			
Staphylococcus saprophyticus subsp. saprophyticus		6.69	-.05				
224443	**Good identification**						**224443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	GLR 99			
224450	**Very good identification**						**224450**
					PYR		
Staphylococcus kloosii		99.01	0.88		52		
Staphylococcus saprophyticus subsp. bovis		0.44	0.22		99		
Staphylococcus saprophyticus subsp. saprophyticus		0.33	0.08		7		
224451	**Low discrimination**						**224451**
Staphylococcus cohnii subsp. urealyticum		57.56	0.38	GLR 99			
Staphylococcus kloosii		38.21	0.46	LAC 91			
Staphylococcus saprophyticus subsp. bovis		4.23	0.09				
224452	**Very good identification**						**224452**
Staphylococcus kloosii		98.05	0.88				
Staphylococcus saprophyticus subsp. saprophyticus		1.70	0.23				
Staphylococcus cohnii subsp. urealyticum		0.14	0.18				
224453	**Good identification**						**224453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		93.12	0.59	GLR 99	99	64	
Staphylococcus kloosii		6.62	0.47		99	99	
Staphylococcus simulans		0.26	0.01		1	1	
224460	**Good identification**						**224460**
Staphylococcus kloosii		100.0	0.34	MAN 99			
224462	**Good identification**						**224462**
Staphylococcus kloosii		100.0	0.35	MAN 99			
224470	**Very good identification**						**224470**
Staphylococcus kloosii		100.0	0.75				
224471	**Good identification**						**224471**
Staphylococcus kloosii		100.0	0.34	MNS 1			
224472	**Very good identification**						**224472**
Staphylococcus kloosii		100.0	0.76				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
224502	**Good identification**							**224502**
Staphylococcus kloosii		100.0	0.27	MAN 99				
224510	**Very good identification**							**224510**
Staphylococcus kloosii		100.0	0.67	MLT 95				
224512	**Very good identification**							**224512**
Staphylococcus kloosii		100.0	0.68	MLT 95				
224513	**Low discrimination**							**224513**
					COA	NOV		
Staphylococcus felis		46.08	0.31	ARG 92	1	1		
Staphylococcus intermedius		24.30	0.32	NIT 99	99	1		
Staphylococcus cohnii subsp. urealyticum		16.17	0.36		1	99		
224530	**Very good identification**							**224530**
Staphylococcus kloosii		100.0	0.55	MLT 95				
224532	**Very good identification**							**224532**
Staphylococcus kloosii		100.0	0.55	MLT 95				
224540	**Good identification**							**224540**
Staphylococcus kloosii		100.0	0.52	MAN 99				
224542	**Very good identification**							**224542**
Staphylococcus kloosii		100.0	0.53	MAN 99				
224543	**Low discrimination**							**224543**
					COA	PYR	NOV	
Staphylococcus intermedius		48.75	0.27	NIT 93	99	90	1	
Staphylococcus cohnii subsp. urealyticum		32.78	0.31	SUC 99	1	53	99	
Staphylococcus lutrae		18.47	0.12		99	10	1	
224550	**Very good identification**							**224550**
					OXI	PYR		
Staphylococcus kloosii		99.04	0.93		1	52		
Staphylococcus saprophyticus subsp. bovis		0.91	0.34		1	99		
Micrococcus equipersicus		0.03	0.04		99	1		
224551	**Low discrimination**							**224551**
Staphylococcus kloosii		56.24	0.52	MNS 1				
Staphylococcus cohnii subsp. urealyticum		18.70	0.31					
Staphylococcus saprophyticus subsp. bovis		12.82	0.22					
224552	**Excellent identification**							**224552**
					PYR			
Staphylococcus kloosii		99.68	0.94		52			
Staphylococcus saprophyticus subsp. bovis		0.21	0.22		99			
Staphylococcus saprophyticus subsp. saprophyticus		0.08	0.02		7			
224553	**Low discrimination**							**224553**
					COA	NOV		
Staphylococcus cohnii subsp. urealyticum		54.04	0.51	GLR 99	1	99		
Staphylococcus intermedius		21.49	0.36		99	1		
Staphylococcus kloosii		17.41	0.53		1	99		
224560	**Good identification**							**224560**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.40	MAN 99			
224562	**Low discrimination**						**224562**
					COA	NOV	
Staphylococcus kloosii		59.10	0.41	MAN 99	1	99	
Staphylococcus lutrae		40.90	0.12		99	1	
224563	**Very good identification**						**224563**
Staphylococcus lutrae		100.0	0.53	NIT 99			
224570	**Very good identification**						**224570**
Staphylococcus kloosii		100.0	0.81				
224572	**Very good identification**						**224572**
					COA	NOV	
Staphylococcus kloosii		98.66	0.82		1	99	
Staphylococcus lutrae		1.34	0.18		99	1	
224573	**Very good identification**						**224573**
					COA	NOV	
Staphylococcus lutrae		98.59	0.59	NIT 99	99	1	
Staphylococcus kloosii		0.74	0.41		1	99	
Staphylococcus xylosus		0.57	0.17		1	99	
224610	**Good identification**						**224610**
Staphylococcus kloosii		100.0	0.41	SUC 9			
224613	**Low discrimination**						**224613**
					COA		
Staphylococcus simulans		36.93	0.36	ARG 92	1		
Staphylococcus felis		32.48	0.26	NIT 96	1		
Staphylococcus intermedius		28.24	0.32		99		
224630	**Good identification**						**224630**
Staphylococcus kloosii		100.0	0.28	SUC 9			
224632	**Good identification**						**224632**
Staphylococcus kloosii		100.0	0.29	SUC 9			
224640	**Low discrimination**						**224640**
					NOV		
Staphylococcus warneri		57.04	0.28	PHS 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		25.07	0.00		99		
Staphylococcus kloosii		17.89	0.26		99		
224643	**Low discrimination**						**224643**
					COA	YEP	
Staphylococcus intermedius		62.78	0.27	NIT 93	99	1	
Staphylococcus simulans		16.62	0.17	GAL 99	1	1	
Staphylococcus chromogenes		14.11	0.12		1	95	
224650	**Low discrimination**						**224650**
					PYR		
Staphylococcus kloosii		64.78	0.67	SUC 9	52		
Staphylococcus saprophyticus subsp. saprophyticus		22.01	0.29		7		
Staphylococcus saprophyticus subsp. bovis		11.71	0.34		99		
224652	**Low discrimination**						**224652**
					PYR		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. saprophyticus		62.25	0.44	PHS 1	7			
Staphylococcus kloosii		35.03	0.68		52			
Staphylococcus saprophyticus subsp. bovis		1.46	0.22		99			
224653	**Low discrimination**							**224653**
					COA	NOV		
Staphylococcus intermedius		56.20	0.36	NIT 93	99	1		
Staphylococcus simulans		22.02	0.29	GAL 99	1	1		
Staphylococcus cohnii subsp. urealyticum		10.20	0.28		1	99		
224670	**Very good identification**							**224670**
Staphylococcus kloosii		100.0	0.54	SUC 9				
224672	**Low discrimination**							**224672**
Staphylococcus kloosii		88.01	0.55	SUC 9				
Staphylococcus saprophyticus subsp. saprophyticus		6.32	0.03					
Staphylococcus arlettae		5.67	0.19					
224701	**Good identification**							**224701**
					COA	YEP		
Staphylococcus intermedius		97.95	0.48	NIT 93	99	1		
Staphylococcus chromogenes		2.05	0.12	LAC 86	1	95		
224703	**Very good identification**							**224703**
					COA	YEP		
Staphylococcus intermedius		95.41	0.64	NIT 93	99	1		
Staphylococcus chromogenes		2.92	0.32		1	95		
Staphylococcus felis		1.50	0.20		1	1		
224710	**Low discrimination**							**224710**
					COA	NOV		
Staphylococcus kloosii		73.90	0.46	SUC 9	1	99		
Staphylococcus intermedius		20.82	0.16	MLT 95	99	1		
Staphylococcus saprophyticus subsp. bovis		5.28	0.05		1	99		
224711	**Very good identification**							**224711**
					COA	YEP		
Staphylococcus intermedius		99.54	0.57	NIT 93	99	1		
Staphylococcus felis		0.24	-.04	LAC 86	1	1		
Staphylococcus chromogenes		0.22	0.01		1	95		
224712	**Low discrimination**							**224712**
					COA	NOV	YEP	
Staphylococcus intermedius		58.86	0.32	NIT 93	99	1	1	
Staphylococcus kloosii		36.85	0.47	MNS 99	1	99	99	
Staphylococcus felis		2.29	-.04		1	1	1	
224713	**Very good identification**							**224713**
					COA			
Staphylococcus intermedius		95.13	0.73	NIT 93	99			
Staphylococcus felis		3.70	0.37		1			
Staphylococcus condimenti		0.71	0.18		1			
224730	**Good identification**							**224730**
Staphylococcus kloosii		100.0	0.34	SUC 9				
224732	**Good identification**							**224732**
Staphylococcus kloosii		100.0	0.35	SUC 9				
224733	**Low discrimination**							**224733**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
					COA	PYR	NOV	
	Staphylococcus intermedius	56.11	0.32	NIT 93	99	90	1	
	Staphylococcus lutrae	22.90	0.18	XYL 1	99	10	1	
	Staphylococcus xylosus	18.81	0.21		1	65	99	
224740	**Low discrimination**							**224740**
					COA	NOV	PYR	
	Staphylococcus kloosii	36.97	0.32	SUC 9	1	99	52	
	Staphylococcus intermedius	30.10	0.11	MAN 99	99	1	90	
	Staphylococcus warneri	19.17	0.18		1	1	8	
224741	**Good identification**							**224741**
					OXI	COA	YEP	
	Staphylococcus intermedius	98.19	0.52	NIT 93	1	99	1	
	Staphylococcus chromogenes	1.37	0.12	LAC 86	1	1	95	
	Kocuria kristinae	0.45	0.00		99	1	15	
224742	**Low discrimination**							**224742**
					COA	NOV		
	Staphylococcus intermedius	77.66	0.27	NIT 93	99	1		
	Staphylococcus kloosii	16.82	0.32	MNS 99	1	99		
	Staphylococcus saprophyticus subsp. saprophyticus	5.52	-.06		1	99		
224743	**Very good identification**							**224743**
					COA	PYR	YEP	
	Staphylococcus intermedius	97.59	0.68	NIT 93	99	90	1	
	Staphylococcus chromogenes	1.99	0.32		1	45	95	
	Staphylococcus lutrae	0.37	0.12		99	10	1	
224750	**Low discrimination**							**224750**
					COA	NOV		
	Staphylococcus kloosii	70.74	0.73	SUC 9	1	99		
	Staphylococcus saprophyticus subsp. bovis	26.35	0.47		1	99		
	Staphylococcus intermedius	1.57	0.20		99	1		
224751	**Good identification**							**224751**
					COA	NOV		
	Staphylococcus intermedius	94.88	0.61	NIT 93	99	1		
	Staphylococcus saprophyticus subsp. bovis	4.01	0.34	LAC 86	1	99		
	Staphylococcus xylosus	0.53	0.13		1	99		
224752	**Low discrimination**							**224752**
					COA	NOV		
	Staphylococcus kloosii	77.42	0.73	SUC 9	1	99		
	Staphylococcus intermedius	9.76	0.36		99	1		
	Staphylococcus saprophyticus subsp. bovis	6.65	0.34		1	99		
224753	**Very good identification**							**224753**
					COA	PYR	NOV	
	Staphylococcus intermedius	98.87	0.77	NIT 93	99	90	1	
	Staphylococcus lutrae	0.27	0.18		99	10	1	
	Staphylococcus xylosus	0.26	0.22		1	65	99	
224763	**Very good identification**							**224763**
					COA	PYR	NOV	
	Staphylococcus lutrae	96.13	0.53	NIT 99	99	10	1	
	Staphylococcus intermedius	2.56	0.27		99	90	1	
	Staphylococcus xylosus	1.31	0.19		1	65	99	
224770	**Very good identification**							**224770**
	Staphylococcus kloosii	98.52	0.60	SUC 9				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. bovis		1.48	0.05					
224771	**Low discrimination**							**224771**
					COA	PYR	NOV	
Staphylococcus xylosus		66.49	0.34	GLR 95	1	65	99	
Staphylococcus lutrae		17.34	0.18	NIT 89	99	10	1	
Staphylococcus intermedius		10.38	0.20		99	90	1	
224772	**Low discrimination**							**224772**
					COA	NOV		
Staphylococcus kloosii		74.61	0.61	SUC 9	1	99		
Staphylococcus arlettae		14.02	0.35		1	99		
Staphylococcus lutrae		10.24	0.18		99	1		
224773	**Low discrimination**							**224773**
					COA	PYR	NOV	
Staphylococcus lutrae		85.24	0.59	NIT 99	99	10	1	
Staphylococcus xylosus		9.40	0.44		1	65	99	
Staphylococcus intermedius		3.17	0.36		99	90	1	
225050	**Low discrimination**							**225050**
					OXI			
Staphylococcus kloosii		77.48	0.46	TRE 99	1			
Macrococcus equipersicus		22.52	0.18		99			
225052	**Good identification**							**225052**
Staphylococcus kloosii		100.0	0.47	TRE 99				
225070	**Good identification**							**225070**
Staphylococcus kloosii		100.0	0.34	TRE 99				
225072	**Good identification**							**225072**
Staphylococcus kloosii		100.0	0.35	TRE 99				
225112	**Good identification**							**225112**
Staphylococcus kloosii		100.0	0.27	TRE 99				
225150	**Acceptable identification**							**225150**
					OXI			
Staphylococcus kloosii		93.97	0.52	TRE 99	1			
Macrococcus equipersicus		6.03	0.10		99			
225152	**Very good identification**							**225152**
Staphylococcus kloosii		100.0	0.53	TRE 99				
225170	**Good identification**							**225170**
Staphylococcus kloosii		100.0	0.40	TRE 99				
225172	**Good identification**							**225172**
Staphylococcus kloosii		100.0	0.41	TRE 99				
225350	**Good identification**							**225350**
Staphylococcus kloosii		100.0	0.32	SUC 9				
225352	**Good identification**							**225352**
Staphylococcus kloosii		100.0	0.32	SUC 9				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
225410	**Very good identification**						**225410**
					OXI		
Staphylococcus kloosii		97.98	0.61	MLT 95	1		
Micrococcus equipersicus		2.02	0.09		99		
225412	**Very good identification**						**225412**
Staphylococcus kloosii		100.0	0.62	MLT 95			
225413	**Acceptable identification**						**225413**
Staphylococcus cohnii subsp. urealyticum		94.83	0.35	GLR 99			
Staphylococcus kloosii		5.17	0.21				
225430	**Good identification**						**225430**
Staphylococcus kloosii		100.0	0.49	MLT 95			
225432	**Good identification**						**225432**
Staphylococcus kloosii		100.0	0.50	MLT 95			
225440	**Good identification**						**225440**
Staphylococcus kloosii		100.0	0.46	MAN 99			
225442	**Good identification**						**225442**
Staphylococcus kloosii		100.0	0.47	MAN 99			
225443	**Good identification**						**225443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	GLR 99			
225450	**Very good identification**						**225450**
					OXI		
Staphylococcus kloosii		99.71	0.88		1		
Micrococcus equipersicus		0.29	0.18		99		
225451	**Low discrimination**						**225451**
Staphylococcus kloosii		63.06	0.46	MNS 1			
Staphylococcus cohnii subsp. urealyticum		36.94	0.30				
225452	**Very good identification**						**225452**
Staphylococcus kloosii		99.94	0.88				
Staphylococcus cohnii subsp. urealyticum		0.06	0.09				
225453	**Low discrimination**						**225453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		84.10	0.51	GLR 99	99	64	
Staphylococcus kloosii		15.38	0.47		99	99	
Aerococcus viridans		0.51	0.16		1	1	
225460	**Good identification**						**225460**
Staphylococcus kloosii		100.0	0.34	MAN 99			
225462	**Good identification**						**225462**
Staphylococcus kloosii		100.0	0.35	MAN 99			
225470	**Very good identification**						**225470**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.75				
225471	**Good identification**						**225471**
Staphylococcus kloosii		100.0	0.34	MNS 1			
225472	**Very good identification**						**225472**
Staphylococcus kloosii		100.0	0.76				
225473	**Low discrimination**						**225473**
					NOV	YEP	
Staphylococcus kloosii		73.82	0.35	MNS 1	99	99	
Staphylococcus cohnii subsp. urealyticum		16.31	0.09		99	64	
Aerococcus viridans		9.87	0.16		1	1	
225502	**Good identification**						**225502**
Staphylococcus kloosii		100.0	0.27	MAN 99			
225510	**Very good identification**						**225510**
					OXI		
Staphylococcus kloosii		99.55	0.67	MLT 95	1		
Macrococcus equipersicus		0.45	0.01		99		
225512	**Very good identification**						**225512**
Staphylococcus kloosii		100.0	0.68	MLT 95			
225513	**Low discrimination**						**225513**
Staphylococcus cohnii subsp. urealyticum		80.19	0.27	GLR 99			
Staphylococcus kloosii		19.81	0.27				
225530	**Very good identification**						**225530**
Staphylococcus kloosii		100.0	0.55	MLT 95			
225532	**Very good identification**						**225532**
Staphylococcus kloosii		100.0	0.55	MLT 95			
225540	**Good identification**						**225540**
Staphylococcus kloosii		100.0	0.52	MAN 99			
225542	**Very good identification**						**225542**
Staphylococcus kloosii		100.0	0.53	MAN 99			
225550	**Very good identification**						**225550**
					OXI		
Staphylococcus kloosii		99.94	0.93		1		
Macrococcus equipersicus		0.07	0.10		99		
225551	**Low discrimination**						**225551**
Staphylococcus kloosii		88.55	0.52	MNS 1			
Staphylococcus cohnii subsp. urealyticum		11.45	0.22				
225552	**Excellent identification**						**225552**
Staphylococcus kloosii		100.0	0.94				
225553	**Low discrimination**						**225553**
					NOV	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		54.27	0.43	GLR 99	99	64	
Staphylococcus kloosii		44.96	0.53		99	99	
Aerococcus viridans		0.77	0.16		1	1	
225560	**Good identification**						**225560**
Staphylococcus kloosii		100.0	0.40	MAN 99			
225562	**Good identification**						**225562**
Staphylococcus kloosii		100.0	0.41	MAN 99			
225570	**Very good identification**						**225570**
Staphylococcus kloosii		100.0	0.81				
225571	**Low discrimination**						**225571**
					YEP		
Staphylococcus kloosii		68.50	0.40	MNS 1	99		
Staphylococcus equorum		15.75	0.07		1		
Staphylococcus gallinarum		10.73	0.01		78		
225572	**Very good identification**						**225572**
Staphylococcus kloosii		100.0	0.82				
225610	**Good identification**						**225610**
Staphylococcus kloosii		100.0	0.41	SUC 9			
225612	**Good identification**						**225612**
Staphylococcus kloosii		100.0	0.41	SUC 9			
225630	**Good identification**						**225630**
Staphylococcus kloosii		100.0	0.28	SUC 9			
225632	**Good identification**						**225632**
Staphylococcus kloosii		100.0	0.29	SUC 9			
225650	**Very good identification**						**225650**
Staphylococcus kloosii		100.0	0.67	SUC 9			
225652	**Very good identification**						**225652**
Staphylococcus kloosii		98.24	0.68	SUC 9			
Staphylococcus saprophyticus subsp. saprophyticus		1.76	0.03				
225670	**Very good identification**						**225670**
Staphylococcus kloosii		100.0	0.54	SUC 9			
225671	**Low discrimination**						**225671**
Staphylococcus equorum		88.43	0.28	GLR 90			
Staphylococcus gallinarum		6.15	0.01	NIT 92			
Staphylococcus xylosus		5.42	0.01	GAL 91			
225672	**Good identification**						**225672**
Staphylococcus kloosii		93.95	0.55	SUC 9			
Staphylococcus arlettae		6.05	0.19				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
225673	**Low discrimination**							**225673**
					YEP			
Staphylococcus equorum		81.47	0.36	GLR 90	1			
Staphylococcus arlettae		6.56	0.22	NIT 92	99			
Staphylococcus xylosus		5.53	0.11	GAL 91	40			
225710	**Good identification**							**225710**
Staphylococcus kloosii		100.0	0.46	SUC 9				
225712	**Good identification**							**225712**
Staphylococcus kloosii		100.0	0.47	SUC 9				
225713	**Good identification**							**225713**
					COA			
Staphylococcus intermedius		96.26	0.32	ESL 1	99			
Staphylococcus felis		3.74	-.04	NIT 93	1			
225730	**Good identification**							**225730**
Staphylococcus kloosii		100.0	0.34	SUC 9				
225731	**Low discrimination**							**225731**
Staphylococcus equorum		89.59	0.28	GLR 90				
Staphylococcus gallinarum		6.23	0.01	NIT 92				
Staphylococcus xylosus		4.18	-.01	MLT 91				
225732	**Good identification**							**225732**
Staphylococcus kloosii		100.0	0.35	SUC 9				
225733	**Low discrimination**							**225733**
Staphylococcus equorum		89.46	0.36	GLR 90				
Staphylococcus gallinarum		5.92	0.09	NIT 92				
Staphylococcus xylosus		4.62	0.08	MLT 91				
225740	**Good identification**							**225740**
Staphylococcus kloosii		100.0	0.32	SUC 9				
225742	**Good identification**							**225742**
Staphylococcus kloosii		100.0	0.32	SUC 9				
225743	**Good identification**							**225743**
Staphylococcus intermedius		100.0	0.27	ESL 1				
225750	**Very good identification**							**225750**
Staphylococcus kloosii		99.63	0.73	SUC 9				
Staphylococcus saprophyticus subsp. bovis		0.38	0.05					
225752	**Very good identification**							**225752**
Staphylococcus kloosii		100.0	0.73	SUC 9				
225753	**Low discrimination**							**225753**
					COA	YEP	NOV	
Staphylococcus intermedius		71.85	0.36	ESL 1	99	1	1	
Staphylococcus equorum		7.94	0.16	NIT 93	1	1	99	
Staphylococcus kloosii		5.76	0.32		1	99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
225763	**Low discrimination**						**225763**
					COA	NOV	
Staphylococcus gallinarum		70.19	0.29	NIT 99	1	99	
Staphylococcus lutrae		14.02	0.12	MAN 91	99	1	
Staphylococcus equorum		11.06	0.16		1	99	
225770	**Very good identification**						**225770**
					YEP		
Staphylococcus kloosii		96.20	0.60	SUC 9	99		
Staphylococcus equorum		2.26	0.07		1		
Staphylococcus gallinarum		1.54	0.01		78		
225771	**Low discrimination**						**225771**
Staphylococcus equorum		57.25	0.48	GLR 90			
Staphylococcus gallinarum		38.99	0.42	NIT 92			
Staphylococcus xylosus		3.51	0.22				
225772	**Low discrimination**						**225772**
					YEP		
Staphylococcus kloosii		78.14	0.61	SUC 9	99		
Staphylococcus arlettae		14.68	0.35		99		
Staphylococcus equorum		4.36	0.16		1		
225773	**Low discrimination**						**225773**
Staphylococcus equorum		55.83	0.57	GLR 90			
Staphylococcus gallinarum		36.20	0.50	NIT 92			
Staphylococcus xylosus		3.79	0.31				
226000	**Low discrimination**						**226000**
					OXI		
Kocuria varians		76.02	0.46	PHS 1	13		
Dermacoccus nishinomiyaensis		23.98	0.28		93		
226002	**Good identification**						**226002**
Kocuria varians		100.0	0.34	PHS 1			
226020	**Good identification**						**226020**
Kocuria varians		100.0	0.36	PHS 1			
226040	**Good identification**						**226040**
Kocuria varians		100.0	0.32	PHS 1			
226100	**Good identification**						**226100**
					OXI		
Kocuria varians		96.15	0.49	PHS 1	13		
Dermacoccus nishinomiyaensis		3.85	0.13		93		
226101	**Low discrimination**						**226101**
					COA	PYR	
Kocuria varians		80.97	0.27	PHS 1	1	10	
Staphylococcus intermedius		13.71	0.07	MNS 8	99	90	
Staphylococcus schleiferi subsp. coagulans		5.32	0.02		99	20	
226102	**Good identification**						**226102**
Kocuria varians		100.0	0.38	PHS 1			
226113	**Low discrimination**						**226113**
					COA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus felis		52.92	0.31	ARG 92	1			
Staphylococcus intermedius		27.90	0.32	TRE 99	99			
Staphylococcus condimenti		13.66	0.14		1			
226120	**Good identification**							**226120**
Kocuria varians		100.0	0.40	PHS 1				
226122	**Good identification**							**226122**
Kocuria varians		100.0	0.28	PHS 1				
226140	**Good identification**							**226140**
Kocuria varians		97.47	0.36	PHS 1				
Staphylococcus epidermidis		2.53	0.01					
226143	**Low discrimination**							**226143**
					COA	PYR		
Staphylococcus intermedius		54.31	0.27	SUC 99	99	90		
Staphylococcus lutrae		20.58	0.12	TRE 93	99	10		
Staphylococcus epidermidis		13.40	0.17		1	1		
226150	**Good identification**							**226150**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	TRE 99				
226153	**Low discrimination**							**226153**
					COA	PYR	NOV	
Staphylococcus intermedius		75.02	0.36	SUC 99	99	90	1	
Staphylococcus lutrae		20.41	0.18	TRE 93	99	10	1	
Staphylococcus saprophyticus subsp. bovis		3.20	0.09		1	99	99	
226163	**Very good identification**							**226163**
Staphylococcus lutrae		100.0	0.53	TRE 99				
226173	**Very good identification**							**226173**
					COA	NOV		
Staphylococcus lutrae		99.81	0.59	TRE 99	99	1		
Staphylococcus xylosus		0.20	0.08		1	99		
226200	**Low discrimination**							**226200**
					OXI			
Kocuria varians		87.13	0.28	PHS 1	13			
Dermacoccus nishinomiyaensis		12.87	0.03	SUC 12	93			
226203	**Low discrimination**							**226203**
					COA	YEP		
Staphylococcus chromogenes		51.61	0.31	ARG 90	1	95		
Staphylococcus simulans		19.86	0.26	GAL 90	1	1		
Staphylococcus intermedius		17.78	0.23	TRE 92	99	1		
226212	**Good identification**							**226212**
Staphylococcus simulans		100.0	0.28	ARG 92				
226213	**Low discrimination**							**226213**
					COA			
Staphylococcus simulans		40.25	0.38	ARG 92	1			
Staphylococcus felis		28.02	0.26	TRE 95	1			
Staphylococcus intermedius		24.36	0.32		99			
226241	**Low discrimination**							**226241**
					COA	YEP	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		78.27	0.32	VPT 96	1	1	1	
Staphylococcus chromogenes		12.37	0.12		1	95	45	
Staphylococcus intermedius		9.36	0.11		99	1	90	
226242	**Low discrimination**							**226242**
Staphylococcus epidermidis		88.59	0.37	VPT 96				
Staphylococcus hominis subsp. hominis		6.01	0.16					
Staphylococcus simulans		3.81	0.08					
226243	**Low discrimination**							**226243**
					COA	YEP	PYR	
Staphylococcus epidermidis		56.86	0.42	VPT 96	1	1	1	
Staphylococcus chromogenes		25.53	0.31		1	95	45	
Staphylococcus intermedius		13.20	0.27		99	1	90	
226250	**Good identification**							**226250**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	TRE 99				
226253	**Low discrimination**							**226253**
					COA			
Staphylococcus intermedius		46.22	0.36	GAL 99	99			
Staphylococcus simulans		22.87	0.31	TRE 93	1			
Staphylococcus delphini		18.86	0.18		99			
226300	**Low discrimination**							**226300**
					COA	PYR		
Kocuria varians		86.98	0.31	PHS 1	1	10		
Staphylococcus intermedius		9.39	0.07	SUC 12	99	90		
Staphylococcus epidermidis		3.63	0.01		1	1		
226301	**Low discrimination**							**226301**
					COA	YEP	PYR	
Staphylococcus intermedius		83.33	0.48	TRE 93	99	1	90	
Staphylococcus chromogenes		15.01	0.31	LAC 86	1	95	45	
Kocuria varians		0.68	0.10		1	80	10	
226303	**Low discrimination**							**226303**
					COA	YEP		
Staphylococcus intermedius		77.27	0.64	TRE 93	99	1		
Staphylococcus chromogenes		20.39	0.51		1	95		
Staphylococcus felis		1.21	0.20		1	1		
226311	**Very good identification**							**226311**
					COA	YEP		
Staphylococcus intermedius		97.92	0.57	TRE 93	99	1		
Staphylococcus chromogenes		1.84	0.20	LAC 86	1	95		
Staphylococcus felis		0.24	-.04		1	1		
226312	**Acceptable identification**							**226312**
					COA			
Staphylococcus intermedius		90.14	0.32	TRE 93	99			
Staphylococcus simulans		3.87	0.06	MNS 99	1			
Staphylococcus felis		3.51	-.04		1			
226313	**Good identification**							**226313**
					COA	YEP		
Staphylococcus intermedius		92.96	0.73	TRE 93	99	1		
Staphylococcus felis		3.62	0.37		1	1		
Staphylococcus chromogenes		2.56	0.40		1	95		
226333	**Low discrimination**							**226333**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	PYR	NOV	
Staphylococcus intermedius		62.99	0.32	TRE 93	99	90	1	
Staphylococcus lutrae		25.71	0.18	XYL 1	99	10	1	
Staphylococcus xylosus		7.12	0.11		1	65	99	
226340	**Low discrimination**							**226340**
					COA	PYR		
Staphylococcus epidermidis		85.96	0.42	VPT 96	1	1		
Kocuria varians		4.57	0.18		1	10		
Staphylococcus intermedius		3.37	0.11		99	90		
226341	**Low discrimination**							**226341**
					COA	YEP	PYR	
Staphylococcus intermedius		62.56	0.52	TRE 93	99	1	90	
Staphylococcus epidermidis		29.93	0.48	LAC 86	1	1	1	
Staphylococcus chromogenes		7.51	0.31		1	95	45	
226342	**Acceptable identification**							**226342**
					COA	YEP	PYR	
Staphylococcus epidermidis		90.07	0.52	VPT 96	1	1	1	
Staphylococcus intermedius		6.85	0.27		99	1	90	
Staphylococcus chromogenes		1.21	0.10		1	95	45	
226343	**Low discrimination**							**226343**
					COA	YEP	PYR	
Staphylococcus intermedius		70.11	0.68	TRE 93	99	1	90	
Staphylococcus epidermidis		17.29	0.58		1	1	1	
Staphylococcus chromogenes		12.33	0.51		1	95	45	
226350	**Acceptable identification**							**226350**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		93.86	0.47	TRE 99	1	99	99	
Staphylococcus intermedius		5.60	0.20		99	1	90	
Staphylococcus epidermidis		0.53	0.01		1	1	1	
226351	**Good identification**							**226351**
					COA	NOV	YEP	
Staphylococcus intermedius		94.47	0.61	TRE 93	99	1	1	
Staphylococcus saprophyticus subsp. bovis		4.00	0.34	LAC 86	1	99	80	
Staphylococcus chromogenes		1.18	0.20		1	1	95	
226352	**Low discrimination**							**226352**
					COA	NOV	PYR	
Staphylococcus intermedius		57.17	0.36	TRE 93	99	1	90	
Staphylococcus saprophyticus subsp. bovis		38.97	0.34	MNS 99	1	99	99	
Staphylococcus epidermidis		2.81	0.11		1	1	1	
226353	**Very good identification**							**226353**
					COA	YEP		
Staphylococcus intermedius		97.14	0.77	TRE 93	99	1		
Staphylococcus chromogenes		1.78	0.40		1	95		
Staphylococcus delphini		0.40	0.18		99	1		
226363	**Very good identification**							**226363**
					COA	PYR		
Staphylococcus lutrae		95.92	0.53	TRE 99	99	10		
Staphylococcus intermedius		2.56	0.27		99	90		
Staphylococcus epidermidis		0.63	0.17		1	1		
226373	**Good identification**							**226373**
					COA	PYR	NOV	
Staphylococcus lutrae		92.80	0.59	TRE 99	99	10	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus xylosus		3.45	0.34		1	65	99	
Staphylococcus intermedius		3.45	0.36		99	90	1	
226403	**Acceptable identification**							**226403**
					YEP			
Staphylococcus felis		94.07	0.44	ARG 92	1			
Staphylococcus simulans		3.67	0.24	MAN 87	1			
Staphylococcus chromogenes		1.40	0.12		95			
226413	**Very good identification**							**226413**
Staphylococcus felis		96.05	0.61	ARG 92				
Staphylococcus simulans		2.24	0.36					
Staphylococcus condimenti		0.84	0.14					
226450	**Very good identification**							**226450**
Staphylococcus saprophyticus subsp. bovis		97.77	0.63					
Staphylococcus kloosii		2.23	0.46					
226451	**Good identification**							**226451**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		97.14	0.50		1	99		
Staphylococcus cohnii subsp. urealyticum		2.36	0.23		1	99		
Staphylococcus intermedius		0.50	0.02		99	1		
226452	**Low discrimination**							**226452**
					NOV	YEP		
Staphylococcus saprophyticus subsp. bovis		88.76	0.50		99	80		
Staphylococcus kloosii		8.78	0.47		99	99		
Staphylococcus simulans		2.46	0.18		1	1		
226453	**Low discrimination**							**226453**
					NOV			
Staphylococcus saprophyticus subsp. bovis		32.06	0.38		99			
Staphylococcus cohnii subsp. urealyticum		31.48	0.43		99			
Staphylococcus simulans		11.91	0.29		1			
226463	**Very good identification**							**226463**
Staphylococcus lutrae		100.0	0.53	GAL 99				
226473	**Very good identification**							**226473**
					COA	NOV		
Staphylococcus lutrae		99.54	0.59	GAL 99	99	1		
Staphylococcus xylosus		0.46	0.15		1	99		
226501	**Low discrimination**							**226501**
					COA	YEP		
Staphylococcus intermedius		62.94	0.30	SUC 99	99	1		
Staphylococcus felis		22.30	0.14	LAC 86	1	1		
Staphylococcus chromogenes		9.81	0.12		1	95		
226503	**Low discrimination**							**226503**
					COA	YEP		
Staphylococcus felis		80.49	0.55	ARG 92	1	1		
Staphylococcus intermedius		14.10	0.46	MAN 87	99	1		
Staphylococcus chromogenes		3.22	0.32		1	95		
226510	**Acceptable identification**							**226510**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		92.67	0.34	MLT 99	1	99		
Staphylococcus kloosii		5.35	0.26		1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		1.98	-.02		99	1		
226511	**Low discrimination**							**226511**
					COA			
Staphylococcus intermedius		44.69	0.39	SUC 99	99			
Staphylococcus felis		39.19	0.31	LAC 86	1			
Staphylococcus condimenti		10.12	0.14		1			
226512	**Low discrimination**							**226512**
					NOV			
Staphylococcus felis		65.99	0.31	ARG 92	1			
Staphylococcus condimenti		17.04	0.14	MNS 99	1			
Staphylococcus saprophyticus subsp. bovis		8.90	0.22		99			
226513	**Low discrimination**							**226513**
					COA			
Staphylococcus felis		74.94	0.72	ARG 92	1			
Staphylococcus condimenti		19.35	0.55		1			
Staphylococcus intermedius		5.30	0.55		99			
226523	**Very good identification**							**226523**
					COA	PYR		
Staphylococcus lutrae		98.43	0.53	MLT 99	99	10		
Staphylococcus felis		1.34	0.14		1	20		
Staphylococcus intermedius		0.24	0.05		99	90		
226533	**Good identification**							**226533**
					COA			
Staphylococcus lutrae		93.82	0.59	MLT 99	99			
Staphylococcus felis		4.40	0.31		1			
Staphylococcus condimenti		1.14	0.14		1			
226540	**Acceptable identification**							**226540**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		93.80	0.34	MAN 99	99	99		
Kocuria varians		6.20	0.12		1	10		
226541	**Low discrimination**							**226541**
					COA	PYR	NOV	
Staphylococcus intermedius		67.02	0.34	SUC 99	99	90	1	
Staphylococcus saprophyticus subsp. bovis		14.28	0.22	LAC 86	1	99	99	
Staphylococcus lutrae		11.74	0.12		99	10	1	
226543	**Low discrimination**							**226543**
					COA	PYR	YEP	
Staphylococcus lutrae		70.03	0.53	XYL 99	99	10	1	
Staphylococcus intermedius		24.80	0.50		99	90	1	
Staphylococcus chromogenes		3.78	0.32		1	45	95	
226550	**Very good identification**							**226550**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		98.87	0.75		1	99		
Staphylococcus kloosii		1.10	0.52		1	99		
Staphylococcus intermedius		0.03	0.02		99	1		
226551	**Low discrimination**							**226551**
					COA	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		86.91	0.63		1	99	99	
Staphylococcus intermedius		11.14	0.43		99	90	1	
Staphylococcus lutrae		1.40	0.18		99	10	1	
226552	**Good identification**							**226552**
					COA	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. bovis		93.28	0.63		1	99		
Staphylococcus kloosii		4.48	0.53		1	99		
Staphylococcus lutrae		1.50	0.18		99	1		
226553	**Low discrimination**							**226553**
					COA	PYR	NOV	
Staphylococcus lutrae		57.52	0.59	XYL 99	99	10	1	
Staphylococcus intermedius		28.37	0.59		99	90	1	
Staphylococcus saprophyticus subsp. bovis		9.01	0.50		1	99	99	
226561	**Very good identification**							**226561**
					COA	NOV		
Staphylococcus lutrae		99.80	0.53	LAC 99	99	1		
Staphylococcus xylosus		0.20	0.02		1	99		
226562	**Very good identification**							**226562**
Staphylococcus lutrae		100.0	0.53	MNS 99				
226563	**Excellent identification**							**226563**
					COA	PYR	NOV	
Staphylococcus lutrae		99.99	0.94		99	10	1	
Staphylococcus xylosus		0.01	0.11		1	65	99	
Staphylococcus intermedius		0.00	0.09		99	90	1	
226570	**Low discrimination**							**226570**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		59.76	0.34	XYL 1	1	99		
Staphylococcus lutrae		23.85	0.18		99	1		
Staphylococcus kloosii		16.39	0.40		1	99		
226571	**Very good identification**							**226571**
					COA	NOV		
Staphylococcus lutrae		97.59	0.59	LAC 99	99	1		
Staphylococcus xylosus		1.61	0.27		1	99		
Staphylococcus saprophyticus subsp. bovis		0.62	0.22		1	99		
226572	**Very good identification**							**226572**
					COA	NOV		
Staphylococcus lutrae		98.63	0.59	MNS 99	99	1		
Staphylococcus kloosii		0.74	0.41		1	99		
Staphylococcus saprophyticus subsp. bovis		0.62	0.22		1	99		
226573	**Excellent identification**							**226573**
					COA	PYR	NOV	
Staphylococcus lutrae		99.94	1.00		99	10	1	
Staphylococcus xylosus		0.05	0.36		1	65	99	
Staphylococcus intermedius		0.01	0.18		99	90	1	
226601	**Low discrimination**							**226601**
					COA	YEP		
Staphylococcus chromogenes		55.98	0.34	ARG 90	1	95		
Staphylococcus intermedius		32.65	0.30	GAL 90	99	1		
Staphylococcus felis		7.01	0.09	LAC 90	1	1		
226602	**Low discrimination**							**226602**
					YEP			
Staphylococcus simulans		87.13	0.42	ARG 92	1			
Staphylococcus felis		6.39	0.09		1			
Staphylococcus chromogenes		4.64	0.12		95			
226603	**Low discrimination**							**226603**
					YEP			

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus felis		40.09	0.50	ARG 92	1			
Staphylococcus chromogenes		29.09	0.53	MAN 87	95			
Staphylococcus simulans		18.49	0.52		1			
226610	**Low discrimination**							**226610**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		89.77	0.34	MLT 99	1	99		
Staphylococcus simulans		8.31	0.13		1	1		
Staphylococcus intermedius		1.92	-.02		99	1		
226611	**Low discrimination**							**226611**
					COA	YEP		
Staphylococcus intermedius		50.62	0.39	GAL 99	99	1		
Staphylococcus felis		26.92	0.26	LAC 86	1	1		
Staphylococcus chromogenes		9.06	0.22		1	95		
226612	**Low discrimination**							**226612**
					NOV			
Staphylococcus simulans		85.62	0.54	ARG 92	1			
Staphylococcus felis		10.50	0.26		1			
Staphylococcus saprophyticus subsp. bovis		2.33	0.22		99			
226613	**Low discrimination**							**226613**
					COA			
Staphylococcus felis		69.92	0.67	ARG 92	1			
Staphylococcus simulans		19.28	0.65		1			
Staphylococcus intermedius		8.16	0.55		99			
226640	**Low discrimination**							**226640**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		76.52	0.34	MAN 99	99	99		
Staphylococcus hominis subsp. hominis		17.64	0.25		1	1		
Staphylococcus warneri		5.84	0.21		1	8		
226641	**Low discrimination**							**226641**
					COA	NOV	YEP	
Staphylococcus chromogenes		47.91	0.34	ARG 90	1	1	95	
Staphylococcus intermedius		41.91	0.34	GAL 90	99	1	1	
Staphylococcus saprophyticus subsp. bovis		8.93	0.22	LAC 90	1	99	80	
226642	**Low discrimination**							**226642**
					NOV	PYR		
Staphylococcus simulans		58.16	0.34	ARG 92	1	70		
Staphylococcus hominis subsp. hominis		15.28	0.26		1	1		
Staphylococcus saprophyticus subsp. bovis		14.12	0.22		99	99		
226643	**Low discrimination**							**226643**
					COA	YEP		
Staphylococcus chromogenes		52.02	0.53	ARG 90	1	95		
Staphylococcus intermedius		31.06	0.50	GAL 90	99	1		
Staphylococcus simulans		14.86	0.45		1	1		
226650	**Very good identification**							**226650**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		99.78	0.75		99	99		
Staphylococcus kloosii		0.06	0.26		99	52		
Staphylococcus warneri		0.06	0.18		1	8		
226651	**Low discrimination**							**226651**
					COA	NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		86.70	0.63		1	99	80	
Staphylococcus intermedius		11.11	0.43		99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus chromogenes		1.33	0.22		1	1	95	
226652	**Low discrimination**							**226652**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		84.44	0.63		1	99		
Staphylococcus simulans		14.06	0.47		1	1		
Staphylococcus intermedius		0.67	0.18		99	1		
226653	**Low discrimination**							**226653**
					COA	NOV		
Staphylococcus intermedius		43.52	0.59	GAL 99	99	1		
Staphylococcus simulans		30.80	0.58		1	1		
Staphylococcus saprophyticus subsp. bovis		13.82	0.50		1	99		
226663	**Very good identification**							**226663**
					COA	NOV	YEP	
Staphylococcus lutrae		97.83	0.53	GAL 99	99	1	1	
Staphylococcus xylosus		1.07	0.17		1	99	40	
Staphylococcus chromogenes		0.59	0.12		1	1	95	
226670	**Good identification**							**226670**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	XYL 1				
226671	**Low discrimination**							**226671**
					COA	NOV		
Staphylococcus xylosus		60.61	0.32	GLR 95	1	99		
Staphylococcus lutrae		19.76	0.18	GAL 91	99	1		
Staphylococcus saprophyticus subsp. bovis		12.38	0.22		1	99		
226673	**Good identification**							**226673**
					COA	PYR	NOV	
Staphylococcus lutrae		90.38	0.59	GAL 99	99	10	1	
Staphylococcus xylosus		7.97	0.42		1	65	99	
Staphylococcus intermedius		0.45	0.18		99	90	1	
226700	**Low discrimination**							**226700**
					COA	YEP	PYR	
Staphylococcus intermedius		79.36	0.30	MNS 99	99	1	90	
Staphylococcus chromogenes		12.37	0.12	LAC 86	1	95	45	
Kocuria varians		4.17	0.08		1	80	10	
226701	**Low discrimination**							**226701**
					COA	YEP		
Staphylococcus intermedius		85.68	0.71	LAC 86	99	1		
Staphylococcus chromogenes		13.36	0.53		1	95		
Staphylococcus felis		0.62	0.20		1	1		
226702	**Low discrimination**							**226702**
					COA	YEP		
Staphylococcus intermedius		71.71	0.46	MNS 99	99	1		
Staphylococcus chromogenes		16.38	0.32		1	95		
Staphylococcus felis		8.40	0.20		1	1		
226703	**Low discrimination**							**226703**
					COA	YEP		
Staphylococcus intermedius		73.73	0.88		99	1		
Staphylococcus chromogenes		16.84	0.73		1	95		
Staphylococcus felis		8.63	0.61		1	1		
226710	**Low discrimination**							**226710**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		64.65	0.47	MLT 99	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		34.18	0.39		99	1		
Staphylococcus felis		0.62	-.04		1	1		
226711	**Very good identification**							**226711**
					COA	YEP		
Staphylococcus intermedius		95.88	0.80	LAC 86	99	1		
Staphylococcus felis		1.72	0.37		1	1		
Staphylococcus chromogenes		1.56	0.42		1	95		
226712	**Low discrimination**							**226712**
					COA	NOV		
Staphylococcus intermedius		66.35	0.55	MNS 99	99	1		
Staphylococcus felis		19.23	0.37		1	1		
Staphylococcus saprophyticus subsp. bovis		5.11	0.34		1	99		
226713	**Low discrimination**							**226713**
					COA			
Staphylococcus intermedius		72.90	0.96		99			
Staphylococcus felis		21.13	0.78		1			
Staphylococcus condimenti		4.03	0.59		1			
226721	**Low discrimination**							**226721**
					COA	PYR	YEP	
Staphylococcus intermedius		66.90	0.30	XYL 1	99	90	1	
Staphylococcus lutrae		17.58	0.12	LAC 86	99	10	1	
Staphylococcus chromogenes		10.43	0.12		1	45	95	
226723	**Low discrimination**							**226723**
					COA	PYR	YEP	
Staphylococcus lutrae		75.31	0.53	MLT 99	99	10	1	
Staphylococcus intermedius		17.78	0.46		99	90	1	
Staphylococcus chromogenes		4.06	0.32		1	45	95	
226731	**Low discrimination**							**226731**
					COA	PYR	NOV	
Staphylococcus intermedius		58.22	0.39	XYL 1	99	90	1	
Staphylococcus xylosus		25.65	0.30	LAC 86	1	65	99	
Staphylococcus lutrae		10.98	0.18		99	10	1	
226733	**Low discrimination**							**226733**
					COA	PYR		
Staphylococcus lutrae		65.56	0.59	MLT 99	99	10		
Staphylococcus intermedius		21.56	0.55		99	90		
Staphylococcus felis		6.25	0.37		1	20		
226740	**Low discrimination**							**226740**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		71.53	0.47	MAN 99	1	99	99	
Staphylococcus intermedius		20.98	0.34		99	1	90	
Staphylococcus hominis subsp. hominis		4.47	0.26		1	1	1	
226741	**Low discrimination**							**226741**
					COA	NOV	YEP	
Staphylococcus intermedius		89.67	0.75	LAC 86	99	1	1	
Staphylococcus chromogenes		9.32	0.53		1	1	95	
Staphylococcus saprophyticus subsp. bovis		0.77	0.34		1	99	80	
226742	**Low discrimination**							**226742**
					COA	NOV	YEP	
Staphylococcus intermedius		72.33	0.50	MNS 99	99	1	1	
Staphylococcus chromogenes		11.01	0.32		1	1	95	
Staphylococcus saprophyticus subsp. bovis		10.03	0.34		1	99	80	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
226743	**Low discrimination**							**226743**
					COA	PYR	YEP	
Staphylococcus intermedius		84.54	0.91		99	90	1	
Staphylococcus chromogenes		12.87	0.73		1	45	95	
Staphylococcus lutrae		2.41	0.53		99	10	1	
226750	**Very good identification**							**226750**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		99.16	0.88		1	99		
Staphylococcus intermedius		0.79	0.43		99	1		
Staphylococcus kloosii		0.03	0.32		1	99		
226751	**Low discrimination**							**226751**
					COA	NOV	YEP	
Staphylococcus intermedius		75.13	0.84	LAC 86	99	1	1	
Staphylococcus saprophyticus subsp. bovis		23.68	0.75		1	99	80	
Staphylococcus chromogenes		0.82	0.42		1	1	95	
226752	**Low discrimination**							**226752**
					COA	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		82.65	0.75		1	99	99	
Staphylococcus intermedius		16.27	0.59		99	90	1	
Staphylococcus lutrae		0.33	0.18		99	10	1	
226753	**Good identification**							**226753**
					COA	PYR	YEP	
Staphylococcus intermedius		94.82	1.00		99	90	1	
Staphylococcus lutrae		1.94	0.59		99	10	1	
Staphylococcus chromogenes		1.51	0.62		1	45	95	
226761	**Low discrimination**							**226761**
					COA	PYR	NOV	
Staphylococcus lutrae		89.04	0.53	LAC 99	99	10	1	
Staphylococcus intermedius		5.13	0.34		99	90	1	
Staphylococcus xylosus		3.46	0.28		1	65	99	
226762	**Very good identification**							**226762**
					COA	PYR	NOV	
Staphylococcus lutrae		99.53	0.53	MNS 99	99	10	1	
Staphylococcus intermedius		0.36	0.09		99	90	1	
Staphylococcus xylosus		0.11	-.03		1	65	99	
226763	**Excellent identification**							**226763**
					COA	PYR	NOV	
Staphylococcus lutrae		99.43	0.94		99	10	1	
Staphylococcus intermedius		0.36	0.50		99	90	1	
Staphylococcus xylosus		0.11	0.38		1	65	99	
226770	**Low discrimination**							**226770**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		86.46	0.47	XYL 1	1	99		
Staphylococcus lutrae		8.63	0.18		99	1		
Staphylococcus xylosus		2.70	0.12		1	99		
226771	**Low discrimination**							**226771**
					COA	NOV		
Staphylococcus lutrae		64.69	0.59	LAC 99	99	1		
Staphylococcus xylosus		20.27	0.53		1	99		
Staphylococcus gallinarum		6.96	0.42		1	99		
226772	**Very good identification**							**226772**
					COA	NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lutrae		95.90	0.59	MNS 99	99	1		
Staphylococcus saprophyticus subsp. bovis		2.43	0.34		1	99		
Staphylococcus xylosus		0.86	0.21		1	99		
226773	**Very good identification**							**226773**
					COA	PYR	NOV	
Staphylococcus lutrae		98.30	1.00		99	10	1	
Staphylococcus xylosus		0.89	0.62		1	65	99	
Staphylococcus intermedius		0.49	0.59		99	90	1	
227313	**Acceptable identification**							**227313**
					COA			
Staphylococcus intermedius		90.36	0.32	ESL 1	99			
Staphylococcus schleiferi subsp. coagulans		3.63	0.07	TRE 93	99			
Staphylococcus felis		3.52	-.04		1			
227343	**Low discrimination**							**227343**
					COA	YEP	PYR	
Staphylococcus intermedius		70.30	0.27	ESL 1	99	1	90	
Staphylococcus epidermidis		17.34	0.17	TRE 93	1	1	1	
Staphylococcus chromogenes		12.37	0.10		1	95	45	
227353	**Low discrimination**							**227353**
					COA	NOV		
Staphylococcus intermedius		88.29	0.36	ESL 1	99	1		
Staphylococcus gallinarum		6.33	0.09	TRE 93	1	99		
Staphylococcus xylosus		1.92	0.00		1	99		
227363	**Low discrimination**							**227363**
					COA	NOV		
Staphylococcus gallinarum		81.80	0.29	TRE 99	1	99		
Staphylococcus lutrae		16.34	0.12	MAN 91	99	1		
Staphylococcus xylosus		1.86	-.03		1	99		
227371	**Low discrimination**							**227371**
Staphylococcus gallinarum		83.27	0.42	TRE 99				
Staphylococcus equorum		14.20	0.29					
Staphylococcus xylosus		2.53	0.12					
227373	**Low discrimination**							**227373**
					COA	NOV		
Staphylococcus gallinarum		79.64	0.50	TRE 99	1	99		
Staphylococcus equorum		14.27	0.38		1	99		
Staphylococcus lutrae		3.05	0.18		99	1		
227450	**Low discrimination**							**227450**
Staphylococcus kloosii		69.33	0.46	NIT 1				
Staphylococcus saprophyticus subsp. bovis		30.67	0.22					
227452	**Acceptable identification**							**227452**
Staphylococcus kloosii		90.74	0.47	NIT 1				
Staphylococcus saprophyticus subsp. bovis		9.26	0.09					
227453	**Good identification**							**227453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	GLR 99				
227470	**Good identification**							**227470**
Staphylococcus kloosii		100.0	0.34	NIT 1				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
227472	**Good identification**						**227472**
	Staphylococcus kloosii	100.0	0.35	NIT 1			
227512	**Good identification**						**227512**
	Staphylococcus kloosii	100.0	0.27	NIT 1			
227513	**Low discrimination**						**227513**
					COA		
	Staphylococcus felis	73.91	0.31	ARG 92	1		
	Staphylococcus condimenti	19.08	0.14	ESL 1	1		
	Staphylococcus intermedius	5.23	0.14		99		
227550	**Low discrimination**						**227550**
	Staphylococcus kloosii	52.31	0.52	NIT 1			
	Staphylococcus saprophyticus subsp. bovis	47.69	0.34				
227552	**Low discrimination**						**227552**
	Staphylococcus kloosii	82.62	0.53	NIT 1			
	Staphylococcus saprophyticus subsp. bovis	17.38	0.22				
227563	**Very good identification**						**227563**
					COA	NOV	
	Staphylococcus lutrae	95.06	0.53	ESL 1	99	1	
	Staphylococcus gallinarum	4.81	0.29		1	99	
	Staphylococcus xylosus	0.14	-.01		1	99	
227570	**Low discrimination**						**227570**
	Staphylococcus kloosii	86.46	0.40	NIT 1			
	Staphylococcus gallinarum	13.54	0.01				
227571	**Low discrimination**						**227571**
					COA	NOV	
	Staphylococcus gallinarum	76.76	0.42	SUC 99	1	99	
	Staphylococcus equorum	13.09	0.29		1	99	
	Staphylococcus lutrae	7.21	0.18		99	1	
227573	**Low discrimination**						**227573**
					COA	NOV	
	Staphylococcus lutrae	75.58	0.59	ESL 1	99	1	
	Staphylococcus gallinarum	19.91	0.50		1	99	
	Staphylococcus equorum	3.57	0.38		1	99	
227631	**Acceptable identification**						**227631**
	Staphylococcus equorum	91.41	0.29	GLR 90			
	Staphylococcus gallinarum	5.59	0.01	GAL 91			
	Staphylococcus xylosus	3.00	-.03	MLT 91			
227633	**Acceptable identification**						**227633**
	Staphylococcus equorum	91.36	0.37	GLR 90			
	Staphylococcus gallinarum	5.32	0.09	GAL 91			
	Staphylococcus xylosus	3.32	0.06	MLT 91			
227650	**Acceptable identification**						**227650**
	Staphylococcus saprophyticus subsp. bovis	94.71	0.34	ESL 1			
	Staphylococcus kloosii	5.29	0.26				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
227663	**Low discrimination**							**227663**
					COA	NOV		
Staphylococcus gallinarum		69.79	0.29	GAL 99	1	99		
Staphylococcus lutrae		13.94	0.12	MAN 91	99	1		
Staphylococcus equorum		12.50	0.17		1	99		
227671	**Low discrimination**							**227671**
Staphylococcus equorum		60.61	0.49	GLR 90				
Staphylococcus gallinarum		36.29	0.42	GAL 91				
Staphylococcus xylosus		2.61	0.20					
227673	**Low discrimination**							**227673**
Staphylococcus equorum		60.93	0.58	GLR 90				
Staphylococcus gallinarum		34.73	0.50	GAL 91				
Staphylococcus xylosus		2.91	0.29					
227701	**Low discrimination**							**227701**
					COA	YEP		
Staphylococcus intermedius		86.51	0.30	ESL 1	99	1		
Staphylococcus chromogenes		13.49	0.12	LAC 86	1	95		
227703	**Low discrimination**							**227703**
					COA	YEP		
Staphylococcus intermedius		73.91	0.46	ESL 1	99	1		
Staphylococcus chromogenes		16.88	0.32		1	95		
Staphylococcus felis		8.65	0.20		1	1		
227711	**Low discrimination**							**227711**
					OXI	COA	NOV	
Staphylococcus intermedius		76.01	0.39	ESL 1	1	99	1	
Staphylococcus lentus		15.12	0.29	LAC 86	99	1	99	
Staphylococcus equorum		2.58	0.08		1	1	99	
227713	**Low discrimination**							**227713**
					OXI	COA	NOV	
Staphylococcus intermedius		66.11	0.55	ESL 1	1	99	1	
Staphylococcus felis		19.16	0.37		1	1	1	
Staphylococcus lentus		7.59	0.40		99	1	99	
227723	**Low discrimination**							**227723**
					COA	NOV		
Staphylococcus gallinarum		68.16	0.29	MAN 91	1	99		
Staphylococcus lutrae		13.61	0.12	MLT 99	99	1		
Staphylococcus equorum		12.21	0.17		1	99		
227731	**Low discrimination**							**227731**
					OXI			
Staphylococcus equorum		58.89	0.49	GLR 90	1			
Staphylococcus gallinarum		35.26	0.42	MLT 91	1			
Staphylococcus lentus		3.49	0.29		99			
227733	**Low discrimination**							**227733**
					OXI			
Staphylococcus equorum		58.17	0.58	GLR 90	1			
Staphylococcus gallinarum		33.16	0.50	MLT 91	1			
Staphylococcus lentus		4.75	0.40		99			
227741	**Low discrimination**							**227741**
					COA	NOV	YEP	
Staphylococcus intermedius		68.39	0.34	ESL 1	99	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus gallinarum		24.50	0.21	LAC 86	1	99	78	
Staphylococcus chromogenes		7.11	0.12		1	1	95	
227743	**Low discrimination**							**227743**
					COA	NOV	YEP	
Staphylococcus intermedius		74.93	0.50	ESL 1	99	1	1	
Staphylococcus chromogenes		11.41	0.32		1	1	95	
Staphylococcus gallinarum		10.70	0.29		1	99	78	
227750	**Good identification**							**227750**
					PYR			
Staphylococcus saprophyticus subsp. bovis		95.62	0.47	ESL 1	99			
Staphylococcus kloosii		2.59	0.32		52			
Staphylococcus gallinarum		1.03	0.01		12			
227751	**Low discrimination**							**227751**
					COA	PYR	NOV	
Staphylococcus gallinarum		40.66	0.42	XYL 99	1	12	99	
Staphylococcus intermedius		30.35	0.43		99	90	1	
Staphylococcus saprophyticus subsp. bovis		9.57	0.34		1	99	99	
227752	**Low discrimination**							**227752**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		68.96	0.34	ESL 1	1	99		
Staphylococcus intermedius		13.58	0.18		99	1		
Staphylococcus kloosii		8.10	0.32		1	99		
227753	**Low discrimination**							**227753**
					OXI	COA	NOV	
Staphylococcus intermedius		53.28	0.59	ESL 1	1	99	1	
Staphylococcus gallinarum		28.45	0.50		1	1	99	
Staphylococcus lentus		8.28	0.47		99	1	99	
227761	**Very good identification**							**227761**
					COA	NOV		
Staphylococcus gallinarum		97.36	0.63	MAN 91	1	99		
Staphylococcus equorum		1.70	0.29		1	99		
Staphylococcus lutrae		0.48	0.12		99	1		
227762	**Low discrimination**							**227762**
					COA	NOV		
Staphylococcus gallinarum		83.35	0.29	MAN 91	1	99		
Staphylococcus lutrae		16.65	0.12	MNS 99	99	1		
227763	**Low discrimination**							**227763**
					COA	NOV		
Staphylococcus gallinarum		81.66	0.71	MAN 91	1	99		
Staphylococcus lutrae		16.31	0.53		99	1		
Staphylococcus equorum		1.49	0.38		1	99		
227770	**Low discrimination**							**227770**
					PYR			
Staphylococcus gallinarum		83.77	0.42	MNS 99	12			
Staphylococcus equorum		14.29	0.29		5			
Staphylococcus saprophyticus subsp. bovis		0.80	0.05		99			
227771	**Low discrimination**							**227771**
Staphylococcus gallinarum		84.66	0.83					
Staphylococcus equorum		14.44	0.70					
Staphylococcus xylosus		0.62	0.41					
227772	**Low discrimination**							**227772**

Code / Taxon		ID score	T index	tests against		Differentiation tests		Code
						COA	NOV	
Staphylococcus gallinarum		81.18	0.50	MNS	99	1	99	
Staphylococcus equorum		14.54	0.38			1	99	
Staphylococcus lutrae		3.11	0.18			99	1	
227773	**Low discrimination**							**227773**
						COA	NOV	
Staphylococcus gallinarum		81.36	0.91			1	99	
Staphylococcus equorum		14.58	0.79			1	99	
Staphylococcus lutrae		3.12	0.59			99	1	
230000	**Low discrimination**							**230000**
						OXI		
Kocuria varians		55.33	0.31	ORN	1	13		
Dermacoccus nishinomiyaensis		29.08	0.18			93		
Micrococcus sp.		15.59	0.03			89		
230100	**Acceptable identification**							**230100**
						OXI		
Kocuria varians		93.74	0.35	ORN	1	13		
Dermacoccus nishinomiyaensis		6.26	0.03			93		
230640	**Low discrimination**							**230640**
						NOV		
Staphylococcus warneri		69.46	0.28	ORN	1	1		
Staphylococcus saprophyticus subsp. saprophyticus		30.54	0.00			99		
230650	**Acceptable identification**							**230650**
						NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		91.73	0.29	ORN	1	99	7	
Staphylococcus warneri		6.30	0.25			1	8	
Staphylococcus saprophyticus subsp. bovis		1.97	0.05			99	99	
230652	**Good identification**							**230652**
						NOV		
Staphylococcus saprophyticus subsp. saprophyticus		99.60	0.44	ORN	1	99		
Staphylococcus warneri		0.40	0.16			1		
232000	**Acceptable identification**							**232000**
						OXI		
Kocuria varians		94.29	0.46	ORN	1	13		
Dermacoccus nishinomiyaensis		5.71	0.13			93		
232002	**Good identification**							**232002**
Kocuria varians		100.0	0.34	ORN	1			
232020	**Good identification**							**232020**
Kocuria varians		100.0	0.36	ORN	1			
232040	**Good identification**							**232040**
Kocuria varians		100.0	0.32	ORN	1			
232100	**Good identification**							**232100**
						OXI		
Kocuria varians		99.24	0.49	ORN	1	13		
Dermacoccus nishinomiyaensis		0.76	-.02			93		
232101	**Good identification**							**232101**
Kocuria varians		100.0	0.27	ORN	1			
232102	**Good identification**							**232102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		100.0	0.38	ORN 1			
232120	**Good identification**						**232120**
Kocuria varians		100.0	0.40	ORN 1			
232122	**Good identification**						**232122**
Kocuria varians		100.0	0.28	ORN 1			
232140	**Good identification**						**232140**
Kocuria varians		100.0	0.36	ORN 1			
232200	**Good identification**						**232200**
Kocuria varians		100.0	0.28	ORN 1			
232300	**Good identification**						**232300**
Kocuria varians		100.0	0.31	ORN 1			
232450	**Good identification**						**232450**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
232550	**Good identification**						**232550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.47	ORN 1			
232551	**Good identification**						**232551**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
232552	**Good identification**						**232552**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
232613	**Good identification**						**232613**
Staphylococcus simulans		100.0	0.34	ARG 92			
232641	**Good identification**						**232641**
Staphylococcus lugdunensis		100.0	0.27	VPT 99			
232643	**Good identification**						**232643**
Staphylococcus lugdunensis		98.36	0.47	VPT 99			
Staphylococcus simulans		1.64	0.14				
232650	**Good identification**						**232650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.64	0.47	ORN 1	99	99	
Staphylococcus warneri		1.36	0.18		1	8	
232651	**Good identification**						**232651**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
232652	**Low discrimination**						**232652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		81.14	0.34	ORN 1	99	99	
Staphylococcus simulans		10.13	0.16		1	70	
Staphylococcus saprophyticus subsp. saprophyticus		8.73	0.03		99	7	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
232741	**Acceptable identification**						**232741**
					COA	NOV	
Staphylococcus lugdunensis		90.08	0.27	VPT 99	1	1	
Staphylococcus saprophyticus subsp. bovis		5.23	0.05	LAC 90	1	99	
Staphylococcus intermedius		4.70	0.03		99	1	
232743	**Good identification**						**232743**
					COA		
Staphylococcus lugdunensis		96.56	0.47	VPT 99	1		
Staphylococcus intermedius		3.44	0.19		99		
232750	**Very good identification**						**232750**
Staphylococcus saprophyticus subsp. bovis		100.0	0.59	ORN 1			
232751	**Good identification**						**232751**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		96.70	0.47	ORN 1	1	99	
Staphylococcus intermedius		2.37	0.12		99	1	
Staphylococcus xylosus		0.93	0.01		1	99	
232752	**Good identification**						**232752**
Staphylococcus saprophyticus subsp. bovis		100.0	0.47	ORN 1			
232753	**Low discrimination**						**232753**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		55.19	0.34	ORN 1	1	99	
Staphylococcus intermedius		33.26	0.28		99	1	
Staphylococcus xylosus		6.06	0.11		1	99	
232773	**Low discrimination**						**232773**
					COA	NOV	
Staphylococcus xylosus		75.97	0.33	ORN 1	1	99	
Staphylococcus lutrae		24.03	0.18	GLR 95	99	1	
233773	**Low discrimination**						**233773**
					OXI		
Staphylococcus equorum		69.60	0.34	ORN 1	1		
Staphylococcus xylosus		17.66	0.20	GLR 90	1		
Staphylococcus lentus		6.85	0.18		99		
234450	**Good identification**						**234450**
Staphylococcus kloosii		100.0	0.46	ORN 1			
234452	**Good identification**						**234452**
Staphylococcus kloosii		100.0	0.47	ORN 1			
234470	**Good identification**						**234470**
Staphylococcus kloosii		100.0	0.34	ORN 1			
234472	**Good identification**						**234472**
Staphylococcus kloosii		100.0	0.35	ORN 1			
234512	**Good identification**						**234512**
Staphylococcus kloosii		100.0	0.27	ORN 1			
234550	**Good identification**						**234550**
Staphylococcus kloosii		100.0	0.52	ORN 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
234552	**Very good identification**					**234552**
Staphylococcus kloosii		100.0	0.53	ORN 1		
234570	**Good identification**					**234570**
Staphylococcus kloosii		100.0	0.40	ORN 1		
234572	**Good identification**					**234572**
Staphylococcus kloosii		100.0	0.41	ORN 1		
234713	**Good identification**					**234713**
					COA	
Staphylococcus intermedius		96.26	0.32	ORN 1	99	
Staphylococcus felis		3.74	-.04	NIT 93	1	
234743	**Good identification**					**234743**
Staphylococcus intermedius		100.0	0.27	ORN 1		
234750	**Low discrimination**					**234750**
Staphylococcus kloosii		72.86	0.32	ORN 1		
Staphylococcus saprophyticus subsp. bovis		27.14	0.05	SUC 9		
234752	**Good identification**					**234752**
Staphylococcus kloosii		100.0	0.32	ORN 1		
234753	**Good identification**					**234753**
Staphylococcus intermedius		100.0	0.36	ORN 1		
235450	**Good identification**					**235450**
Staphylococcus kloosii		100.0	0.46	ORN 1		
235452	**Good identification**					**235452**
Staphylococcus kloosii		100.0	0.47	ORN 1		
235470	**Good identification**					**235470**
Staphylococcus kloosii		100.0	0.34	ORN 1		
235472	**Good identification**					**235472**
Staphylococcus kloosii		100.0	0.35	ORN 1		
235512	**Good identification**					**235512**
Staphylococcus kloosii		100.0	0.27	ORN 1		
235550	**Good identification**					**235550**
Staphylococcus kloosii		100.0	0.52	ORN 1		
235552	**Very good identification**					**235552**
Staphylococcus kloosii		100.0	0.53	ORN 1		
235570	**Good identification**					**235570**
Staphylococcus kloosii		100.0	0.40	ORN 1		
235572	**Good identification**					**235572**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.41	ORN 1				
235750	**Good identification**							**235750**
Staphylococcus kloosii		100.0	0.32	ORN 1				
235752	**Good identification**							**235752**
Staphylococcus kloosii		100.0	0.32	ORN 1				
236313	**Acceptable identification**							**236313**
					COA	YEP		
Staphylococcus intermedius		93.77	0.32	ORN 1	99	1		
Staphylococcus felis		3.65	-.04	TRE 93	1	1		
Staphylococcus chromogenes		2.58	-.01		1	95		
236341	**Low discrimination**							**236341**
					COA	PYR		
Staphylococcus epidermidis		86.59	0.30	VPT 96	1	1		
Staphylococcus intermedius		13.41	0.11	ORN 12	99	90		
236342	**Good identification**							**236342**
Staphylococcus epidermidis		100.0	0.34	VPT 96				
236343	**Low discrimination**							**236343**
					COA	YEP	PYR	
Staphylococcus epidermidis		73.90	0.40	VPT 96	1	1	1	
Staphylococcus intermedius		22.19	0.27	ORN 12	99	1	90	
Staphylococcus chromogenes		3.90	0.10		1	95	45	
236353	**Good identification**							**236353**
					COA	YEP		
Staphylococcus intermedius		98.20	0.36	ORN 1	99	1		
Staphylococcus chromogenes		1.80	-.01	TRE 93	1	95		
236513	**Low discrimination**							**236513**
					COA			
Staphylococcus felis		75.25	0.31	ARG 92	1			
Staphylococcus condimenti		19.43	0.14	ORN 1	1			
Staphylococcus intermedius		5.32	0.14		99			
236550	**Good identification**							**236550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1				
236563	**Very good identification**							**236563**
Staphylococcus lutrae		100.0	0.53	ORN 1				
236573	**Very good identification**							**236573**
Staphylococcus lutrae		100.0	0.59	ORN 1				
236650	**Good identification**							**236650**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1				
236701	**Low discrimination**							**236701**
					COA	YEP		
Staphylococcus intermedius		86.51	0.30	ORN 1	99	1		
Staphylococcus chromogenes		13.49	0.12	LAC 86	1	95		
236703	**Low discrimination**							**236703**
					COA	YEP		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		73.91	0.46	ORN 1	99	1		
Staphylococcus chromogenes		16.88	0.32		1	95		
Staphylococcus felis		8.65	0.20		1	1		
236711	**Good identification**							**236711**
					COA	YEP		
Staphylococcus intermedius		96.69	0.39	ORN 1	99	1		
Staphylococcus felis		1.74	-.04	LAC 86	1	1		
Staphylococcus chromogenes		1.57	0.01		1	95		
236713	**Low discrimination**							**236713**
					COA			
Staphylococcus intermedius		72.94	0.55	ORN 1	99			
Staphylococcus felis		21.14	0.37		1			
Staphylococcus condimenti		4.03	0.18		1			
236741	**Acceptable identification**							**236741**
					COA	YEP		
Staphylococcus intermedius		90.59	0.34	ORN 1	99	1		
Staphylococcus chromogenes		9.41	0.12	LAC 86	1	95		
236743	**Low discrimination**							**236743**
					COA	PYR	YEP	
Staphylococcus intermedius		84.06	0.50	ORN 1	99	90	1	
Staphylococcus chromogenes		12.80	0.32		1	45	95	
Staphylococcus lutrae		2.40	0.12		99	10	1	
236750	**Good identification**							**236750**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		99.21	0.47	ORN 1	1	99		
Staphylococcus intermedius		0.80	0.02		99	1		
236751	**Low discrimination**							**236751**
					COA	NOV	YEP	
Staphylococcus intermedius		75.41	0.43	ORN 1	99	1	1	
Staphylococcus saprophyticus subsp. bovis		23.77	0.34	LAC 86	1	99	80	
Staphylococcus chromogenes		0.82	0.01		1	1	95	
236752	**Low discrimination**							**236752**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		83.55	0.34	ORN 1	1	99		
Staphylococcus intermedius		16.45	0.18		99	1		
236753	**Good identification**							**236753**
					COA	PYR	YEP	
Staphylococcus intermedius		94.99	0.59	ORN 1	99	90	1	
Staphylococcus lutrae		1.95	0.18		99	10	1	
Staphylococcus chromogenes		1.51	0.20		1	45	95	
236763	**Very good identification**							**236763**
					COA	PYR	NOV	
Staphylococcus lutrae		99.53	0.53	ORN 1	99	10	1	
Staphylococcus intermedius		0.36	0.09		99	90	1	
Staphylococcus xylosus		0.11	-.03		1	65	99	
236773	**Very good identification**							**236773**
					COA	PYR	NOV	
Staphylococcus lutrae		98.36	0.59	ORN 1	99	10	1	
Staphylococcus xylosus		0.89	0.21		1	65	99	
Staphylococcus intermedius		0.49	0.18		99	90	1	
237763	**Low discrimination**							**237763**
					COA	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus gallinarum		83.35	0.29	ORN 1	1	99	
Staphylococcus lutrae		16.65	0.12	MAN 91	99	1	
237771	**Low discrimination**						**237771**
Staphylococcus gallinarum		84.90	0.42	ORN 1			
Staphylococcus equorum		14.48	0.29				
Staphylococcus xylosus		0.62	-.01				
237773	**Low discrimination**						**237773**
					COA	NOV	
Staphylococcus gallinarum		81.37	0.50	ORN 1	1	99	
Staphylococcus equorum		14.58	0.38		1	99	
Staphylococcus lutrae		3.12	0.18		99	1	
240000	**Low discrimination**						**240000**
Dermacoccus nishinomiyaensis		60.11	0.59	GLR 1			
Micrococcus sp.		32.23	0.44				
Kocuria rosea		3.79	0.33				
240002	**Low discrimination**						**240002**
					OXI		
Kocuria varians		51.17	0.30	GLR 1	13		
Dermacoccus nishinomiyaensis		31.78	0.18		93		
Micrococcus sp.		17.04	0.03		89		
240010	**Low discrimination**						**240010**
					YEP		
Macrococcus equipersicus		58.11	0.43	GLR 1	99		
Macrococcus bovicus		28.14	0.35		99		
Dermacoccus nishinomiyaensis		5.89	0.18		1		
240013	**Good identification**						**240013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	TRE 99			
240042	**Low discrimination**						**240042**
Staphylococcus warneri		71.98	0.32	SUC 95			
Kocuria varians		28.03	0.16	TRE 95			
240043	**Good identification**						**240043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99			
240050	**Low discrimination**						**240050**
					OXI	NOV	
Macrococcus equipersicus		67.03	0.52	GLR 1	99	99	
Macrococcus bovicus		28.01	0.43		99	99	
Staphylococcus warneri		4.96	0.39		1	1	
240051	**Low discrimination**						**240051**
					OXI		
Staphylococcus cohnii subsp. urealyticum		77.54	0.31	TRE 99	1		
Macrococcus equipersicus		15.84	0.11	LAC 91	99		
Macrococcus bovicus		6.62	0.02		99		
240052	**Low discrimination**						**240052**
					OXI	NOV	
Staphylococcus warneri		56.00	0.29	SUC 95	1	1	
Macrococcus equipersicus		22.94	0.11	TRE 95	99	99	
Staphylococcus cohnii subsp. urealyticum		11.47	0.11		1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
240053	**Good identification**						**240053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	TRE 99			
240100	**Low discrimination**						**240100**
					OXI		
Dermacoccus nishinomiyaensis		66.92	0.44	GLR 1	93		
Kocuria varians		30.39	0.45		13		
Micrococcus sp.		1.90	0.03		89		
240102	**Acceptable identification**						**240102**
					OXI		
Kocuria varians		92.69	0.33	GLR 1	13		
Dermacoccus nishinomiyaensis		7.31	0.03		93		
240110	**Low discrimination**						**240110**
					YEP		
Macrococcus equipersicus		64.53	0.36	GLR 1	99		
Macrococcus bovicus		31.25	0.28		99		
Dermacoccus nishinomiyaensis		2.91	0.03		1		
240113	**Good identification**						**240113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.29	TRE 99			
240120	**Acceptable identification**						**240120**
					OXI		
Kocuria varians		94.05	0.35	GLR 1	13		
Dermacoccus nishinomiyaensis		5.95	0.03		93		
240140	**Low discrimination**						**240140**
					NOV	OXI	
Kocuria varians		53.23	0.31	GLR 1	1	13	
Staphylococcus warneri		24.35	0.31		1	1	
Macrococcus equipersicus		17.03	0.18		99	99	
240150	**Low discrimination**						**240150**
					OXI	NOV	
Macrococcus equipersicus		67.92	0.45	GLR 1	99	99	
Macrococcus bovicus		28.38	0.35		99	99	
Staphylococcus warneri		3.70	0.28		1	1	
240153	**Good identification**						**240153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	TRE 99			
240200	**Low discrimination**						**240200**
					OXI		
Dermacoccus nishinomiyaensis		39.91	0.34	GLR 1	93		
Micrococcus sp.		33.16	0.23	SUC 6	89		
Staphylococcus warneri		17.20	0.42		1		
240202	**Low discrimination**						**240202**
Staphylococcus warneri		80.52	0.32	TRE 95			
Kocuria varians		19.48	0.12	MLT 95			
240210	**Low discrimination**						**240210**
					OXI	NOV	
Staphylococcus warneri		42.37	0.39	TRE 95	1	1	
Macrococcus bovicus		25.64	0.23	MLT 95	99	99	
Macrococcus equipersicus		20.95	0.23		99	99	
240212	**Low discrimination**						**240212**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		78.65	0.29	TRE 95			
Staphylococcus pasteuri		21.36	0.03	MLT 95			
240240	**Very good identification**						**240240**
					NOV	YEP	
Staphylococcus warneri		96.74	0.68	TRE 95	1	43	
Staphylococcus hominis subsp. novobiosepticus		0.97	0.10		99	1	
Staphylococcus pasteuri		0.82	0.11		1	86	
240241	**Low discrimination**						**240241**
					NOV		
Staphylococcus warneri		89.05	0.33	TRE 95	1		
Staphylococcus hominis subsp. novobiosepticus		10.95	-.03	MNS 2	99		
240242	**Very good identification**						**240242**
					NOV	YEP	
Staphylococcus warneri		96.88	0.58	TRE 95	1	43	
Staphylococcus hominis subsp. novobiosepticus		1.57	0.04		99	1	
Staphylococcus pasteuri		1.00	0.03		1	86	
240250	**Low discrimination**						**240250**
					OXI	NOV	
Staphylococcus warneri		82.21	0.65	TRE 95	1	1	
Macrococcus bovicus		6.11	0.31		99	99	
Staphylococcus pasteuri		5.89	0.27		1	1	
240251	**Good identification**						**240251**
Staphylococcus warneri		100.0	0.30	TRE 95			
240252	**Good identification**						**240252**
					NOV		
Staphylococcus warneri		91.03	0.55	TRE 95	1		
Staphylococcus pasteuri		7.99	0.19		1		
Staphylococcus saprophyticus subsp. saprophyticus		0.97	-.06		99		
240260	**Good identification**						**240260**
Staphylococcus warneri		100.0	0.27	TRE 95			
240310	**Low discrimination**						**240310**
					OXI	NOV	
Staphylococcus warneri		40.12	0.28	TRE 95	1	1	
Macrococcus bovicus		32.96	0.15	MLT 95	99	99	
Macrococcus equipersicus		26.93	0.15		99	99	
240340	**Low discrimination**						**240340**
					NOV		
Staphylococcus warneri		89.56	0.58	TRE 95	1		
Staphylococcus hominis subsp. novobiosepticus		8.54	0.20		99		
Kocuria varians		1.41	0.13		1		
240342	**Low discrimination**						**240342**
					NOV		
Staphylococcus warneri		84.27	0.48	TRE 95	1		
Staphylococcus hominis subsp. novobiosepticus		12.98	0.14		99		
Staphylococcus hominis subsp. hominis		1.63	0.07		1		
240350	**Low discrimination**						**240350**
					OXI	NOV	
Staphylococcus warneri		83.04	0.55	TRE 95	1	1	
Macrococcus bovicus		8.38	0.23		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		7.93	0.24		99	99	
240352	**Good identification**						**240352**
Staphylococcus warneri		100.0	0.45	TRE 95			
240371	**Good identification**						**240371**
Staphylococcus xylosus		100.0	0.31	NIT 89			
240373	**Good identification**						**240373**
Staphylococcus xylosus		98.20	0.41	NIT 89			
Staphylococcus arlettae		1.80	0.14	TRE 96			
240400	**Low discrimination**						**240400**
					OXI	NOV	
Staphylococcus warneri		40.20	0.42	SUC 95	1	1	
Staphylococcus cohnii subsp. cohnii		21.00	0.31	MLT 95	1	99	
Dermacoccus nishinomiyaensis		14.76	0.18		93	80	
240401	**Low discrimination**						**240401**
Staphylococcus cohnii subsp. urealyticum		59.63	0.36	MAN 91			
Staphylococcus cohnii subsp. cohnii		40.37	0.32	LAC 91			
240402	**Low discrimination**						**240402**
					NOV		
Staphylococcus warneri		71.96	0.32	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		18.44	0.16	MLT 95	99		
Kocuria varians		9.61	0.07		1		
240403	**Very good identification**						**240403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.57	MAN 91			
240410	**Low discrimination**						**240410**
					OXI	NOV	
Macrococcus equipersicus		45.69	0.43	GLR 1	99	99	
Macrococcus bovicus		33.18	0.39		99	99	
Staphylococcus warneri		9.14	0.39		1	1	
240411	**Good identification**						**240411**
					OXI	YEP	
Staphylococcus cohnii subsp. urealyticum		91.52	0.57	LAC 91	1	64	
Staphylococcus cohnii subsp. cohnii		7.80	0.34		1	12	
Macrococcus equipersicus		0.40	0.02		99	99	
240412	**Low discrimination**						**240412**
					NOV		
Staphylococcus cohnii subsp. urealyticum		67.65	0.36	MNS 99	99		
Staphylococcus warneri		18.91	0.29		1		
Staphylococcus pasteuri		5.13	0.03		1		
240413	**Very good identification**						**240413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.96	0.77		99		
Staphylococcus simulans		0.04	0.02		1		
240433	**Good identification**						**240433**
Staphylococcus cohnii subsp. urealyticum		95.92	0.36	XYL 1			
Staphylococcus xylosus		4.08	-.01				
240440	**Low discrimination**						**240440**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	YEP	
Staphylococcus warneri		88.34	0.68	SUC 95	1	1	43	
Staphylococcus cohnii subsp. cohnii		7.69	0.41		1	99	12	
Macrococcus equipersicus		2.40	0.26		99	99	99	
240441	**Low discrimination**							**240441**
					NOV			
Staphylococcus cohnii subsp. urealyticum		68.49	0.51	MAN 91	99			
Staphylococcus cohnii subsp. cohnii		25.91	0.42	LAC 91	99			
Staphylococcus warneri		5.60	0.33		1			
240442	**Good identification**							**240442**
					NOV			
Staphylococcus warneri		92.02	0.58	SUC 95	1			
Staphylococcus cohnii subsp. urealyticum		7.03	0.31		99			
Staphylococcus pasteuri		0.95	0.03		1			
240443	**Very good identification**							**240443**
					NOV			
Staphylococcus cohnii subsp. urealyticum		99.73	0.72	MAN 91	99			
Staphylococcus warneri		0.27	0.23		1			
240450	**Low discrimination**							**240450**
					OXI	NOV		
Staphylococcus warneri		40.10	0.65	SUC 95	1	1		
Macrococcus equipersicus		28.53	0.52		99	99		
Macrococcus bovicus		17.88	0.47		99	99		
240451	**Good identification**							**240451**
					NOV			
Staphylococcus cohnii subsp. urealyticum		94.52	0.72	LAC 91	99			
Staphylococcus cohnii subsp. cohnii		4.50	0.44		99			
Staphylococcus warneri		0.55	0.30		1			
240452	**Low discrimination**							**240452**
					NOV			
Staphylococcus cohnii subsp. urealyticum		42.86	0.52	MNS 99	99			
Staphylococcus warneri		40.16	0.55		1			
Staphylococcus kloosii		7.14	0.48		99			
240453	**Very good identification**							**240453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		99.96	0.93		99	53		
Staphylococcus warneri		0.02	0.20		1	8		
Aerococcus viridans		0.01	0.24		1	95		
240460	**Good identification**							**240460**
Staphylococcus warneri		100.0	0.27	SUC 95				
240463	**Acceptable identification**							**240463**
Staphylococcus cohnii subsp. urealyticum		93.95	0.31	MAN 91				
Staphylococcus xylosus		6.05	-.03	XYL 1				
240470	**Low discrimination**							**240470**
					OXI	NOV		
Staphylococcus kloosii		38.55	0.34	PHS 99	1	99		
Staphylococcus warneri		28.48	0.24		1	1		
Macrococcus equipersicus		20.27	0.11		99	99		
240471	**Low discrimination**							**240471**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		73.83	0.31	XYL 1				
Staphylococcus xylosus		26.17	0.13	LAC 91				
240472	**Low discrimination**							**240472**
					NOV			
Staphylococcus kloosii		62.48	0.35	PHS 99	99			
Staphylococcus cohnii subsp. urealyticum		15.16	0.11		99			
Staphylococcus warneri		14.21	0.14		1			
240473	**Low discrimination**							**240473**
					NOV			
Staphylococcus cohnii subsp. urealyticum		89.89	0.52	XYL 1	99			
Staphylococcus xylosus		8.97	0.22		99			
Aerococcus viridans		1.14	0.24		1			
240500	**Low discrimination**							**240500**
					NOV	OXI		
Staphylococcus warneri		41.44	0.31	SUC 95	1	1		
Kocuria varians		31.05	0.22	MLT 95	1	13		
Macrococcus equipersicus		10.72	0.09		99	99		
240501	**Low discrimination**							**240501**
Staphylococcus cohnii subsp. urealyticum		85.07	0.28	MAN 91				
Staphylococcus cohnii subsp. cohnii		14.93	0.12	LAC 91				
240503	**Good identification**							**240503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.49	MAN 91				
240510	**Low discrimination**							**240510**
					OXI	NOV		
Macrococcus equipersicus		50.14	0.36	GLR 1	99	99		
Macrococcus bovicus		36.42	0.31		99	99		
Staphylococcus warneri		7.39	0.28		1	1		
240511	**Good identification**							**240511**
Staphylococcus cohnii subsp. urealyticum		97.84	0.49	LAC 91				
Staphylococcus cohnii subsp. cohnii		2.16	0.14					
240512	**Low discrimination**							**240512**
					NOV			
Staphylococcus cohnii subsp. urealyticum		69.88	0.29	MNS 99	99			
Staphylococcus kloosii		15.72	0.27		99			
Staphylococcus warneri		14.39	0.19		1			
240513	**Very good identification**							**240513**
Staphylococcus cohnii subsp. urealyticum		99.91	0.70					
Staphylococcus xylosus		0.09	-.02					
240540	**Good identification**							**240540**
					OXI	NOV	YEP	
Staphylococcus warneri		92.55	0.58	SUC 95	1	1	43	
Macrococcus equipersicus		3.41	0.18		99	99	99	
Staphylococcus cohnii subsp. cohnii		2.84	0.22		1	99	12	
240541	**Low discrimination**							**240541**
					NOV			
Staphylococcus cohnii subsp. urealyticum		86.33	0.44	MAN 91	99			
Staphylococcus cohnii subsp. cohnii		8.47	0.22	LAC 91	99			
Staphylococcus warneri		5.21	0.23		1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
240542	**Acceptable identification**						**240542**
					NOV		
Staphylococcus warneri		90.60	0.48	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		9.40	0.23		99		
240543	**Very good identification**						**240543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.67	0.65	MAN 91	99		
Staphylococcus warneri		0.20	0.13		1		
Staphylococcus xylosus		0.13	-.04		99		
240550	**Low discrimination**						**240550**
					OXI	NOV	
Staphylococcus warneri		31.20	0.55	SUC 95	1	1	
Macrococcus equipersicus		30.13	0.45		99	99	
Macrococcus bovicus		18.89	0.39		99	99	
240551	**Very good identification**						**240551**
					PYR		
Staphylococcus cohnii subsp. urealyticum		96.36	0.65	LAC 91	53		
Staphylococcus cohnii subsp. cohnii		1.19	0.25		1		
Staphylococcus saprophyticus subsp. bovis		1.00	0.18		99		
240552	**Low discrimination**						**240552**
					NOV		
Staphylococcus cohnii subsp. urealyticum		38.28	0.44	MNS 99	99		
Staphylococcus kloosii		28.88	0.53		99		
Staphylococcus warneri		26.44	0.45		1		
240553	**Very good identification**						**240553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.72	0.85		99		
Staphylococcus xylosus		0.21	0.21		99		
Aerococcus viridans		0.03	0.24		1		
240570	**Low discrimination**						**240570**
					OXI	NOV	
Staphylococcus kloosii		76.39	0.40	PHS 99	1	99	
Staphylococcus warneri		9.18	0.14		1	1	
Macrococcus equipersicus		8.87	0.04		99	99	
240571	**Low discrimination**						**240571**
Staphylococcus xylosus		89.32	0.33	NIT 89			
Staphylococcus cohnii subsp. urealyticum		10.68	0.23	SUC 95			
240572	**Low discrimination**						**240572**
Staphylococcus kloosii		75.17	0.41	PHS 99			
Staphylococcus arlettae		10.29	0.12				
Staphylococcus xylosus		9.48	0.02				
240573	**Low discrimination**						**240573**
Staphylococcus xylosus		68.88	0.43	NIT 89			
Staphylococcus cohnii subsp. urealyticum		29.26	0.44	SUC 95			
Staphylococcus arlettae		1.00	0.14				
240600	**Low discrimination**						**240600**
Staphylococcus warneri		88.02	0.68	MLT 95			
Staphylococcus pasteuri		11.98	0.35				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
240601	**Low discrimination**							**240601**
					OXI	PYR		
Staphylococcus warneri		79.07	0.33	MLT 95	1	8		
Kocuria kristinae		10.33	0.00	MNS 2	99	93		
Staphylococcus pasteuri		5.33	-.06		1	1		
240602	**Low discrimination**							**240602**
					YEP			
Staphylococcus warneri		85.00	0.58	MLT 95	43			
Staphylococcus pasteuri		14.18	0.27		86			
Staphylococcus simulans		0.82	0.07		1			
240610	**Low discrimination**							**240610**
					OXI	NOV		
Staphylococcus pasteuri		51.81	0.52	VPT 86	1	1		
Staphylococcus warneri		44.87	0.65	MLT 86	1	1		
Micrococcus bovicus		2.14	0.27		99	99		
240611	**Low discrimination**							**240611**
					NOV			
Staphylococcus warneri		39.96	0.30	MLT 95	1			
Staphylococcus cohnii subsp. urealyticum		37.21	0.26	MNS 2	99			
Staphylococcus pasteuri		22.83	0.11		1			
240612	**Low discrimination**							**240612**
					YEP			
Staphylococcus pasteuri		56.94	0.44	VPT 86	86			
Staphylococcus warneri		40.25	0.55	MLT 86	43			
Staphylococcus simulans		2.15	0.19		1			
240613	**Low discrimination**							**240613**
					NOV			
Staphylococcus cohnii subsp. urealyticum		71.81	0.46	SUC 3	99			
Staphylococcus simulans		22.22	0.30		1			
Staphylococcus warneri		2.54	0.20		1			
240620	**Low discrimination**							**240620**
Staphylococcus warneri		88.02	0.27	XYL 1				
Staphylococcus pasteuri		11.98	-.06	MLT 95				
240640	**Very good identification**							**240640**
					PYR			
Staphylococcus warneri		95.64	0.94		8			
Staphylococcus pasteuri		4.21	0.52		1			
Staphylococcus petrasii subsp. croceilyticus		0.08	0.05		90			
240641	**Good identification**							**240641**
					OXI	PYR		
Staphylococcus warneri		90.61	0.59	MNS 2	1	8		
Kocuria kristinae		5.61	0.20		99	93		
Staphylococcus pasteuri		1.97	0.11		1	1		
240642	**Good identification**							**240642**
					NOV			
Staphylococcus warneri		94.63	0.84		1			
Staphylococcus pasteuri		5.10	0.44		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.16	0.07		99			
240643	**Low discrimination**							**240643**
					OXI	NOV	PYR	
Staphylococcus warneri		57.64	0.50	MNS 2	1	1	8	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		34.78	0.41		1	99	53	
Kocuria kristinae		2.68	0.07		99	1	93	
240650	**Low discrimination**							**240650**
					OXI	NOV		
Staphylococcus warneri		71.96	0.91		1	1		
Staphylococcus pasteuri		26.86	0.68		1	1		
Macrococcus bovicus		0.42	0.34		99	99		
240651	**Low discrimination**							**240651**
					NOV			
Staphylococcus warneri		64.28	0.57	MNS 2	1			
Staphylococcus cohnii subsp. urealyticum		17.86	0.41		99			
Staphylococcus pasteuri		11.88	0.27		1			
240652	**Low discrimination**							**240652**
					NOV			
Staphylococcus warneri		65.80	0.82		1			
Staphylococcus pasteuri		30.10	0.60		1			
Staphylococcus saprophyticus subsp. saprophyticus		3.67	0.35		99			
240653	**Low discrimination**							**240653**
					NOV			
Staphylococcus cohnii subsp. urealyticum		80.98	0.62	SUC 3	99			
Staphylococcus warneri		9.61	0.47		1			
Staphylococcus xylosus		4.36	0.26		99			
240660	**Very good identification**							**240660**
Staphylococcus warneri		95.79	0.53	XYL 1				
Staphylococcus pasteuri		4.22	0.11					
240662	**Acceptable identification**							**240662**
Staphylococcus warneri		94.88	0.43	XYL 1				
Staphylococcus pasteuri		5.12	0.03					
240670	**Low discrimination**							**240670**
					NOV			
Staphylococcus warneri		72.16	0.50	XYL 1	1			
Staphylococcus pasteuri		26.94	0.27		1			
Staphylococcus xylosus		0.90	-.02		99			
240671	**Good identification**							**240671**
					NOV			
Staphylococcus xylosus		96.67	0.39	NIT 89	99			
Staphylococcus equorum		1.74	0.04	GAL 91	99			
Staphylococcus warneri		1.60	0.15		1			
240672	**Low discrimination**							**240672**
					NOV			
Staphylococcus arlettae		39.45	0.37	URE 1	99			
Staphylococcus warneri		36.70	0.40		1			
Staphylococcus pasteuri		16.79	0.19		1			
240673	**Low discrimination**							**240673**
Staphylococcus xylosus		85.15	0.48	NIT 89				
Staphylococcus arlettae		11.51	0.40	GAL 91				
Staphylococcus cohnii subsp. urealyticum		1.39	0.21					
240700	**Very good identification**							**240700**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		99.04	0.58	MLT 95				
Kocuria varians		0.53	0.04					
Staphylococcus pasteuri		0.43	-.06					
240702	**Good identification**							**240702**
Staphylococcus warneri		100.0	0.48	MLT 95				
240710	**Low discrimination**							**240710**
					OXI	NOV		
Staphylococcus warneri		87.33	0.55	MLT 95	1	1		
Macrococcus bovicus		5.66	0.19		99	99		
Staphylococcus pasteuri		3.23	0.11		1	1		
240712	**Good identification**							**240712**
Staphylococcus warneri		95.67	0.45	MLT 95				
Staphylococcus pasteuri		4.33	0.03					
240713	**Low discrimination**							**240713**
					NOV			
Staphylococcus cohnii subsp. urealyticum		61.32	0.39	SUC 3	99			
Staphylococcus xylosus		33.23	0.24		99			
Staphylococcus simulans		3.85	0.08		1			
240731	**Good identification**							**240731**
Staphylococcus xylosus		97.70	0.36	NIT 89				
Staphylococcus equorum		2.30	0.04	MLT 93				
240733	**Good identification**							**240733**
					YEP			
Staphylococcus xylosus		96.95	0.46	NIT 89	40			
Staphylococcus equorum		2.07	0.13	MLT 93	1			
Staphylococcus arlettae		0.99	0.14		99			
240740	**Very good identification**							**240740**
Staphylococcus warneri		99.74	0.84					
Staphylococcus pasteuri		0.14	0.11					
Staphylococcus hominis subsp. hominis		0.09	0.16					
240741	**Low discrimination**							**240741**
					OXI	NOV	PYR	
Staphylococcus warneri		85.40	0.49	MNS 2	1	1	8	
Staphylococcus xylosus		6.72	0.13		1	99	65	
Kocuria kristinae		5.58	0.10		99	1	93	
240742	**Very good identification**							**240742**
Staphylococcus warneri		99.51	0.74					
Staphylococcus hominis subsp. hominis		0.32	0.17					
Staphylococcus pasteuri		0.17	0.03					
240743	**Low discrimination**							**240743**
					NOV			
Staphylococcus warneri		39.38	0.39	MNS 2	1			
Staphylococcus cohnii subsp. urealyticum		32.25	0.33		99			
Staphylococcus xylosus		26.44	0.23		99			
240750	**Good identification**							**240750**
					NOV	PYR		
Staphylococcus warneri		93.52	0.81		1	8		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		3.87	0.43		99	99	
Staphylococcus pasteuri		1.12	0.27		1	1	
240751	**Low discrimination**						**240751**
					NOV	PYR	
Staphylococcus xylosus		47.46	0.38	NIT 89	99	65	
Staphylococcus warneri		27.90	0.46	XYL 92	1	8	
Staphylococcus saprophyticus subsp. bovis		14.12	0.31		99	99	
240752	**Good identification**						**240752**
					NOV	PYR	
Staphylococcus warneri		93.03	0.71		1	8	
Staphylococcus saprophyticus subsp. bovis		2.88	0.31		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		1.43	0.13		99	7	
240753	**Low discrimination**						**240753**
					NOV		
Staphylococcus xylosus		52.56	0.47	NIT 89	99		
Staphylococcus cohnii subsp. urealyticum		41.38	0.54	XYL 92	99		
Staphylococcus warneri		3.62	0.36		1		
240760	**Good identification**						**240760**
Staphylococcus warneri		100.0	0.43	XYL 1			
240761	**Good identification**						**240761**
Staphylococcus xylosus		100.0	0.35	NIT 89			
240762	**Low discrimination**						**240762**
					NOV		
Staphylococcus warneri		77.00	0.33	XYL 1	1		
Staphylococcus xylosus		12.13	0.03		99		
Staphylococcus arlettae		10.86	0.12		99		
240763	**Good identification**						**240763**
Staphylococcus xylosus		98.82	0.44	NIT 89			
Staphylococcus arlettae		1.19	0.14	MAN 94			
240770	**Low discrimination**						**240770**
					NOV		
Staphylococcus warneri		64.56	0.40	XYL 1	1		
Staphylococcus xylosus		25.78	0.19		99		
Staphylococcus arlettae		4.19	0.12		99		
240771	**Very good identification**						**240771**
					YEP		
Staphylococcus xylosus		98.03	0.60	NIT 89	40		
Staphylococcus equorum		1.76	0.25		1		
Staphylococcus arlettae		0.21	0.14		99		
240772	**Low discrimination**						**240772**
					NOV		
Staphylococcus arlettae		80.59	0.53	URE 1	99		
Staphylococcus xylosus		14.25	0.28		99		
Staphylococcus warneri		4.18	0.30		1		
240773	**Good identification**						**240773**
					YEP		
Staphylococcus xylosus		91.47	0.69	NIT 89	40		
Staphylococcus arlettae		6.93	0.55		99		
Staphylococcus equorum		1.48	0.34		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
241010	**Good identification**						**241010**
Micrococcus equipersicus		97.53	0.50	GLR 1			
Micrococcus bovicus		2.47	0.16				
241013	**Good identification**						**241013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	TRE 99			
241040	**Good identification**						**241040**
Micrococcus equipersicus		100.0	0.33	GLR 1			
241050	**Very good identification**						**241050**
Micrococcus equipersicus		97.86	0.59	GLR 1			
Micrococcus bovicus		2.14	0.23				
241053	**Good identification**						**241053**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.48	0.43	TRE 99	99		
Aerococcus viridans		1.52	0.17		1		
241110	**Good identification**						**241110**
Micrococcus equipersicus		97.53	0.43	GLR 1			
Micrococcus bovicus		2.47	0.08				
241150	**Good identification**						**241150**
Micrococcus equipersicus		97.86	0.51	GLR 1			
Micrococcus bovicus		2.14	0.16				
241153	**Good identification**						**241153**
					NOV		
Staphylococcus cohnii subsp. urealyticum		96.52	0.36	TRE 99	99		
Aerococcus viridans		3.48	0.17		1		
241210	**Acceptable identification**						**241210**
Micrococcus equipersicus		93.99	0.29	GLR 1			
Micrococcus bovicus		6.02	0.03	SUC 9			
241240	**Low discrimination**						**241240**
					OXI	NOV	
Staphylococcus warneri		63.93	0.27	ESL 1	1	1	
Micrococcus equipersicus		36.07	0.12	TRE 95	99	99	
241250	**Low discrimination**						**241250**
					OXI	NOV	
Micrococcus equipersicus		89.06	0.38	GLR 1	99	99	
Staphylococcus warneri		6.02	0.24	SUC 9	1	1	
Micrococcus bovicus		4.92	0.11		99	99	
241350	**Acceptable identification**						**241350**
					OXI	NOV	
Micrococcus equipersicus		90.50	0.31	GLR 1	99	99	
Micrococcus bovicus		5.00	0.03	SUC 9	99	99	
Staphylococcus warneri		4.50	0.14		1	1	
241373	**Low discrimination**						**241373**
					YEP		
Staphylococcus equorum		58.51	0.34	NIT 92	1		
Staphylococcus xylosus		37.60	0.28	TRE 99	40		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		2.76	0.14		99		
241401	**Good identification**						**241401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91			
241403	**Good identification**						**241403**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.60	0.48	MAN 91	99		
Aerococcus viridans		1.40	0.21		1		
241410	**Acceptable identification**						**241410**
Macrococcus equipersicus		94.06	0.50	GLR 1			
Macrococcus bovicus		3.57	0.19				
Macrococcus carouselicus		1.47	0.08				
241411	**Good identification**						**241411**
					OXI		
Staphylococcus cohnii subsp. urealyticum		97.69	0.48	LAC 91	1		
Macrococcus equipersicus		2.31	0.09		99		
241412	**Low discrimination**						**241412**
					OXI	NOV	
Staphylococcus cohnii subsp. urealyticum		65.04	0.28	MNS 99	1	99	
Macrococcus equipersicus		15.05	0.09		99	99	
Aerococcus viridans		11.60	0.23		1	1	
241413	**Very good identification**						**241413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.58	0.69		99		
Aerococcus viridans		0.42	0.31		1		
241433	**Low discrimination**						**241433**
					NOV		
Staphylococcus cohnii subsp. urealyticum		70.62	0.28	XYL 1	99		
Aerococcus viridans		29.38	0.31		1		
241440	**Low discrimination**						**241440**
					OXI	NOV	
Macrococcus equipersicus		85.09	0.33	GLR 1	99	99	
Staphylococcus warneri		14.92	0.27	MAN 95	1	1	
241441	**Good identification**						**241441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91			
241443	**Very good identification**						**241443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.01	0.64	MAN 91	99		
Aerococcus viridans		0.99	0.34		1		
241450	**Good identification**						**241450**
					OXI		
Macrococcus equipersicus		92.09	0.59	GLR 1	99		
Staphylococcus kloosii		3.33	0.47		1		
Macrococcus bovicus		3.02	0.27		99		
241451	**Very good identification**						**241451**
					OXI	NOV	
Staphylococcus cohnii subsp. urealyticum		98.56	0.64	LAC 91	1	99	
Macrococcus equipersicus		1.11	0.18		99	99	
Aerococcus viridans		0.33	0.24		1	1	
241452	**Low discrimination**						**241452**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		59.21	0.43	MNS 99	99	64	
Staphylococcus kloosii		25.36	0.48		99	99	
Aerococcus viridans		7.45	0.36		1	1	
241453	**Very good identification**						**241453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.70	0.84		99		
Aerococcus viridans		0.30	0.43		1		
241472	**Low discrimination**						**241472**
					NOV	YEP	
Aerococcus viridans		54.03	0.36	URE 1	1	1	
Staphylococcus kloosii		45.97	0.35		99	99	
241473	**Low discrimination**						**241473**
					NOV	PYR	
Staphylococcus cohnii subsp. urealyticum		69.60	0.43	XYL 1	99	53	
Aerococcus viridans		20.44	0.43		1	95	
Staphylococcus equorum		5.50	0.13		99	5	
241503	**Good identification**						**241503**
					NOV		
Staphylococcus cohnii subsp. urealyticum		96.80	0.41	MAN 91	99		
Aerococcus viridans		3.20	0.21		1		
241510	**Acceptable identification**						**241510**
					OXI		
Macrococcus equipersicus		92.86	0.43	GLR 1	99		
Macrococcus bovicus		3.53	0.12		99		
Staphylococcus kloosii		2.17	0.26		1		
241511	**Good identification**						**241511**
					OXI		
Staphylococcus cohnii subsp. urealyticum		97.69	0.41	LAC 91	1		
Macrococcus equipersicus		2.31	0.01		99		
241513	**Very good identification**						**241513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.03	0.61		99		
Aerococcus viridans		0.97	0.31		1		
241541	**Good identification**						**241541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91			
241543	**Very good identification**						**241543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.72	0.56	MAN 91	99		
Aerococcus viridans		2.28	0.34		1		
241550	**Low discrimination**						**241550**
					OXI		
Macrococcus equipersicus		82.40	0.51	GLR 1	99		
Staphylococcus kloosii		13.50	0.53		1		
Macrococcus bovicus		2.70	0.19		99		
241551	**Very good identification**						**241551**
					OXI	NOV	
Staphylococcus cohnii subsp. urealyticum		97.68	0.56	LAC 91	1	99	
Macrococcus equipersicus		1.10	0.10		99	99	
Aerococcus viridans		0.76	0.24		1	1	
241552	**Low discrimination**						**241552**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	YEP		
Staphylococcus kloosii		58.01	0.53	PHS 99	99	99		
Staphylococcus cohnii subsp. urealyticum		29.90	0.36		99	64		
Aerococcus viridans		8.78	0.36		1	1		
241553	**Very good identification**							**241553**
					NOV			
Staphylococcus cohnii subsp. urealyticum		99.18	0.77		99			
Aerococcus viridans		0.69	0.43		1			
Staphylococcus xylosus		0.13	0.08		99			
241563	**Low discrimination**							**241563**
					NOV			
Aerococcus viridans		53.88	0.34	URE 1	1			
Staphylococcus cohnii subsp. urealyticum		23.33	0.15		99			
Staphylococcus xylosus		22.79	0.06		99			
241570	**Low discrimination**							**241570**
					OXI	NOV	YEP	
Staphylococcus kloosii		75.79	0.40	PHS 99	1	99	99	
Macrococcus equipersicus		18.69	0.10		99	99	99	
Aerococcus viridans		5.52	0.16		1	1	1	
241572	**Low discrimination**							**241572**
					NOV	YEP		
Staphylococcus kloosii		57.40	0.41	PHS 99	99	99		
Aerococcus viridans		34.75	0.36		1	1		
Staphylococcus arlettae		7.86	0.12		99	99		
241573	**Low discrimination**							**241573**
Staphylococcus equorum		36.20	0.34	NIT 92				
Staphylococcus xylosus		29.38	0.30	SUC 99				
Staphylococcus cohnii subsp. urealyticum		19.41	0.36					
241610	**Low discrimination**							**241610**
					OXI	NOV		
Macrococcus equipersicus		55.22	0.29	GLR 1	99	99		
Macrococcus carouselicus		17.75	0.14	SUC 9	99	99		
Staphylococcus pasteuri		11.64	0.11		1	1		
241613	**Low discrimination**							**241613**
					NOV			
Staphylococcus cohnii subsp. urealyticum		88.04	0.38	SUC 3	99			
Aerococcus viridans		11.96	0.31		1			
241633	**Low discrimination**							**241633**
					NOV	PYR		
Staphylococcus equorum		82.41	0.33	NIT 92	99	5		
Staphylococcus xylosus		9.77	0.13	GAL 91	99	65		
Aerococcus viridans		7.82	0.31	MLT 91	1	95		
241640	**Good identification**							**241640**
					OXI	NOV		
Staphylococcus warneri		93.14	0.53	ESL 1	1	1		
Staphylococcus pasteuri		4.10	0.11		1	1		
Macrococcus equipersicus		2.77	0.12		99	99		
241642	**Low discrimination**							**241642**
					PYR	YEP		
Staphylococcus warneri		89.32	0.43	ESL 1	8	43		
Aerococcus viridans		5.87	0.26		95	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus pasteuri		4.82	0.03		1	86		
241643	**Low discrimination**							**241643**
					OXI	NOV		
Staphylococcus cohnii subsp. urealyticum		70.64	0.33	SUC 3	1	99		
Aerococcus viridans		22.83	0.34	MAN 91	1	1		
Kocuria kristinae		3.49	-.05		99	1		
241650	**Low discrimination**							**241650**
					OXI	NOV		
Staphylococcus warneri		44.34	0.50	ESL 1	1	1		
Macrococcus equipersicus		34.55	0.38		99	99		
Staphylococcus pasteuri		16.55	0.27		1	1		
241651	**Low discrimination**							**241651**
					NOV			
Staphylococcus cohnii subsp. urealyticum		70.14	0.33	SUC 3	99			
Staphylococcus xylosus		8.63	0.05	LAC 91	99			
Aerococcus viridans		7.56	0.24		1			
241652	**Low discrimination**							**241652**
					PYR	YEP		
Staphylococcus warneri		51.45	0.40	ESL 1	8	43		
Staphylococcus pasteuri		23.53	0.19		1	86		
Aerococcus viridans		14.00	0.36		95	1		
241653	**Low discrimination**							**241653**
					NOV			
Staphylococcus cohnii subsp. urealyticum		86.48	0.53	SUC 3	99			
Aerococcus viridans		8.29	0.43		1			
Staphylococcus xylosus		2.99	0.14		99			
241663	**Low discrimination**							**241663**
					NOV	PYR		
Aerococcus viridans		38.42	0.34	URE 1	1	95		
Staphylococcus equorum		31.03	0.13		99	5		
Staphylococcus xylosus		30.54	0.11		99	65		
241671	**Low discrimination**							**241671**
					NOV	PYR		
Staphylococcus equorum		86.85	0.45	NIT 92	99	5		
Staphylococcus xylosus		12.22	0.27	GAL 91	99	65		
Aerococcus viridans		0.93	0.24		1	95		
241672	**Low discrimination**							**241672**
					NOV	YEP	PYR	
Staphylococcus arlettae		70.82	0.37	URE 1	99	99	1	
Aerococcus viridans		17.93	0.36		1	1	95	
Staphylococcus equorum		11.26	0.13		99	1	5	
241673	**Low discrimination**							**241673**
					YEP			
Staphylococcus equorum		78.04	0.54	NIT 92	1			
Staphylococcus xylosus		12.16	0.36	GAL 91	40			
Staphylococcus arlettae		6.57	0.40		99			
241713	**Low discrimination**							**241713**
Staphylococcus cohnii subsp. urealyticum		49.83	0.30	SUC 3				
Staphylococcus xylosus		17.36	0.12					
Staphylococcus equorum		17.01	0.13					
241731	**Acceptable identification**							**241731**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus equorum		90.33	0.45	NIT 92				
Staphylococcus xylosus		9.67	0.24	MLT 91				
241733	**Low discrimination**							**241733**
					NOV	PYR		
Staphylococcus equorum		88.28	0.54	NIT 92	99	5		
Staphylococcus xylosus		10.47	0.33	MLT 91	99	65		
Aerococcus viridans		0.83	0.31		1	95		
241740	**Good identification**							**241740**
					OXI	NOV		
Staphylococcus warneri		96.13	0.43	ESL 1	1	1		
Macrococcus equipersicus		3.87	0.04		99	99		
241742	**Low discrimination**							**241742**
					PYR			
Staphylococcus warneri		82.79	0.33	ESL 1	8			
Aerococcus viridans		17.22	0.26		95			
241750	**Low discrimination**							**241750**
					OXI	NOV		
Macrococcus equipersicus		44.22	0.31	GLR 1	99	99		
Staphylococcus warneri		41.82	0.40	SUC 9	1	1		
Staphylococcus kloosii		7.24	0.32		1	99		
241752	**Low discrimination**							**241752**
					NOV	PYR	YEP	
Staphylococcus warneri		36.90	0.30	ESL 1	1	8	43	
Aerococcus viridans		31.79	0.36		1	95	1	
Staphylococcus kloosii		20.77	0.33		99	52	99	
241753	**Low discrimination**							**241753**
Staphylococcus cohnii subsp. urealyticum		37.34	0.46	SUC 3				
Staphylococcus xylosus		30.49	0.35					
Staphylococcus equorum		22.74	0.34					
241763	**Low discrimination**							**241763**
					NOV	PYR		
Staphylococcus equorum		45.93	0.34	NIT 92	99	5		
Staphylococcus xylosus		45.20	0.32	MAN 99	99	65		
Aerococcus viridans		5.62	0.34		1	95		
241771	**Low discrimination**							**241771**
Staphylococcus equorum		87.26	0.66	NIT 92				
Staphylococcus xylosus		12.28	0.47					
Staphylococcus gallinarum		0.22	0.10					
241772	**Low discrimination**							**241772**
					NOV	YEP	PYR	
Staphylococcus arlettae		72.47	0.53	URE 1	99	99	1	
Staphylococcus equorum		20.56	0.34		99	1	5	
Aerococcus viridans		3.24	0.36		1	1	95	
241773	**Low discrimination**							**241773**
					YEP			
Staphylococcus equorum		82.70	0.75	NIT 92	1			
Staphylococcus xylosus		12.88	0.57		40			
Staphylococcus arlettae		3.90	0.55		99			
242000	**Low discrimination**							**242000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI		
Kocuria rosea		39.39	0.53	URE 1	35		
Dermacoccus nishinomiyaensis		39.09	0.54		93		
Kocuria varians		19.56	0.55		13		
242002	**Low discrimination**						**242002**
					OXI		
Kocuria varians		87.44	0.44	GLR 1	13		
Kocuria rosea		6.31	0.12		35		
Dermacoccus nishinomiyaensis		6.26	0.13		93		
242010	**Low discrimination**						**242010**
					YEP	NOV	
Kocuria rosea		80.88	0.33	URE 1	1	1	
Dermacoccus nishinomiyaensis		7.30	0.13	MAN 10	1	80	
Macrococcus bovicus		6.88	0.16		99	99	
242020	**Low discrimination**						**242020**
					OXI		
Kocuria rosea		69.91	0.45	URE 1	35		
Kocuria varians		28.46	0.46		13		
Dermacoccus nishinomiyaensis		1.64	0.13		93		
242022	**Acceptable identification**						**242022**
Kocuria varians		91.91	0.35	GLR 1			
Kocuria rosea		8.09	0.04				
242040	**Low discrimination**						**242040**
Kocuria varians		69.71	0.42	GLR 1			
Staphylococcus warneri		13.30	0.34				
Kocuria rosea		6.46	0.12				
242042	**Low discrimination**						**242042**
Kocuria varians		78.02	0.30	GLR 1			
Staphylococcus warneri		17.59	0.24				
Staphylococcus hominis subsp. hominis		4.40	0.06				
242053	**Good identification**						**242053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	TRE 99			
242060	**Low discrimination**						**242060**
Kocuria varians		89.85	0.32	GLR 1			
Kocuria rosea		10.16	0.04				
242100	**Low discrimination**						**242100**
					OXI		
Kocuria varians		78.91	0.59	GLR 1	13		
Dermacoccus nishinomiyaensis		20.02	0.39		93		
Kocuria rosea		1.07	0.12		35		
242101	**Good identification**						**242101**
					OXI		
Kocuria varians		97.14	0.37	GLR 1	13		
Dermacoccus nishinomiyaensis		2.86	-.02	MNS 8	93		
242102	**Good identification**						**242102**
					OXI		
Kocuria varians		99.10	0.48	GLR 1	13		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Dermacoccus nishinomiyaensis		0.90	-.02		93			
242120	**Good identification**							**242120**
					OXI			
Kocuria varians		97.67	0.50	GLR 1	13			
Kocuria rosea		1.62	0.04		35			
Dermacoccus nishinomiyaensis		0.71	-.02		93			
242121	**Good identification**							**242121**
Kocuria varians		100.0	0.28	GLR 1				
242122	**Good identification**							**242122**
Kocuria varians		100.0	0.38	GLR 1				
242140	**Acceptable identification**							**242140**
					NOV			
Kocuria varians		92.85	0.45	GLR 1	1			
Staphylococcus warneri		3.73	0.24		1			
Staphylococcus hominis subsp. novobiosepticus		1.43	-.02		99			
242142	**Low discrimination**							**242142**
Kocuria varians		89.48	0.34	GLR 1				
Staphylococcus warneri		4.25	0.14					
Staphylococcus hominis subsp. hominis		3.64	0.07					
242150	**Low discrimination**							**242150**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		69.79	0.31	TRE 99	1	99	99	
Macrococcus bovicus		15.94	0.16		99	99	1	
Staphylococcus warneri		8.02	0.21		1	1	8	
242153	**Low discrimination**							**242153**
Staphylococcus cohnii subsp. urealyticum		65.14	0.29	TRE 99				
Staphylococcus xylosus		25.48	0.11					
Staphylococcus saprophyticus subsp. bovis		9.38	0.06					
242160	**Good identification**							**242160**
Kocuria varians		100.0	0.36	GLR 1				
242173	**Good identification**							**242173**
Staphylococcus xylosus		100.0	0.33	SUC 95				
242200	**Low discrimination**							**242200**
					OXI			
Kocuria rosea		40.78	0.33	URE 1	35			
Kocuria varians		24.85	0.38	SUC 10	13			
Dermacoccus nishinomiyaensis		23.25	0.29		93			
242213	**Good identification**							**242213**
Staphylococcus simulans		100.0	0.32	ARG 92				
242240	**Low discrimination**							**242240**
					NOV			
Staphylococcus warneri		60.01	0.60	TRE 95	1			
Staphylococcus hominis subsp. hominis		22.19	0.46		1			
Staphylococcus hominis subsp. novobiosepticus		12.65	0.30		99			
242242	**Low discrimination**							**242242**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus hominis subsp. hominis		47.42	0.47	GLR 1	1		
Staphylococcus warneri		36.41	0.51		1		
Staphylococcus hominis subsp. novobiosepticus		12.40	0.24		99		
242250	**Low discrimination**						**242250**
					NOV	PYR	
Staphylococcus warneri		60.63	0.57	TRE 95	1	8	
Staphylococcus pasteuri		24.81	0.35		1	1	
Staphylococcus saprophyticus subsp. bovis		8.77	0.31		99	99	
242252	**Low discrimination**						**242252**
Staphylococcus warneri		51.30	0.48	TRE 95			
Staphylococcus pasteuri		25.73	0.27				
Staphylococcus hominis subsp. hominis		13.79	0.30				
242271	**Good identification**						**242271**
Staphylococcus xylosus		100.0	0.29	GAL 91			
242273	**Good identification**						**242273**
Staphylococcus xylosus		100.0	0.39	GAL 91			
242300	**Low discrimination**						**242300**
					OXI		
Kocuria varians		81.70	0.41	GLR 1	13		
Dermacoccus nishinomiyaensis		9.70	0.14	SUC 12	93		
Staphylococcus warneri		5.28	0.24		1		
242302	**Low discrimination**						**242302**
Kocuria varians		84.08	0.30	GLR 1			
Staphylococcus warneri		6.42	0.14	SUC 12			
Staphylococcus hominis subsp. hominis		5.51	0.07				
242303	**Good identification**						**242303**
Staphylococcus hyicus		96.50	0.28	ARG 96			
Kocuria varians		3.50	0.08	PHS 92			
242320	**Good identification**						**242320**
Kocuria varians		100.0	0.32	GLR 1			
242331	**Good identification**						**242331**
Staphylococcus xylosus		100.0	0.27	TRE 96			
242333	**Good identification**						**242333**
Staphylococcus xylosus		100.0	0.36	TRE 96			
242340	**Low discrimination**						**242340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		44.86	0.39	VPT 90	99		
Staphylococcus hominis subsp. hominis		28.43	0.47	GLR 1	1		
Staphylococcus warneri		22.41	0.50		1		
242341	**Low discrimination**						**242341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		83.26	0.27	VPT 90	99		
Staphylococcus hominis subsp. hominis		4.31	0.12	GLR 1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		3.99	0.03		99		
242342	**Low discrimination**						**242342**
					NOV		
Staphylococcus hominis subsp. hominis		50.04	0.48	GLR 1	1		
Staphylococcus hominis subsp. novobiosepticus		36.22	0.34		99		
Staphylococcus warneri		11.20	0.40		1		
242350	**Low discrimination**						**242350**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		58.20	0.43	TRE 99	99	99	
Staphylococcus warneri		31.77	0.47		1	8	
Staphylococcus hominis subsp. hominis		8.32	0.30		1	1	
242351	**Low discrimination**						**242351**
					NOV	PYR	
Staphylococcus xylosus		52.03	0.28	TRE 96	99	65	
Staphylococcus saprophyticus subsp. bovis		45.92	0.31	XYL 92	99	99	
Staphylococcus warneri		2.05	0.12		1	8	
242352	**Low discrimination**						**242352**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		40.58	0.31	TRE 99	99	99	
Staphylococcus warneri		29.54	0.37		1	8	
Staphylococcus hominis subsp. hominis		27.23	0.31		1	1	
242353	**Acceptable identification**						**242353**
					NOV		
Staphylococcus xylosus		91.56	0.37	TRE 96	99		
Staphylococcus saprophyticus subsp. bovis		7.10	0.18	XYL 92	99		
Staphylococcus simulans		1.34	0.03		1		
242363	**Good identification**						**242363**
Staphylococcus xylosus		100.0	0.35	TRE 96			
242371	**Good identification**						**242371**
Staphylococcus xylosus		99.39	0.50	TRE 96			
Staphylococcus equorum		0.62	0.06				
242373	**Very good identification**						**242373**
Staphylococcus xylosus		99.44	0.59	TRE 96			
Staphylococcus equorum		0.56	0.14				
242400	**Low discrimination**						**242400**
Kocuria varians		41.81	0.32	GLR 1			
Staphylococcus warneri		23.26	0.34	TRE 7			
Kocuria rosea		11.30	0.12				
242403	**Low discrimination**						**242403**
					NOV		
Staphylococcus cohnii subsp. urealyticum		87.95	0.41	MAN 91	99		
Staphylococcus simulans		12.06	0.18		1		
242411	**Acceptable identification**						**242411**
					PYR		
Staphylococcus cohnii subsp. urealyticum		93.28	0.41	LAC 91	53		
Staphylococcus cohnii subsp. cohnii		3.39	0.11		1		
Staphylococcus saprophyticus subsp. bovis		3.33	0.06		99		
242413	**Good identification**						**242413**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus cohnii subsp. urealyticum		94.70	0.62		99		
Staphylococcus simulans		5.14	0.30		1		
Staphylococcus xylosus		0.16	-.04		99		
242440	**Low discrimination**						**242440**
					YEP		
Staphylococcus warneri		86.34	0.60	SUC 95	43		
Staphylococcus pasteuri		4.17	0.19		86		
Staphylococcus auricularis		3.10	0.22		1		
242441	**Low discrimination**						**242441**
					NOV		
Staphylococcus cohnii subsp. urealyticum		68.37	0.36	MAN 91	99		
Staphylococcus warneri		13.59	0.26	LAC 91	1		
Staphylococcus cohnii subsp. cohnii		11.03	0.19		99		
242442	**Low discrimination**						**242442**
Staphylococcus warneri		83.92	0.51	SUC 95			
Staphylococcus pasteuri		4.96	0.11				
Staphylococcus hominis subsp. hominis		3.50	0.16				
242443	**Very good identification**						**242443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.29	0.57	MAN 91	99		
Staphylococcus simulans		1.07	0.11		1		
Staphylococcus warneri		0.64	0.16		1		
242450	**Low discrimination**						**242450**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		69.34	0.59		99	99	
Staphylococcus warneri		19.37	0.57		1	8	
Staphylococcus pasteuri		7.93	0.35		1	1	
242451	**Low discrimination**						**242451**
					PYR		
Staphylococcus cohnii subsp. urealyticum		59.78	0.57	LAC 91	53		
Staphylococcus saprophyticus subsp. bovis		37.31	0.47		99		
Staphylococcus cohnii subsp. cohnii		1.21	0.21		1		
242452	**Low discrimination**						**242452**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		57.35	0.47		99	99	
Staphylococcus warneri		21.36	0.48		1	8	
Staphylococcus pasteuri		10.71	0.27		1	1	
242453	**Very good identification**						**242453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.64	0.77		99		
Staphylococcus saprophyticus subsp. bovis		1.51	0.34		99		
Staphylococcus simulans		0.42	0.23		1		
242471	**Acceptable identification**						**242471**
Staphylococcus xylosus		90.92	0.31	GAL 91			
Staphylococcus cohnii subsp. urealyticum		5.59	0.16	SUC 95			
Staphylococcus saprophyticus subsp. bovis		3.49	0.06				
242473	**Low discrimination**						**242473**
Staphylococcus xylosus		82.06	0.41	GAL 91			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		17.94	0.36	SUC 95				
242500	**Low discrimination**							**242500**
					OXI			
Kocuria varians		86.87	0.36	GLR 1	13			
Staphylococcus warneri		10.17	0.24	TRE 7	1			
Dermacoccus nishinomiyaensis		2.96	-.02		93			
242503	**Low discrimination**							**242503**
					NOV			
Staphylococcus cohnii subsp. urealyticum		76.83	0.34	MAN 91	99			
Staphylococcus hyicus		19.77	0.09		1			
Kocuria varians		3.41	0.03		1			
242510	**Low discrimination**							**242510**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		79.26	0.31	MLT 99	1	99	99	
Micrococcus bovicus		11.64	0.12		99	99	1	
Staphylococcus warneri		9.11	0.21		1	1	8	
242511	**Low discrimination**							**242511**
Staphylococcus cohnii subsp. urealyticum		67.70	0.34	LAC 91				
Staphylococcus saprophyticus subsp. bovis		22.57	0.18					
Staphylococcus xylosus		9.73	0.07					
242513	**Good identification**							**242513**
					NOV			
Staphylococcus cohnii subsp. urealyticum		94.37	0.54		99			
Staphylococcus xylosus		3.82	0.16		99			
Staphylococcus simulans		1.04	0.08		1			
242520	**Good identification**							**242520**
Kocuria varians		100.0	0.26	GLR 1				
242531	**Good identification**							**242531**
Staphylococcus xylosus		100.0	0.29	SUC 95				
242533	**Good identification**							**242533**
Staphylococcus xylosus		97.87	0.38	SUC 95				
Staphylococcus cohnii subsp. urealyticum		2.13	0.13	MLT 93				
242540	**Low discrimination**							**242540**
					NOV	PYR		
Staphylococcus warneri		67.41	0.50	SUC 95	1	8		
Staphylococcus saprophyticus subsp. bovis		22.36	0.31		99	99		
Kocuria varians		6.65	0.22		1	10		
242541	**Low discrimination**							**242541**
Staphylococcus cohnii subsp. urealyticum		51.05	0.28	MAN 91				
Staphylococcus saprophyticus subsp. bovis		30.37	0.18	LAC 91				
Staphylococcus xylosus		11.10	0.06					
242542	**Low discrimination**							**242542**
					NOV	PYR		
Staphylococcus warneri		65.88	0.40	SUC 95	1	8		
Staphylococcus saprophyticus subsp. bovis		16.39	0.18		99	99		
Staphylococcus hominis subsp. hominis		9.42	0.17		1	1		
242543	**Acceptable identification**							**242543**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		92.95	0.49	MAN 91				
Staphylococcus xylosus		5.69	0.15					
Staphylococcus saprophyticus subsp. bovis		1.37	0.06					
242550	**Very good identification**							**242550**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		97.42	0.72		1	99	99	
Staphylococcus warneri		2.15	0.47		1	1	8	
Macrococcus bovicus		0.34	0.19		99	99	1	
242551	**Low discrimination**							**242551**
Staphylococcus saprophyticus subsp. bovis		81.22	0.59					
Staphylococcus cohnii subsp. urealyticum		13.94	0.49					
Staphylococcus xylosus		4.70	0.30					
242552	**Very good identification**							**242552**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		95.37	0.59		99	99		
Staphylococcus warneri		2.80	0.37		1	8		
Staphylococcus cohnii subsp. urealyticum		1.67	0.29		99	53		
242553	**Low discrimination**							**242553**
Staphylococcus cohnii subsp. urealyticum		80.66	0.70					
Staphylococcus saprophyticus subsp. bovis		11.62	0.47					
Staphylococcus xylosus		7.65	0.40					
242561	**Good identification**							**242561**
Staphylococcus xylosus		100.0	0.27	SUC 95				
242563	**Good identification**							**242563**
Staphylococcus xylosus		98.59	0.37	SUC 95				
Staphylococcus cohnii subsp. urealyticum		1.42	0.08	MAN 94				
242570	**Low discrimination**							**242570**
Staphylococcus saprophyticus subsp. bovis		85.75	0.31	XYL 1				
Staphylococcus xylosus		14.25	0.11					
242571	**Good identification**							**242571**
					PYR			
Staphylococcus xylosus		97.78	0.52	SUC 95	65			
Staphylococcus saprophyticus subsp. bovis		1.49	0.18		99			
Staphylococcus equorum		0.48	0.06		5			
242573	**Very good identification**							**242573**
Staphylococcus xylosus		98.52	0.61	SUC 95				
Staphylococcus cohnii subsp. urealyticum		0.91	0.29					
Staphylococcus equorum		0.44	0.14					
242600	**Low discrimination**							**242600**
					YEP			
Staphylococcus warneri		55.05	0.60	MLT 95	43			
Staphylococcus pasteuri		42.81	0.43		86			
Staphylococcus auricularis		0.78	0.14		1			
242602	**Low discrimination**							**242602**
					YEP			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		39.28	0.51	MLT 95	43		
Staphylococcus pasteuri		37.43	0.35		86		
Staphylococcus simulans		21.22	0.35		1		
242603	**Low discrimination**						**242603**
Staphylococcus simulans		75.71	0.46	ARG 92			
Staphylococcus hyicus		21.84	0.28	BGA 90			
Staphylococcus warneri		0.85	0.16				
242610	**Low discrimination**						**242610**
					NOV	PYR	
Staphylococcus pasteuri		84.90	0.60	VPT 86	1	1	
Staphylococcus warneri		12.87	0.57	MLT 86	1	8	
Staphylococcus saprophyticus subsp. bovis		1.86	0.31		99	99	
242612	**Low discrimination**						**242612**
					YEP		
Staphylococcus pasteuri		66.39	0.52	VPT 86	86		
Staphylococcus simulans		24.51	0.48	MLT 86	1		
Staphylococcus warneri		8.21	0.48		43		
242613	**Good identification**						**242613**
					NOV		
Staphylococcus simulans		94.19	0.59	ARG 92	1		
Staphylococcus xylosus		2.36	0.22	BGA 90	99		
Staphylococcus cohnii subsp. urealyticum		2.24	0.31		99		
242631	**Good identification**						**242631**
Staphylococcus xylosus		96.76	0.34	GAL 91			
Staphylococcus equorum		3.24	0.06	MLT 93			
242633	**Acceptable identification**						**242633**
					NOV		
Staphylococcus xylosus		93.86	0.44	GAL 91	99		
Staphylococcus simulans		3.30	0.17	MLT 93	1		
Staphylococcus equorum		2.84	0.14		99		
242640	**Low discrimination**						**242640**
Staphylococcus warneri		75.27	0.87				
Staphylococcus pasteuri		18.92	0.60				
Staphylococcus hominis subsp. hominis		4.64	0.56				
242641	**Low discrimination**						**242641**
					NOV	PYR	
Staphylococcus warneri		75.27	0.52	MNS 2	1	8	
Staphylococcus pasteuri		9.37	0.19		1	1	
Staphylococcus saprophyticus subsp. bovis		5.08	0.18		99	99	
242642	**Low discrimination**						**242642**
Staphylococcus warneri		64.80	0.77				
Staphylococcus pasteuri		19.96	0.52				
Staphylococcus hominis subsp. hominis		14.07	0.58				
242643	**Low discrimination**						**242643**
					NOV		
Staphylococcus simulans		50.37	0.39	ARG 92	1		
Staphylococcus warneri		24.04	0.42	BGA 90	1		
Staphylococcus xylosus		9.52	0.21		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
242650	**Low discrimination**						**242650**
					NOV	PYR	
Staphylococcus pasteuri		54.67	0.76	VPT 86	1	1	
Staphylococcus warneri		25.64	0.84		1	8	
Staphylococcus saprophyticus subsp. bovis		19.32	0.72		99	99	
242651	**Low discrimination**						**242651**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		74.11	0.59		99	99	
Staphylococcus pasteuri		8.47	0.35		1	1	
Staphylococcus xylosus		8.05	0.36		99	65	
242652	**Low discrimination**						**242652**
					NOV	PYR	
Staphylococcus pasteuri		60.81	0.68	VPT 86	1	1	
Staphylococcus warneri		23.27	0.74		1	8	
Staphylococcus saprophyticus subsp. bovis		13.15	0.59		99	99	
242653	**Low discrimination**						**242653**
					NOV		
Staphylococcus simulans		35.21	0.51	ARG 92	1		
Staphylococcus xylosus		26.06	0.45	BGA 90	99		
Staphylococcus saprophyticus subsp. bovis		21.06	0.47		99		
242660	**Low discrimination**						**242660**
Staphylococcus warneri		76.16	0.46	XYL 1			
Staphylococcus pasteuri		19.15	0.19				
Staphylococcus hominis subsp. hominis		4.69	0.15				
242661	**Good identification**						**242661**
					NOV		
Staphylococcus xylosus		98.14	0.33	GAL 91	99		
Staphylococcus warneri		1.86	0.11	MAN 94	1		
242662	**Low discrimination**						**242662**
Staphylococcus warneri		61.80	0.36	XYL 1			
Staphylococcus pasteuri		19.04	0.11				
Staphylococcus hominis subsp. hominis		13.42	0.16				
242663	**Good identification**						**242663**
Staphylococcus xylosus		100.0	0.42	GAL 91			
242670	**Low discrimination**						**242670**
					NOV	PYR	
Staphylococcus pasteuri		51.74	0.35	VPT 86	1	1	
Staphylococcus warneri		24.26	0.43	XYL 1	1	8	
Staphylococcus saprophyticus subsp. bovis		18.29	0.31		99	99	
242671	**Very good identification**						**242671**
					PYR		
Staphylococcus xylosus		96.75	0.58	GAL 91	65		
Staphylococcus equorum		2.47	0.26		5		
Staphylococcus saprophyticus subsp. bovis		0.78	0.18		99		
242672	**Low discrimination**						**242672**
					NOV		
Staphylococcus pasteuri		41.75	0.27	VPT 86	1		
Staphylococcus xylosus		32.12	0.26	XYL 1	99		
Staphylococcus warneri		15.98	0.33		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
242673	**Very good identification**						**242673**
					NOV		
Staphylococcus xylosus		97.53	0.67	GAL 91	99		
Staphylococcus equorum		2.25	0.35		99		
Staphylococcus simulans		0.12	0.10		1		
242700	**Low discrimination**						**242700**
Staphylococcus warneri		87.71	0.50	MLT 95			
Kocuria varians		5.38	0.18				
Staphylococcus hominis subsp. hominis		3.56	0.16				
242701	**Low discrimination**						**242701**
					NOV		
Staphylococcus hyicus		89.63	0.28	ARG 96	1		
Staphylococcus xylosus		7.05	0.09	PHS 92	99		
Staphylococcus warneri		3.32	0.15	LAC 92	1		
242702	**Low discrimination**						**242702**
Staphylococcus warneri		63.86	0.40	MLT 95			
Staphylococcus hyicus		12.26	0.09				
Staphylococcus simulans		9.50	0.14				
242703	**Good identification**						**242703**
					NOV		
Staphylococcus hyicus		95.45	0.50	ARG 96	1		
Staphylococcus simulans		2.50	0.24	PHS 92	1		
Staphylococcus xylosus		1.86	0.18		99		
242710	**Low discrimination**						**242710**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		59.87	0.43	MLT 99	99	99	
Staphylococcus warneri		32.68	0.47		1	8	
Staphylococcus pasteuri		6.90	0.19		1	1	
242711	**Low discrimination**						**242711**
					NOV	PYR	
Staphylococcus xylosus		66.21	0.33	XYL 92	99	65	
Staphylococcus saprophyticus subsp. bovis		32.35	0.31	MLT 93	99	99	
Staphylococcus warneri		1.44	0.12		1	8	
242712	**Low discrimination**						**242712**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		38.87	0.31	MLT 99	99	99	
Staphylococcus warneri		28.29	0.37		1	8	
Staphylococcus simulans		23.24	0.26		1	70	
242713	**Low discrimination**						**242713**
					NOV		
Staphylococcus xylosus		68.49	0.43	XYL 92	99		
Staphylococcus simulans		23.54	0.37	MLT 93	1		
Staphylococcus saprophyticus subsp. bovis		2.94	0.18		99		
242721	**Good identification**						**242721**
Staphylococcus xylosus		100.0	0.31	MAN 94			
242723	**Good identification**						**242723**
					NOV		
Staphylococcus xylosus		95.68	0.40	MAN 94	99		
Staphylococcus hyicus		4.32	0.09	MLT 93	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
242731	**Very good identification**						**242731**
	Staphylococcus xylosus	96.76	0.55	MLT 93			
	Staphylococcus equorum	3.24	0.26				
242733	**Very good identification**						**242733**
	Staphylococcus xylosus	97.06	0.65	MLT 93			
	Staphylococcus equorum	2.94	0.35				
242740	**Low discrimination**						**242740**
					NOV	PYR	
	Staphylococcus warneri	77.34	0.76		1	8	
	Staphylococcus hominis subsp. hominis	16.35	0.57		1	1	
	Staphylococcus saprophyticus subsp. bovis	5.40	0.43		99	99	
242741	**Low discrimination**						**242741**
					NOV	PYR	
	Staphylococcus xylosus	41.80	0.32	MAN 94	99	65	
	Staphylococcus warneri	28.16	0.42	XYL 92	1	8	
	Staphylococcus saprophyticus subsp. bovis	24.09	0.31		99	99	
242742	**Low discrimination**						**242742**
					NOV	PYR	
	Staphylococcus warneri	55.04	0.67		1	8	
	Staphylococcus hominis subsp. hominis	40.99	0.58		1	1	
	Staphylococcus saprophyticus subsp. bovis	2.88	0.31		99	99	
242743	**Low discrimination**						**242743**
					NOV		
	Staphylococcus xylosus	76.50	0.41	MAN 94	99		
	Staphylococcus warneri	6.04	0.32	XYL 92	1		
	Staphylococcus hominis subsp. hominis	4.49	0.23		1		
242750	**Low discrimination**						**242750**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	89.49	0.84		99	99	
	Staphylococcus warneri	9.37	0.73		1	8	
	Staphylococcus pasteuri	0.64	0.35		1	1	
242751	**Low discrimination**						**242751**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	77.78	0.72		99	99	
	Staphylococcus xylosus	21.36	0.56		99	65	
	Staphylococcus warneri	0.67	0.39		1	8	
242752	**Low discrimination**						**242752**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	84.50	0.72		99	99	
	Staphylococcus warneri	11.80	0.64		1	8	
	Staphylococcus hominis subsp. hominis	1.81	0.41		1	1	
242753	**Low discrimination**						**242753**
	Staphylococcus xylosus	73.77	0.66	XYL 92			
	Staphylococcus saprophyticus subsp. bovis	23.59	0.59				
	Staphylococcus cohnii subsp. urealyticum	1.27	0.39				
242760	**Low discrimination**						**242760**
					NOV		
	Staphylococcus warneri	61.36	0.35	XYL 1	1		
	Staphylococcus xylosus	21.38	0.13		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		12.97	0.16		1		
242761	**Very good identification**						**242761**
Staphylococcus xylosus		99.60	0.54	MAN 94			
Staphylococcus equorum		0.40	0.06				
242763	**Very good identification**						**242763**
Staphylococcus xylosus		99.56	0.63	MAN 94			
Staphylococcus equorum		0.36	0.14				
Staphylococcus gallinarum		0.07	-.03				
242770	**Low discrimination**						**242770**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		52.23	0.43	XYL 1	99	99	
Staphylococcus xylosus		41.24	0.37		99	65	
Staphylococcus warneri		5.47	0.32		1	8	
242771	**Very good identification**						**242771**
					PYR		
Staphylococcus xylosus		97.16	0.78		65		
Staphylococcus equorum		2.48	0.47		5		
Staphylococcus saprophyticus subsp. bovis		0.31	0.31		99		
242772	**Low discrimination**						**242772**
					PYR		
Staphylococcus xylosus		86.44	0.47	MNS 99	65		
Staphylococcus saprophyticus subsp. bovis		9.62	0.31		99		
Staphylococcus equorum		1.99	0.14		5		
242773	**Very good identification**						**242773**
Staphylococcus xylosus		97.67	0.88				
Staphylococcus equorum		2.25	0.55				
Staphylococcus gallinarum		0.05	0.18				
243053	**Good identification**						**243053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	TRE 99			
243273	**Low discrimination**						**243273**
Staphylococcus equorum		68.87	0.35	GAL 91			
Staphylococcus xylosus		31.13	0.26	TRE 99			
243333	**Low discrimination**						**243333**
Staphylococcus equorum		74.40	0.35	TRE 99			
Staphylococcus xylosus		25.60	0.24	MLT 91			
243371	**Low discrimination**						**243371**
Staphylococcus equorum		69.62	0.47	TRE 99			
Staphylococcus xylosus		28.43	0.38				
Staphylococcus gallinarum		1.51	0.10				
243373	**Low discrimination**						**243373**
Staphylococcus equorum		67.90	0.55	TRE 99			
Staphylococcus xylosus		30.69	0.47				
Staphylococcus gallinarum		1.41	0.18				
243403	**Good identification**						**243403**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.33	MAN 91			
243411	**Good identification**						**243411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.33	LAC 91			
243413	**Very good identification**						**243413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.53				
243441	**Good identification**						**243441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91			
243443	**Good identification**						**243443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	MAN 91			
243451	**Good identification**						**243451**
Staphylococcus cohnii subsp. urealyticum		97.52	0.48	LAC 91			
Staphylococcus saprophyticus subsp. bovis		1.58	0.06				
Staphylococcus xylosus		0.90	-.03				
243452	**Low discrimination**						**243452**
Staphylococcus cohnii subsp. urealyticum		86.30	0.28	MNS 99			
Staphylococcus saprophyticus subsp. bovis		13.70	0.06				
243453	**Very good identification**						**243453**
Staphylococcus cohnii subsp. urealyticum		99.74	0.69				
Staphylococcus xylosus		0.26	0.06				
243473	**Low discrimination**						**243473**
Staphylococcus equorum		56.65	0.35	GAL 91			
Staphylococcus xylosus		32.35	0.28	SUC 99			
Staphylococcus cohnii subsp. urealyticum		11.00	0.28				
243513	**Good identification**						**243513**
Staphylococcus cohnii subsp. urealyticum		97.47	0.46				
Staphylococcus xylosus		2.53	0.04				
243533	**Low discrimination**						**243533**
Staphylococcus equorum		69.00	0.35	SUC 99			
Staphylococcus xylosus		29.99	0.26	MLT 91			
Staphylococcus cohnii subsp. urealyticum		1.01	0.05				
243543	**Good identification**						**243543**
Staphylococcus cohnii subsp. urealyticum		96.22	0.41	MAN 91			
Staphylococcus xylosus		3.79	0.03				
243550	**Low discrimination**						**243550**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		87.44	0.31	ESL 1	1	99	
Macrococcus equipersicus		9.23	0.10		99	1	
Macrococcus bovicus		3.33	0.00		99	1	
243551	**Low discrimination**						**243551**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		70.59	0.41	LAC 91			
Staphylococcus xylosus		15.29	0.18				
Staphylococcus saprophyticus subsp. bovis		10.68	0.18				
243553	**Good identification**						**243553**
Staphylococcus cohnii subsp. urealyticum		92.84	0.61				
Staphylococcus xylosus		5.66	0.27				
Staphylococcus equorum		1.15	0.14				
243571	**Low discrimination**						**243571**
Staphylococcus equorum		65.04	0.47	SUC 99			
Staphylococcus xylosus		33.55	0.40				
Staphylococcus gallinarum		1.41	0.10				
243573	**Low discrimination**						**243573**
Staphylococcus equorum		62.50	0.55	SUC 99			
Staphylococcus xylosus		35.69	0.49				
Staphylococcus gallinarum		1.29	0.18				
243631	**Acceptable identification**						**243631**
					OXI		
Staphylococcus equorum		90.61	0.47	GAL 91	1		
Staphylococcus xylosus		6.83	0.22	MLT 91	1		
Staphylococcus sciuri		2.56	0.18		99		
243633	**Good identification**						**243633**
					OXI		
Staphylococcus equorum		92.15	0.55	GAL 91	1		
Staphylococcus xylosus		7.69	0.31	MLT 91	1		
Staphylococcus sciuri		0.17	0.01		99		
243640	**Low discrimination**						**243640**
Staphylococcus warneri		75.18	0.46	ESL 1			
Staphylococcus pasteuri		18.90	0.19				
Staphylococcus hominis subsp. hominis		4.63	0.15				
243642	**Low discrimination**						**243642**
Staphylococcus warneri		62.41	0.36	ESL 1			
Staphylococcus pasteuri		19.22	0.11				
Staphylococcus hominis subsp. hominis		13.55	0.16				
243650	**Low discrimination**						**243650**
					NOV	PYR	
Staphylococcus pasteuri		53.48	0.35	VPT 86	1	1	
Staphylococcus warneri		25.08	0.43	ESL 1	1	8	
Staphylococcus saprophyticus subsp. bovis		18.90	0.31		99	99	
243651	**Low discrimination**						**243651**
					OXI		
Staphylococcus sciuri		37.31	0.32	URE 1	99		
Staphylococcus equorum		27.02	0.26	GLR 14	1		
Staphylococcus xylosus		23.03	0.23		1		
243652	**Low discrimination**						**243652**
					NOV	PYR	
Staphylococcus pasteuri		61.50	0.27	VPT 86	1	1	
Staphylococcus warneri		23.54	0.33	ESL 1	1	8	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		13.30	0.18		99	99	
243653	**Low discrimination**						**243653**
Staphylococcus equorum		36.93	0.35	GAL 91			
Staphylococcus xylosus		34.84	0.33	XYL 99			
Staphylococcus cohnii subsp. urealyticum		21.95	0.38				
243663	**Low discrimination**						**243663**
Staphylococcus equorum		58.37	0.35	GAL 91			
Staphylococcus xylosus		40.42	0.30	MAN 99			
Staphylococcus gallinarum		1.21	-.03				
243671	**Low discrimination**						**243671**
					OXI		
Staphylococcus equorum		89.95	0.67	GAL 91	1		
Staphylococcus xylosus		8.91	0.45		1		
Staphylococcus sciuri		0.95	0.30		99		
243672	**Acceptable identification**						**243672**
Staphylococcus equorum		90.12	0.35	GAL 91			
Staphylococcus xylosus		9.88	0.14	MNS 99			
243673	**Low discrimination**						**243673**
Staphylococcus equorum		89.90	0.76	GAL 91			
Staphylococcus xylosus		9.85	0.55				
Staphylococcus gallinarum		0.19	0.18				
243713	**Low discrimination**						**243713**
Staphylococcus equorum		53.98	0.35	XYL 99			
Staphylococcus xylosus		38.75	0.30	MLT 91			
Staphylococcus cohnii subsp. urealyticum		2.43	0.15				
243723	**Low discrimination**						**243723**
Staphylococcus equorum		64.61	0.35	MAN 99			
Staphylococcus xylosus		34.05	0.28	MLT 91			
Staphylococcus gallinarum		1.34	-.03				
243731	**Good identification**						**243731**
					OXI		
Staphylococcus equorum		92.34	0.67	MLT 91	1		
Staphylococcus xylosus		6.96	0.43		1		
Staphylococcus sciuri		0.50	0.24		99		
243732	**Acceptable identification**						**243732**
Staphylococcus equorum		92.30	0.35	MLT 91			
Staphylococcus xylosus		7.70	0.11	MNS 99			
243733	**Good identification**						**243733**
Staphylococcus equorum		92.06	0.76	MLT 91			
Staphylococcus xylosus		7.68	0.52				
Staphylococcus gallinarum		0.20	0.18				
243740	**Low discrimination**						**243740**
Staphylococcus warneri		73.38	0.35	ESL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		15.52	0.16					
Staphylococcus piscifermentans		5.98	0.14					
243743	**Low discrimination**							**243743**
Staphylococcus xylosus		80.44	0.29	MAN 94				
Staphylococcus equorum		13.49	0.14	XYL 92				
Staphylococcus cohnii subsp. urealyticum		3.33	0.10					
243750	**Low discrimination**							**243750**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		82.82	0.43	ESL 1	1	99	99	
Staphylococcus warneri		8.68	0.32		1	1	8	
Staphylococcus sciuri		5.41	0.19		99	99	1	
243751	**Low discrimination**							**243751**
					OXI			
Staphylococcus equorum		43.91	0.47	XYL 99	1			
Staphylococcus xylosus		37.43	0.44		1			
Staphylococcus sciuri		11.64	0.38		99			
243752	**Low discrimination**							**243752**
					PYR			
Staphylococcus saprophyticus subsp. bovis		61.91	0.31	ESL 1	99			
Staphylococcus equorum		12.83	0.14		5			
Staphylococcus xylosus		12.10	0.12		65			
243753	**Low discrimination**							**243753**
Staphylococcus equorum		49.33	0.55	XYL 99				
Staphylococcus xylosus		46.54	0.53					
Staphylococcus cohnii subsp. urealyticum		1.24	0.30					
243761	**Low discrimination**							**243761**
Staphylococcus equorum		54.21	0.47	MAN 99				
Staphylococcus xylosus		33.92	0.41					
Staphylococcus gallinarum		11.54	0.30					
243763	**Low discrimination**							**243763**
Staphylococcus equorum		52.77	0.55	MAN 99				
Staphylococcus xylosus		36.54	0.51					
Staphylococcus gallinarum		10.69	0.38					
243770	**Low discrimination**							**243770**
					OXI			
Staphylococcus equorum		86.92	0.47	MNS 99	1			
Staphylococcus xylosus		8.61	0.25		1			
Staphylococcus sciuri		2.15	0.17		99			
243771	**Low discrimination**							**243771**
Staphylococcus equorum		89.06	0.88					
Staphylococcus xylosus		8.82	0.66					
Staphylococcus gallinarum		1.94	0.51					
243772	**Low discrimination**							**243772**
Staphylococcus equorum		88.11	0.55	MNS 99				
Staphylococcus xylosus		9.66	0.34					
Staphylococcus gallinarum		1.82	0.18					

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
243773	**Low discrimination**					**243773**
Staphylococcus equorum		88.45	0.96			
Staphylococcus xylosus		9.69	0.75			
Staphylococcus gallinarum		1.83	0.59			
244000	**Acceptable identification**					**244000**
Dermacoccus nishinomiyaensis		90.67	0.33	GLR 1		
Micrococcus sp.		9.33	0.03	PHS 5		
244013	**Good identification**					**244013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	TRE 99		
244043	**Good identification**					**244043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
244050	**Low discrimination**					**244050**
					OXI	
Staphylococcus kloosii		84.29	0.47	TRE 99	1	
Macrococcus equipersicus		11.08	0.11		99	
Macrococcus bovicus		4.63	0.02		99	
244051	**Good identification**					**244051**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
244052	**Acceptable identification**					**244052**
Staphylococcus kloosii		91.66	0.48	TRE 99		
Staphylococcus cohnii subsp. urealyticum		8.34	0.14			
244053	**Very good identification**					**244053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	TRE 99		
244070	**Good identification**					**244070**
Staphylococcus kloosii		100.0	0.34	TRE 99		
244072	**Good identification**					**244072**
Staphylococcus kloosii		100.0	0.35	TRE 99		
244112	**Good identification**					**244112**
Staphylococcus kloosii		100.0	0.27	TRE 99		
244113	**Good identification**					**244113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	TRE 99		
244143	**Good identification**					**244143**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99		
244150	**Good identification**					**244150**
					OXI	
Staphylococcus kloosii		97.18	0.53	TRE 99	1	
Macrococcus equipersicus		2.82	0.04		99	
244151	**Good identification**					**244151**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99		
244152	**Very good identification**					**244152**
Staphylococcus kloosii		98.03	0.53	TRE 99		
Staphylococcus cohnii subsp. urealyticum		1.97	0.07			
244153	**Good identification**					**244153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	TRE 99		
244170	**Good identification**					**244170**
Staphylococcus kloosii		100.0	0.40	TRE 99		
244172	**Good identification**					**244172**
Staphylococcus kloosii		100.0	0.41	TRE 99		
244252	**Low discrimination**					**244252**
					NOV	
Staphylococcus kloosii		63.51	0.27	SUC 9	99	
Staphylococcus warneri		36.50	0.14	TRE 99	1	
244350	**Low discrimination**					**244350**
					NOV	
Staphylococcus kloosii		76.70	0.32	SUC 9	99	
Staphylococcus warneri		23.30	0.14	TRE 99	1	
244352	**Good identification**					**244352**
Staphylococcus kloosii		100.0	0.33	SUC 9		
244373	**Acceptable identification**					**244373**
Staphylococcus xylosus		91.10	0.29	NIT 89		
Staphylococcus arlettae		8.90	0.18	TRE 96		
244401	**Good identification**					**244401**
Staphylococcus cohnii subsp. urealyticum		95.73	0.40	MAN 91		
Staphylococcus cohnii subsp. cohnii		4.27	0.11	LAC 91		
244403	**Very good identification**					**244403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.60	MAN 91		
244410	**Good identification**					**244410**
					YEP	
Staphylococcus kloosii		93.85	0.62	MLT 95	99	
Staphylococcus cohnii subsp. urealyticum		3.07	0.19		64	
Staphylococcus cohnii subsp. cohnii		1.58	0.12		12	
244411	**Very good identification**					**244411**
					YEP	
Staphylococcus cohnii subsp. urealyticum		99.13	0.60	LAC 91	64	
Staphylococcus cohnii subsp. cohnii		0.56	0.13		12	
Staphylococcus kloosii		0.31	0.20		99	
244412	**Low discrimination**					**244412**
Staphylococcus kloosii		76.63	0.62	MLT 95		
Staphylococcus cohnii subsp. urealyticum		23.37	0.40			
244413	**Very good identification**					**244413**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		99.97	0.81				
Staphylococcus kloosii		0.03	0.21				
244430	**Good identification**						**244430**
Staphylococcus kloosii		100.0	0.49	MLT 95			
244432	**Good identification**						**244432**
Staphylococcus kloosii		100.0	0.50	MLT 95			
244433	**Good identification**						**244433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	XYL 1			
244440	**Low discrimination**						**244440**
					NOV	YEP	
Staphylococcus kloosii		63.78	0.47	MAN 99	99	99	
Staphylococcus warneri		16.27	0.27		1	43	
Staphylococcus cohnii subsp. cohnii		13.87	0.21		99	12	
244441	**Very good identification**						**244441**
Staphylococcus cohnii subsp. urealyticum		97.57	0.55	MAN 91			
Staphylococcus cohnii subsp. cohnii		2.43	0.21	LAC 91			
244442	**Low discrimination**						**244442**
					NOV		
Staphylococcus kloosii		50.78	0.48	MAN 99	99		
Staphylococcus cohnii subsp. urealyticum		45.23	0.35		99		
Staphylococcus warneri		3.99	0.17		1		
244443	**Very good identification**						**244443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.76	MAN 91			
244450	**Very good identification**						**244450**
					YEP		
Staphylococcus kloosii		98.30	0.88		99		
Staphylococcus cohnii subsp. urealyticum		0.96	0.35		64		
Staphylococcus cohnii subsp. cohnii		0.28	0.23		12		
244451	**Very good identification**						**244451**
					YEP		
Staphylococcus cohnii subsp. urealyticum		98.66	0.76	LAC 91	64		
Staphylococcus kloosii		1.03	0.47		99		
Staphylococcus cohnii subsp. cohnii		0.31	0.23		12		
244452	**Good identification**						**244452**
					NOV		
Staphylococcus kloosii		91.61	0.89		99		
Staphylococcus cohnii subsp. urealyticum		8.33	0.55		99		
Staphylococcus warneri		0.05	0.14		1		
244453	**Excellent identification**						**244453**
Staphylococcus cohnii subsp. urealyticum		99.89	0.96				
Staphylococcus kloosii		0.11	0.48				
244460	**Good identification**						**244460**
Staphylococcus kloosii		100.0	0.34	MAN 99			
244462	**Good identification**						**244462**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.35	MAN 99		
244463	**Good identification**					**244463**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91		
244470	**Very good identification**					**244470**
Staphylococcus kloosii		100.0	0.75			
244471	**Low discrimination**					**244471**
Staphylococcus cohnii subsp. urealyticum		75.43	0.35	XYL 1		
Staphylococcus kloosii		19.54	0.34	LAC 91		
Staphylococcus xylosus		5.03	0.01			
244472	**Very good identification**					**244472**
Staphylococcus kloosii		99.63	0.76			
Staphylococcus cohnii subsp. urealyticum		0.37	0.14			
244473	**Very good identification**					**244473**
Staphylococcus cohnii subsp. urealyticum		95.55	0.55	XYL 1		
Staphylococcus kloosii		2.65	0.35			
Staphylococcus xylosus		1.79	0.11			
244501	**Good identification**					**244501**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91		
244502	**Low discrimination**					**244502**
Staphylococcus kloosii		60.27	0.27	MAN 99		
Staphylococcus cohnii subsp. urealyticum		39.74	0.12	MLT 95		
244503	**Good identification**					**244503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.53	MAN 91		
244510	**Very good identification**					**244510**
Staphylococcus kloosii		99.28	0.67	MLT 95		
Staphylococcus cohnii subsp. urealyticum		0.72	0.12			
244511	**Good identification**					**244511**
Staphylococcus cohnii subsp. urealyticum		98.61	0.53	LAC 91		
Staphylococcus kloosii		1.39	0.26			
244512	**Good identification**					**244512**
Staphylococcus kloosii		93.69	0.68	MLT 95		
Staphylococcus cohnii subsp. urealyticum		6.31	0.32			
244513	**Very good identification**					**244513**
Staphylococcus cohnii subsp. urealyticum		99.85	0.73			
Staphylococcus kloosii		0.15	0.27			
244530	**Very good identification**					**244530**
Staphylococcus kloosii		100.0	0.55	MLT 95		
244532	**Very good identification**					**244532**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.56	MLT 95		
244533	**Low discrimination**					**244533**
Staphylococcus cohnii subsp. urealyticum		84.12	0.32	XYL 1		
Staphylococcus xylosus		15.88	0.08			
244540	**Acceptable identification**					**244540**
					NOV	
Staphylococcus kloosii		94.11	0.53	MAN 99	99	
Staphylococcus warneri		3.91	0.17		1	
Staphylococcus cohnii subsp. urealyticum		1.98	0.06		99	
244541	**Good identification**					**244541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	MAN 91		
244542	**Low discrimination**					**244542**
Staphylococcus kloosii		83.57	0.53	MAN 99		
Staphylococcus cohnii subsp. urealyticum		16.43	0.27			
244543	**Very good identification**					**244543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.68	MAN 91		
244550	**Excellent identification**					**244550**
Staphylococcus kloosii		99.56	0.94			
Staphylococcus cohnii subsp. urealyticum		0.21	0.27			
Staphylococcus saprophyticus subsp. bovis		0.15	0.18			
244551	**Very good identification**					**244551**
Staphylococcus cohnii subsp. urealyticum		95.12	0.68	LAC 91		
Staphylococcus kloosii		4.51	0.53			
Staphylococcus saprophyticus subsp. bovis		0.17	0.06			
244552	**Very good identification**					**244552**
Staphylococcus kloosii		98.00	0.95			
Staphylococcus cohnii subsp. urealyticum		1.97	0.48			
Staphylococcus saprophyticus subsp. bovis		0.03	0.06			
244553	**Very good identification**					**244553**
Staphylococcus cohnii subsp. urealyticum		99.46	0.89			
Staphylococcus kloosii		0.51	0.53			
Staphylococcus xylosus		0.04	0.09			
244560	**Good identification**					**244560**
Staphylococcus kloosii		100.0	0.40	MAN 99		
244562	**Good identification**					**244562**
Staphylococcus kloosii		100.0	0.41	MAN 99		
244563	**Low discrimination**					**244563**
Staphylococcus cohnii subsp. urealyticum		77.78	0.27	MAN 91		
Staphylococcus xylosus		22.22	0.07	XYL 1		
244570	**Very good identification**					**244570**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.81				
244572	**Very good identification**						**244572**
Staphylococcus kloosii		99.71	0.82				
Staphylococcus arlettae		0.21	0.15				
Staphylococcus cohnii subsp. urealyticum		0.08	0.07				
244573	**Low discrimination**						**244573**
Staphylococcus cohnii subsp. urealyticum		62.40	0.48	XYL 1			
Staphylococcus xylosus		27.62	0.31				
Staphylococcus kloosii		7.85	0.41				
244600	**Low discrimination**						**244600**
Staphylococcus warneri		88.02	0.27	PHS 1			
Staphylococcus pasteuri		11.98	-.06	MLT 95			
244603	**Low discrimination**						**244603**
					NOV		
Staphylococcus cohnii subsp. urealyticum		62.77	0.29	SUC 3	99		
Staphylococcus hyicus		26.33	0.09	MAN 91	1		
Staphylococcus simulans		10.91	0.08		1		
244610	**Low discrimination**						**244610**
					NOV		
Staphylococcus kloosii		56.42	0.41	SUC 9	99		
Staphylococcus pasteuri		23.35	0.11	MLT 95	1		
Staphylococcus warneri		20.23	0.24		1		
244611	**Good identification**						**244611**
Staphylococcus cohnii subsp. urealyticum		100.0	0.29	SUC 3			
244612	**Low discrimination**						**244612**
					NOV	YEP	
Staphylococcus kloosii		64.28	0.42	SUC 9	99	99	
Staphylococcus simulans		12.47	0.10	MLT 95	1	1	
Staphylococcus pasteuri		10.03	0.03		1	86	
244613	**Acceptable identification**						**244613**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.57	0.50	SUC 3	99		
Staphylococcus simulans		6.43	0.20		1		
244630	**Good identification**						**244630**
Staphylococcus kloosii		100.0	0.28	SUC 9			
244632	**Good identification**						**244632**
Staphylococcus kloosii		100.0	0.29	SUC 9			
244640	**Good identification**						**244640**
					NOV		
Staphylococcus warneri		93.95	0.53	PHS 1	1		
Staphylococcus pasteuri		4.13	0.11		1		
Staphylococcus kloosii		1.92	0.26		99		
244642	**Low discrimination**						**244642**
					NOV		
Staphylococcus warneri		89.27	0.43	PHS 1	1		
Staphylococcus kloosii		5.92	0.27		99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		4.81	0.03		1		
244643	**Good identification**						**244643**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.57	0.45	SUC 3	99		
Staphylococcus simulans		1.35	0.01	MAN 91	1		
Staphylococcus warneri		1.09	0.09		1		
244650	**Low discrimination**						**244650**
					NOV		
Staphylococcus kloosii		65.74	0.67	SUC 9	99		
Staphylococcus warneri		23.56	0.50		1		
Staphylococcus pasteuri		8.80	0.27		1		
244651	**Low discrimination**						**244651**
Staphylococcus cohnii subsp. urealyticum		89.47	0.45	SUC 3			
Staphylococcus xylosus		3.22	0.06	LAC 91			
Staphylococcus kloosii		3.00	0.26				
244652	**Low discrimination**						**244652**
					NOV		
Staphylococcus kloosii		82.88	0.68	SUC 9	99		
Staphylococcus warneri		9.14	0.40		1		
Staphylococcus pasteuri		4.18	0.19		1		
244653	**Very good identification**						**244653**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.12	0.65	SUC 3	99		
Staphylococcus xylosus		0.99	0.15		99		
Staphylococcus simulans		0.54	0.13		1		
244670	**Very good identification**						**244670**
					NOV		
Staphylococcus kloosii		98.57	0.55	SUC 9	99		
Staphylococcus warneri		1.43	0.09		1		
244671	**Acceptable identification**						**244671**
Staphylococcus xylosus		91.29	0.28	NIT 89			
Staphylococcus equorum		8.71	0.08	GAL 91			
244672	**Low discrimination**						**244672**
Staphylococcus kloosii		57.99	0.56	SUC 9			
Staphylococcus arlettae		41.25	0.41				
Staphylococcus xylosus		0.77	-.04				
244673	**Low discrimination**						**244673**
Staphylococcus xylosus		52.86	0.37	NIT 89			
Staphylococcus arlettae		38.00	0.43	GAL 91			
Staphylococcus cohnii subsp. urealyticum		4.59	0.24				
244703	**Acceptable identification**						**244703**
					NOV	YEP	
Staphylococcus hyicus		90.39	0.31	ARG 96	1	1	
Staphylococcus cohnii subsp. urealyticum		8.03	0.22	NIT 99	99	64	
Staphylococcus chromogenes		1.58	-.02		1	95	
244710	**Acceptable identification**						**244710**
					NOV		
Staphylococcus kloosii		94.49	0.47	SUC 9	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		5.51	0.14	MLT 95	1		
244712	**Good identification**						**244712**
Staphylococcus kloosii		100.0	0.48	SUC 9			
244713	**Low discrimination**						**244713**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		88.92	0.42	SUC 3	1	99	
Staphylococcus xylosus		9.06	0.13		1	99	
Staphylococcus intermedius		2.02	0.01		99	1	
244730	**Good identification**						**244730**
Staphylococcus kloosii		100.0	0.34	SUC 9			
244732	**Low discrimination**						**244732**
Staphylococcus kloosii		71.51	0.35	SUC 9			
Staphylococcus arlettae		28.50	0.15	MLT 95			
244733	**Low discrimination**						**244733**
					YEP		
Staphylococcus xylosus		85.66	0.35	NIT 89	40		
Staphylococcus equorum		9.71	0.16	MLT 93	1		
Staphylococcus arlettae		4.63	0.18		99		
244740	**Low discrimination**						**244740**
					NOV		
Staphylococcus warneri		88.85	0.43	PHS 1	1		
Staphylococcus kloosii		11.15	0.32		99		
244742	**Low discrimination**						**244742**
					NOV		
Staphylococcus warneri		71.04	0.33	PHS 1	1		
Staphylococcus kloosii		28.96	0.33		99		
244743	**Low discrimination**						**244743**
					NOV		
Staphylococcus cohnii subsp. urealyticum		84.11	0.37	SUC 3	99		
Staphylococcus xylosus		12.97	0.11	MAN 91	99		
Staphylococcus chromogenes		2.92	-.02		1		
244750	**Low discrimination**						**244750**
					NOV	PYR	
Staphylococcus kloosii		89.45	0.73	SUC 9	99	52	
Staphylococcus saprophyticus subsp. bovis		5.34	0.31		99	99	
Staphylococcus warneri		5.22	0.40		1	8	
244751	**Low discrimination**						**244751**
Staphylococcus cohnii subsp. urealyticum		44.96	0.37	SUC 3			
Staphylococcus xylosus		38.16	0.26	LAC 91			
Staphylococcus saprophyticus subsp. bovis		10.06	0.18				
244752	**Very good identification**						**244752**
					NOV	PYR	
Staphylococcus kloosii		95.86	0.74	SUC 9	99	52	
Staphylococcus warneri		1.72	0.30		1	8	
Staphylococcus saprophyticus subsp. bovis		1.32	0.18		99	99	
244753	**Low discrimination**						**244753**
Staphylococcus cohnii subsp. urealyticum		78.26	0.58	SUC 3			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		18.69	0.36			
Staphylococcus kloosii		1.27	0.33			
244763	**Acceptable identification**					**244763**
Staphylococcus xylosus		92.17	0.33	NIT 89		
Staphylococcus arlettae		5.88	0.18	MAN 94		
Staphylococcus gallinarum		1.95	-.03			
244770	**Very good identification**					**244770**
Staphylococcus kloosii		95.33	0.61	SUC 9		
Staphylococcus xylosus		2.50	0.07			
Staphylococcus arlettae		2.17	0.15			
244771	**Low discrimination**					**244771**
Staphylococcus xylosus		88.75	0.48	NIT 89		
Staphylococcus equorum		8.47	0.29			
Staphylococcus gallinarum		1.41	0.10			
244772	**Low discrimination**					**244772**
Staphylococcus arlettae		66.01	0.56	URE 1		
Staphylococcus kloosii		31.79	0.61			
Staphylococcus xylosus		2.19	0.17			
244773	**Low discrimination**					**244773**
					YEP	
Staphylococcus xylosus		66.32	0.58	NIT 89	40	
Staphylococcus arlettae		26.72	0.59		99	
Staphylococcus equorum		5.72	0.37		1	
245013	**Good identification**					**245013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
245050	**Low discrimination**					**245050**
					OXI	
Staphylococcus kloosii		78.17	0.47	TRE 99	1	
Macrococcus equipersicus		21.83	0.18		99	
245052	**Good identification**					**245052**
Staphylococcus kloosii		96.58	0.48	TRE 99		
Staphylococcus cohnii subsp. urealyticum		3.42	0.06			
245053	**Good identification**					**245053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	TRE 99		
245070	**Good identification**					**245070**
Staphylococcus kloosii		100.0	0.34	TRE 99		
245072	**Good identification**					**245072**
Staphylococcus kloosii		100.0	0.35	TRE 99		
245112	**Good identification**					**245112**
Staphylococcus kloosii		100.0	0.27	TRE 99		
245150	**Acceptable identification**					**245150**
					OXI	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		94.19	0.53	TRE 99	1	
Macrococcus equipersicus		5.81	0.10		99	
245152	**Very good identification**					**245152**
Staphylococcus kloosii		100.0	0.53	TRE 99		
245153	**Good identification**					**245153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	TRE 99		
245170	**Good identification**					**245170**
Staphylococcus kloosii		100.0	0.40	TRE 99		
245172	**Good identification**					**245172**
Staphylococcus kloosii		100.0	0.41	TRE 99		
245252	**Good identification**					**245252**
Staphylococcus kloosii		100.0	0.27	SUC 9		
245350	**Good identification**					**245350**
Staphylococcus kloosii		100.0	0.32	SUC 9		
245352	**Good identification**					**245352**
Staphylococcus kloosii		100.0	0.33	SUC 9		
245371	**Low discrimination**					**245371**
Staphylococcus equorum		78.38	0.29	NIT 92		
Staphylococcus gallinarum		13.06	0.10	TRE 99		
Staphylococcus xylosus		8.56	0.07			
245373	**Low discrimination**					**245373**
Staphylococcus equorum		75.38	0.37	NIT 92		
Staphylococcus gallinarum		11.96	0.18	TRE 99		
Staphylococcus xylosus		9.11	0.17			
245401	**Good identification**					**245401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	MAN 91		
245403	**Good identification**					**245403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	MAN 91		
245410	**Very good identification**					**245410**
					OXI	
Staphylococcus kloosii		96.85	0.62	MLT 95	1	
Macrococcus equipersicus		1.92	0.09		99	
Staphylococcus cohnii subsp. urealyticum		1.23	0.11		1	
245411	**Good identification**					**245411**
Staphylococcus cohnii subsp. urealyticum		99.20	0.52	LAC 91		
Staphylococcus kloosii		0.80	0.20			
245412	**Low discrimination**					**245412**
Staphylococcus kloosii		89.40	0.62	MLT 95		
Staphylococcus cohnii subsp. urealyticum		10.60	0.31			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
245413	**Very good identification**					**245413**
Staphylococcus cohnii subsp. urealyticum		99.91	0.72			
Staphylococcus kloosii		0.09	0.21			
245430	**Good identification**					**245430**
Staphylococcus kloosii		100.0	0.49	MLT 95		
245432	**Good identification**					**245432**
Staphylococcus kloosii		100.0	0.50	MLT 95		
245433	**Good identification**					**245433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	XYL 1		
245440	**Good identification**					**245440**
Staphylococcus kloosii		96.42	0.47	MAN 99		
Staphylococcus cohnii subsp. urealyticum		3.58	0.06			
245441	**Good identification**					**245441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	MAN 91		
245442	**Low discrimination**					**245442**
Staphylococcus kloosii		74.27	0.48	MAN 99		
Staphylococcus cohnii subsp. urealyticum		25.73	0.26			
245443	**Very good identification**					**245443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.67	MAN 91		
245450	**Very good identification**					**245450**
					OXI	
Staphylococcus kloosii		99.34	0.88		1	
Staphylococcus cohnii subsp. urealyticum		0.38	0.26		1	
Macrococcus equipersicus		0.28	0.18		99	
245451	**Very good identification**					**245451**
Staphylococcus cohnii subsp. urealyticum		97.38	0.67	LAC 91		
Staphylococcus kloosii		2.62	0.47			
245452	**Very good identification**					**245452**
Staphylococcus kloosii		96.58	0.89			
Staphylococcus cohnii subsp. urealyticum		3.42	0.47			
245453	**Very good identification**					**245453**
Staphylococcus cohnii subsp. urealyticum		99.71	0.88			
Staphylococcus kloosii		0.29	0.48			
245460	**Good identification**					**245460**
Staphylococcus kloosii		100.0	0.34	MAN 99		
245462	**Good identification**					**245462**
Staphylococcus kloosii		100.0	0.35	MAN 99		
245470	**Very good identification**					**245470**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.75			
245472	**Very good identification**					**245472**
Staphylococcus kloosii		99.86	0.76			
Staphylococcus cohnii subsp. urealyticum		0.14	0.06			
245473	**Low discrimination**					**245473**
					YEP	
Staphylococcus cohnii subsp. urealyticum		86.02	0.47	XYL 1	64	
Staphylococcus equorum		6.80	0.16		1	
Staphylococcus kloosii		6.14	0.35		99	
245502	**Good identification**					**245502**
Staphylococcus kloosii		100.0	0.27	MAN 99		
245503	**Good identification**					**245503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	MAN 91		
245510	**Very good identification**					**245510**
					OXI	
Staphylococcus kloosii		99.56	0.67	MLT 95	1	
Micrococcus equipersicus		0.44	0.01		99	
245511	**Good identification**					**245511**
Staphylococcus cohnii subsp. urealyticum		96.49	0.44	LAC 91		
Staphylococcus kloosii		3.51	0.26			
245512	**Very good identification**					**245512**
Staphylococcus kloosii		97.45	0.68	MLT 95		
Staphylococcus cohnii subsp. urealyticum		2.55	0.24			
245513	**Very good identification**					**245513**
Staphylococcus cohnii subsp. urealyticum		99.61	0.65			
Staphylococcus kloosii		0.39	0.27			
245530	**Very good identification**					**245530**
Staphylococcus kloosii		100.0	0.55	MLT 95		
245532	**Very good identification**					**245532**
Staphylococcus kloosii		100.0	0.56	MLT 95		
245540	**Good identification**					**245540**
Staphylococcus kloosii		100.0	0.53	MAN 99		
245541	**Good identification**					**245541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	MAN 91		
245542	**Good identification**					**245542**
Staphylococcus kloosii		92.90	0.53	MAN 99		
Staphylococcus cohnii subsp. urealyticum		7.10	0.19			
245543	**Very good identification**					**245543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.60	MAN 91		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
245550	**Excellent identification**					**245550**
					OXI	
Staphylococcus kloosii		99.85	0.94		1	
Staphylococcus cohnii subsp. urealyticum		0.08	0.19		1	
Macrococcus equipersicus		0.06	0.10		99	
245551	**Low discrimination**					**245551**
Staphylococcus cohnii subsp. urealyticum		89.13	0.60	LAC 91		
Staphylococcus kloosii		10.87	0.53			
245552	**Excellent identification**					**245552**
Staphylococcus kloosii		99.23	0.95			
Staphylococcus cohnii subsp. urealyticum		0.78	0.39			
245553	**Very good identification**					**245553**
Staphylococcus cohnii subsp. urealyticum		98.69	0.80			
Staphylococcus kloosii		1.29	0.53			
Staphylococcus xylosus		0.03	-.03			
245560	**Good identification**					**245560**
Staphylococcus kloosii		100.0	0.40	MAN 99		
245562	**Good identification**					**245562**
Staphylococcus kloosii		100.0	0.41	MAN 99		
245570	**Very good identification**					**245570**
Staphylococcus kloosii		100.0	0.81			
245571	**Low discrimination**					**245571**
					YEP	
Staphylococcus equorum		53.98	0.29	NIT 92	1	
Staphylococcus kloosii		22.22	0.40	SUC 99	99	
Staphylococcus gallinarum		9.00	0.10		78	
245572	**Very good identification**					**245572**
Staphylococcus kloosii		99.79	0.82			
Staphylococcus arlettae		0.21	0.15			
245573	**Low discrimination**					**245573**
					YEP	
Staphylococcus equorum		48.36	0.37	NIT 92	1	
Staphylococcus cohnii subsp. urealyticum		25.93	0.39	SUC 99	64	
Staphylococcus kloosii		8.38	0.41		99	
245610	**Good identification**					**245610**
Staphylococcus kloosii		100.0	0.41	SUC 9		
245612	**Good identification**					**245612**
Staphylococcus kloosii		100.0	0.42	SUC 9		
245613	**Good identification**					**245613**
Staphylococcus cohnii subsp. urealyticum		100.0	0.41	SUC 3		
245630	**Good identification**					**245630**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Staphylococcus kloosii	100.0	0.28	SUC 9		
245631	**Good identification**					**245631**
	Staphylococcus equorum	100.0	0.28	NIT 92		
245632	**Good identification**					**245632**
	Staphylococcus kloosii	100.0	0.29	SUC 9		
245633	**Good identification**					**245633**
	Staphylococcus equorum	97.82	0.37	NIT 92		
	Staphylococcus xylosus	2.18	0.01	GAL 91		
245642	**Good identification**					**245642**
	Staphylococcus kloosii	100.0	0.27	SUC 9		
245643	**Good identification**					**245643**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.36	SUC 3		
245650	**Very good identification**					**245650**
					NOV	
	Staphylococcus kloosii	99.64	0.67	SUC 9	99	
	Staphylococcus warneri	0.36	0.09		1	
245651	**Low discrimination**					**245651**
					YEP	
	Staphylococcus cohnii subsp. urealyticum	81.64	0.36	SUC 3	64	
	Staphylococcus equorum	8.29	0.08	LAC 91	1	
	Staphylococcus kloosii	7.03	0.26		99	
245652	**Very good identification**					**245652**
	Staphylococcus kloosii	98.91	0.68	SUC 9		
	Staphylococcus cohnii subsp. urealyticum	1.09	0.16			
245653	**Very good identification**					**245653**
					YEP	
	Staphylococcus cohnii subsp. urealyticum	96.01	0.57	SUC 3	64	
	Staphylococcus equorum	2.48	0.16		1	
	Staphylococcus kloosii	0.89	0.27		99	
245670	**Very good identification**					**245670**
					YEP	
	Staphylococcus kloosii	95.45	0.55	SUC 9	99	
	Staphylococcus equorum	4.55	0.08		1	
245671	**Good identification**					**245671**
	Staphylococcus equorum	95.83	0.49	NIT 92		
	Staphylococcus xylosus	2.54	0.15	GAL 91		
	Staphylococcus gallinarum	1.63	0.10			
245672	**Low discrimination**					**245672**
					YEP	
	Staphylococcus kloosii	54.81	0.56	SUC 9	99	
	Staphylococcus arlettae	38.99	0.41		99	
	Staphylococcus equorum	6.20	0.16		1	
245673	**Low discrimination**					**245673**
					YEP	
	Staphylococcus equorum	88.21	0.58	NIT 92	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus arlettae		7.43	0.43	GAL 91	99	
Staphylococcus xylosus		2.58	0.25		40	
245710	**Good identification**					**245710**
Staphylococcus kloosii		100.0	0.47	SUC 9		
245712	**Good identification**					**245712**
Staphylococcus kloosii		100.0	0.48	SUC 9		
245713	**Low discrimination**					**245713**
Staphylococcus cohnii subsp. urealyticum		71.08	0.34	SUC 3		
Staphylococcus equorum		24.26	0.16			
Staphylococcus xylosus		4.66	0.00			
245730	**Low discrimination**					**245730**
					YEP	
Staphylococcus kloosii		68.20	0.34	SUC 9	99	
Staphylococcus equorum		31.80	0.08	MLT 95	1	
245731	**Good identification**					**245731**
Staphylococcus equorum		96.42	0.49	NIT 92		
Staphylococcus xylosus		1.94	0.13	MLT 91		
Staphylococcus gallinarum		1.64	0.10			
245733	**Very good identification**					**245733**
Staphylococcus equorum		95.85	0.58	NIT 92		
Staphylococcus xylosus		2.14	0.22	MLT 91		
Staphylococcus gallinarum		1.55	0.18			
245740	**Good identification**					**245740**
Staphylococcus kloosii		100.0	0.32	SUC 9		
245742	**Good identification**					**245742**
Staphylococcus kloosii		100.0	0.33	SUC 9		
245743	**Low discrimination**					**245743**
Staphylococcus cohnii subsp. urealyticum		83.63	0.29	SUC 3		
Staphylococcus xylosus		8.29	-.01	MAN 91		
Staphylococcus gallinarum		8.08	-.03			
245750	**Very good identification**					**245750**
Staphylococcus kloosii		100.0	0.73	SUC 9		
245751	**Low discrimination**					**245751**
Staphylococcus equorum		48.89	0.29	NIT 92		
Staphylococcus cohnii subsp. urealyticum		20.41	0.29	XYL 99		
Staphylococcus xylosus		11.14	0.14			
245752	**Very good identification**					**245752**
Staphylococcus kloosii		99.24	0.74	SUC 9		
Staphylococcus arlettae		0.52	0.15			
Staphylococcus cohnii subsp. urealyticum		0.24	0.08			
245753	**Low discrimination**					**245753**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		51.82	0.49	SUC 3		
Staphylococcus equorum		31.56	0.37			
Staphylococcus xylosus		7.96	0.23			
245761	**Low discrimination**					**245761**
Staphylococcus gallinarum		58.29	0.30	NIT 99		
Staphylococcus equorum		35.73	0.29	MAN 91		
Staphylococcus xylosus		5.98	0.11			
245763	**Low discrimination**					**245763**
Staphylococcus gallinarum		55.76	0.38	NIT 99		
Staphylococcus equorum		35.90	0.37	MAN 91		
Staphylococcus xylosus		6.64	0.21			
245770	**Low discrimination**					**245770**
					YEP	
Staphylococcus kloosii		76.21	0.61	SUC 9	99	
Staphylococcus equorum		18.91	0.29		1	
Staphylococcus gallinarum		3.15	0.10		78	
245771	**Low discrimination**					**245771**
Staphylococcus equorum		83.66	0.70	NIT 92		
Staphylococcus gallinarum		13.94	0.51			
Staphylococcus xylosus		2.21	0.36			
245772	**Low discrimination**					**245772**
					YEP	
Staphylococcus arlettae		54.99	0.56	URE 1	99	
Staphylococcus kloosii		26.48	0.61		99	
Staphylococcus equorum		15.60	0.37		1	
245773	**Low discrimination**					**245773**
					YEP	
Staphylococcus equorum		80.94	0.78	NIT 92	1	
Staphylococcus gallinarum		12.84	0.59		78	
Staphylococcus arlettae		3.82	0.59		99	
246000	**Low discrimination**					**246000**
					OXI	
Dermacoccus nishinomiyaensis		77.55	0.28	GLR 1	93	
Kocuria rosea		15.00	0.12	PHS 5	35	
Kocuria varians		7.45	0.14		13	
246053	**Good identification**					**246053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	TRE 99		
246153	**Acceptable identification**					**246153**
Staphylococcus cohnii subsp. urealyticum		93.15	0.32	TRE 99		
Staphylococcus xylosus		6.85	0.00			
246203	**Acceptable identification**					**246203**
					YEP	
Staphylococcus hyicus		92.32	0.28	ARG 96	1	
Staphylococcus simulans		5.61	0.10	GAL 92	1	
Staphylococcus chromogenes		2.06	-.02	TRE 92	95	
246273	**Good identification**					**246273**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		97.12	0.27	GAL 91		
Staphylococcus equorum		2.89	-.03	TRE 96		
246301	**Good identification**					**246301**
					YEP	
Staphylococcus hyicus		97.81	0.28	ARG 96	1	
Staphylococcus chromogenes		2.19	-.02	TRE 92	95	
246303	**Good identification**					**246303**
					YEP	
Staphylococcus hyicus		98.28	0.50	ARG 96	1	
Staphylococcus chromogenes		1.72	0.17	TRE 92	95	
246343	**Low discrimination**					**246343**
					YEP	
Staphylococcus epidermidis		57.85	0.29	VPT 96	1	
Staphylococcus chromogenes		24.07	0.17	GLR 1	95	
Staphylococcus hyicus		13.90	0.09		1	
246350	**Good identification**					**246350**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	TRE 99		
246371	**Acceptable identification**					**246371**
Staphylococcus xylosus		92.59	0.39	TRE 96		
Staphylococcus gallinarum		4.37	0.10			
Staphylococcus equorum		3.05	0.09			
246373	**Acceptable identification**					**246373**
Staphylococcus xylosus		93.43	0.48	TRE 96		
Staphylococcus gallinarum		3.79	0.18			
Staphylococcus equorum		2.78	0.18			
246403	**Acceptable identification**					**246403**
					NOV	
Staphylococcus cohnii subsp. urealyticum		90.58	0.45	MAN 91	99	
Staphylococcus hyicus		6.66	0.09		1	
Staphylococcus simulans		2.76	0.08		1	
246411	**Good identification**					**246411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.45	LAC 91		
246413	**Very good identification**					**246413**
					NOV	
Staphylococcus cohnii subsp. urealyticum		98.27	0.65		99	
Staphylococcus simulans		1.18	0.20		1	
Staphylococcus felis		0.54	0.04		1	
246441	**Good identification**					**246441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	MAN 91		
246443	**Very good identification**					**246443**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.76	0.60	MAN 91	99	
Staphylococcus simulans		0.24	0.01		1	
246450	**Low discrimination**					**246450**
Staphylococcus saprophyticus subsp. bovis		84.88	0.47			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		12.10	0.47			
Staphylococcus cohnii subsp. urealyticum		2.06	0.19			
246451	**Good identification**					**246451**
Staphylococcus cohnii subsp. urealyticum		90.36	0.60	LAC 91		
Staphylococcus saprophyticus subsp. bovis		9.40	0.34			
Staphylococcus xylosus		0.24	-.02			
246452	**Low discrimination**					**246452**
Staphylococcus saprophyticus subsp. bovis		38.06	0.34			
Staphylococcus cohnii subsp. urealyticum		37.37	0.40			
Staphylococcus kloosii		23.51	0.48			
246453	**Very good identification**					**246453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.57	0.81		99	
Staphylococcus saprophyticus subsp. bovis		0.26	0.22		99	
Staphylococcus simulans		0.10	0.13		1	
246470	**Low discrimination**					**246470**
Staphylococcus kloosii		77.92	0.34	NIT 1		
Staphylococcus saprophyticus subsp. bovis		22.08	0.06			
246472	**Good identification**					**246472**
Staphylococcus kloosii		100.0	0.35	NIT 1		
246473	**Low discrimination**					**246473**
Staphylococcus cohnii subsp. urealyticum		53.18	0.40	XYL 1		
Staphylococcus xylosus		45.75	0.29			
Staphylococcus equorum		1.08	-.03			
246503	**Low discrimination**					**246503**
					NOV	
Staphylococcus hyicus		64.20	0.31	ARG 96	1	
Staphylococcus cohnii subsp. urealyticum		32.55	0.37	SUC 99	99	
Staphylococcus felis		2.12	-.02		1	
246511	**Acceptable identification**					**246511**
Staphylococcus cohnii subsp. urealyticum		92.37	0.37	LAC 91		
Staphylococcus saprophyticus subsp. bovis		5.13	0.06			
Staphylococcus xylosus		2.50	-.04			
246513	**Very good identification**					**246513**
					NOV	
Staphylococcus cohnii subsp. urealyticum		95.17	0.58		99	
Staphylococcus felis		4.11	0.15		1	
Staphylococcus xylosus		0.72	0.05		99	
246533	**Low discrimination**					**246533**
Staphylococcus xylosus		87.23	0.27	SUC 95		
Staphylococcus cohnii subsp. urealyticum		10.08	0.17	MLT 93		
Staphylococcus equorum		2.70	-.03			
246541	**Acceptable identification**					**246541**
Staphylococcus cohnii subsp. urealyticum		90.98	0.32	MAN 91		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		9.02	0.06	LAC 91			
246543	**Good identification**						**246543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.27	0.53	MAN 91	99		
Staphylococcus xylosus		1.13	0.04		99		
Staphylococcus chromogenes		0.60	-.02		1		
246550	**Good identification**						**246550**
Staphylococcus saprophyticus subsp. bovis		93.30	0.59				
Staphylococcus kloosii		6.46	0.53				
Staphylococcus cohnii subsp. urealyticum		0.24	0.12				
246551	**Low discrimination**						**246551**
Staphylococcus cohnii subsp. urealyticum		49.16	0.53	LAC 91			
Staphylococcus saprophyticus subsp. bovis		47.73	0.47				
Staphylococcus xylosus		3.11	0.19				
246552	**Low discrimination**						**246552**
Staphylococcus saprophyticus subsp. bovis		71.17	0.47				
Staphylococcus kloosii		21.34	0.53				
Staphylococcus cohnii subsp. urealyticum		7.49	0.32				
246553	**Very good identification**						**246553**
Staphylococcus cohnii subsp. urealyticum		95.99	0.73				
Staphylococcus saprophyticus subsp. bovis		2.30	0.34				
Staphylococcus xylosus		1.71	0.28				
246570	**Low discrimination**						**246570**
Staphylococcus kloosii		59.05	0.40	NIT 1			
Staphylococcus saprophyticus subsp. bovis		34.48	0.18				
Staphylococcus xylosus		6.47	-.01				
246571	**Acceptable identification**						**246571**
Staphylococcus xylosus		91.68	0.41	SUC 95			
Staphylococcus gallinarum		3.42	0.10				
Staphylococcus equorum		2.39	0.09				
246572	**Low discrimination**						**246572**
Staphylococcus kloosii		70.30	0.41	NIT 1			
Staphylococcus xylosus		20.23	0.09				
Staphylococcus saprophyticus subsp. bovis		9.47	0.06				
246573	**Low discrimination**						**246573**
					COA	NOV	
Staphylococcus xylosus		86.94	0.50	SUC 95	1	99	
Staphylococcus cohnii subsp. urealyticum		4.28	0.32		1	99	
Staphylococcus lutrae		3.94	0.18		99	1	
246601	**Good identification**						**246601**
					YEP		
Staphylococcus hyicus		97.22	0.28	ARG 96	1		
Staphylococcus chromogenes		2.78	0.00	GAL 92	95		
246603	**Low discrimination**						**246603**
					YEP		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hyicus		88.54	0.50	ARG 96	1			
Staphylococcus simulans		8.90	0.36	GAL 92	1			
Staphylococcus chromogenes		1.98	0.20		95			
246612	**Acceptable identification**							**246612**
					NOV	YEP	PYR	
Staphylococcus simulans		90.14	0.38	ARG 92	1	1	70	
Staphylococcus pasteuri		7.40	0.11	BGA 90	1	86	1	
Staphylococcus saprophyticus subsp. bovis		2.46	0.06		99	80	99	
246613	**Low discrimination**							**246613**
					NOV			
Staphylococcus simulans		82.80	0.49	ARG 92	1			
Staphylococcus cohnii subsp. urealyticum		8.85	0.34	BGA 90	99			
Staphylococcus felis		3.21	0.11		1			
246620	**Good identification**							**246620**
Staphylococcus muscae		100.0	0.39	URE 1				
246633	**Low discrimination**							**246633**
					NOV			
Staphylococcus xylosus		83.16	0.33	GAL 91	99			
Staphylococcus equorum		13.39	0.18	MLT 93	99			
Staphylococcus simulans		3.45	0.08		1			
246640	**Low discrimination**							**246640**
Staphylococcus warneri		63.43	0.46	PHS 1				
Staphylococcus pasteuri		15.95	0.19					
Staphylococcus hominis subsp. hominis		11.97	0.25					
246642	**Low discrimination**							**246642**
Staphylococcus warneri		40.06	0.36	PHS 1				
Staphylococcus hominis subsp. hominis		26.62	0.26					
Staphylococcus simulans		16.89	0.18					
246643	**Low discrimination**							**246643**
					NOV	YEP		
Staphylococcus simulans		41.45	0.29	ARG 92	1	1		
Staphylococcus cohnii subsp. urealyticum		22.12	0.29	BGA 90	99	64		
Staphylococcus chromogenes		20.53	0.20		1	95		
246650	**Low discrimination**							**246650**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		85.21	0.59		99	99		
Staphylococcus pasteuri		9.74	0.35		1	1		
Staphylococcus warneri		4.57	0.43		1	8		
246651	**Low discrimination**							**246651**
Staphylococcus saprophyticus subsp. bovis		83.16	0.47					
Staphylococcus xylosus		10.19	0.24					
Staphylococcus cohnii subsp. urealyticum		6.18	0.29					
246652	**Low discrimination**							**246652**
					NOV	YEP	PYR	
Staphylococcus saprophyticus subsp. bovis		68.62	0.47		99	80	99	
Staphylococcus pasteuri		12.82	0.27		1	86	1	
Staphylococcus simulans		11.42	0.31		1	1	70	
246653	**Low discrimination**							**246653**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV			
Staphylococcus cohnii subsp. urealyticum		38.90	0.50	SUC 3	99			
Staphylococcus simulans		28.84	0.42		1			
Staphylococcus xylosus		18.06	0.34		99			
246663	**Good identification**							**246663**
Staphylococcus xylosus		95.61	0.31	GAL 91				
Staphylococcus gallinarum		2.53	-.03	MAN 94				
Staphylococcus equorum		1.85	-.03					
246671	**Low discrimination**							**246671**
Staphylococcus xylosus		86.01	0.46	GAL 91				
Staphylococcus equorum		11.67	0.30					
Staphylococcus gallinarum		1.71	0.10					
246673	**Low discrimination**							**246673**
Staphylococcus xylosus		87.49	0.56	GAL 91				
Staphylococcus equorum		10.72	0.38					
Staphylococcus gallinarum		1.50	0.18					
246701	**Good identification**							**246701**
					NOV	YEP		
Staphylococcus hyicus		97.45	0.50	ARG 96	1	1		
Staphylococcus chromogenes		2.18	0.20	LAC 92	1	95		
Staphylococcus xylosus		0.19	-.03		99	40		
246702	**Good identification**							**246702**
					YEP			
Staphylococcus hyicus		96.16	0.31	ARG 96	1			
Staphylococcus simulans		2.16	0.04	MNS 99	1			
Staphylococcus chromogenes		1.68	-.02		95			
246703	**Very good identification**							**246703**
					YEP	PYR	COA	
Staphylococcus hyicus		97.98	0.72	ARG 96	1	1	24	
Staphylococcus chromogenes		1.71	0.39		95	45	1	
Staphylococcus intermedius		0.10	0.15		1	90	99	
246710	**Good identification**							**246710**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	MLT 99				
246713	**Low discrimination**							**246713**
					NOV	YEP		
Staphylococcus hyicus		28.97	0.31	ARG 96	1	1		
Staphylococcus xylosus		21.46	0.31	MAN 1	99	40		
Staphylococcus chromogenes		14.15	0.28		1	95		
246723	**Low discrimination**							**246723**
					NOV			
Staphylococcus hyicus		62.74	0.31	ARG 96	1			
Staphylococcus xylosus		34.11	0.29	XYL 1	99			
Staphylococcus gallinarum		1.19	-.03		99			
246731	**Low discrimination**							**246731**
Staphylococcus xylosus		83.03	0.44	MLT 93				
Staphylococcus equorum		14.80	0.30					
Staphylococcus gallinarum		2.17	0.10					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
246733	**Low discrimination**						**246733**
Staphylococcus xylosus		84.50	0.53	MLT 93			
Staphylococcus equorum		13.61	0.38				
Staphylococcus gallinarum		1.90	0.18				
246740	**Low discrimination**						**246740**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		51.20	0.31	MAN 99	99	99	
Staphylococcus warneri		29.63	0.35		1	8	
Staphylococcus hominis subsp. hominis		19.18	0.26		1	1	
246742	**Low discrimination**						**246742**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		47.73	0.27	GLR 1	1	1	
Staphylococcus saprophyticus subsp. bovis		27.13	0.18	PHS 3	99	99	
Staphylococcus warneri		20.93	0.25		1	8	
246743	**Low discrimination**						**246743**
					NOV	YEP	
Staphylococcus chromogenes		47.05	0.39	ARG 90	1	95	
Staphylococcus hyicus		27.17	0.31	GLR 1	1	1	
Staphylococcus xylosus		17.07	0.30		99	40	
246750	**Very good identification**						**246750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		99.28	0.72		99	99	
Staphylococcus warneri		0.42	0.32		1	8	
Staphylococcus kloosii		0.17	0.32		99	52	
246751	**Low discrimination**						**246751**
Staphylococcus saprophyticus subsp. bovis		74.52	0.59				
Staphylococcus xylosus		23.09	0.45				
Staphylococcus cohnii subsp. urealyticum		0.59	0.22				
246752	**Very good identification**						**246752**
Staphylococcus saprophyticus subsp. bovis		97.25	0.59				
Staphylococcus xylosus		0.87	0.13				
Staphylococcus kloosii		0.72	0.33				
246753	**Low discrimination**						**246753**
Staphylococcus xylosus		65.67	0.55	XYL 92			
Staphylococcus saprophyticus subsp. bovis		18.61	0.47				
Staphylococcus cohnii subsp. urealyticum		6.00	0.42				
246761	**Low discrimination**						**246761**
Staphylococcus xylosus		75.59	0.42	MAN 94			
Staphylococcus gallinarum		22.79	0.30				
Staphylococcus equorum		1.62	0.09				
246763	**Low discrimination**						**246763**
Staphylococcus xylosus		76.90	0.52	MAN 94			
Staphylococcus gallinarum		19.94	0.38				
Staphylococcus equorum		1.49	0.18				
246770	**Low discrimination**						**246770**
					PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus saprophyticus subsp. bovis		44.90	0.31	XYL 1	99	
Staphylococcus xylosus		40.00	0.26		65	
Staphylococcus gallinarum		7.78	0.10		12	
246771	**Low discrimination**					**246771**
Staphylococcus xylosus		75.02	0.67			
Staphylococcus gallinarum		14.60	0.51			
Staphylococcus equorum		10.18	0.51			
246772	**Low discrimination**					**246772**
Staphylococcus xylosus		69.24	0.35	MNS 99		
Staphylococcus gallinarum		11.59	0.18			
Staphylococcus equorum		8.49	0.18			
246773	**Low discrimination**					**246773**
Staphylococcus xylosus		77.34	0.76			
Staphylococcus gallinarum		12.94	0.59			
Staphylococcus equorum		9.48	0.59			
247053	**Good identification**					**247053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
247271	**Low discrimination**					**247271**
Staphylococcus equorum		79.37	0.30	GAL 91		
Staphylococcus gallinarum		11.63	0.10	TRE 99		
Staphylococcus xylosus		6.10	0.06			
247273	**Low discrimination**					**247273**
Staphylococcus equorum		81.67	0.38	GAL 91		
Staphylococcus gallinarum		11.39	0.18	TRE 99		
Staphylococcus xylosus		6.94	0.15			
247331	**Low discrimination**					**247331**
Staphylococcus equorum		82.99	0.30	TRE 99		
Staphylococcus gallinarum		12.16	0.10	MLT 91		
Staphylococcus xylosus		4.85	0.03			
247333	**Low discrimination**					**247333**
Staphylococcus equorum		83.05	0.38	TRE 99		
Staphylococcus gallinarum		11.58	0.18	MLT 91		
Staphylococcus xylosus		5.37	0.12			
247361	**Acceptable identification**					**247361**
Staphylococcus gallinarum		90.44	0.30	TRE 99		
Staphylococcus equorum		6.44	0.09	MAN 91		
Staphylococcus xylosus		3.13	0.02			
247363	**Low discrimination**					**247363**
Staphylococcus gallinarum		89.69	0.38	TRE 99		
Staphylococcus equorum		6.71	0.18	MAN 91		
Staphylococcus xylosus		3.60	0.11			
247371	**Low discrimination**					**247371**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus gallinarum		56.96	0.51	TRE 99		
Staphylococcus equorum		39.71	0.51			
Staphylococcus xylosus		3.05	0.26			
247373	**Low discrimination**					**247373**
Staphylococcus gallinarum		55.73	0.59	TRE 99		
Staphylococcus equorum		40.81	0.59			
Staphylococcus xylosus		3.47	0.36			
247403	**Good identification**					**247403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91		
247411	**Good identification**					**247411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	LAC 91		
247413	**Very good identification**					**247413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.57			
247441	**Good identification**					**247441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	MAN 91		
247443	**Good identification**					**247443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	MAN 91		
247450	**Low discrimination**					**247450**
Staphylococcus kloosii		87.95	0.47	NIT 1		
Staphylococcus saprophyticus subsp. bovis		6.23	0.06			
Staphylococcus cohnii subsp. urealyticum		5.82	0.11			
247451	**Good identification**					**247451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	LAC 91		
247452	**Low discrimination**					**247452**
Staphylococcus kloosii		61.80	0.48	NIT 1		
Staphylococcus cohnii subsp. urealyticum		38.20	0.31			
247453	**Very good identification**					**247453**
Staphylococcus cohnii subsp. urealyticum		99.95	0.72			
Staphylococcus equorum		0.05	-.03			
247470	**Good identification**					**247470**
Staphylococcus kloosii		100.0	0.34	NIT 1		
247471	**Low discrimination**					**247471**
Staphylococcus equorum		77.35	0.30	GAL 91		
Staphylococcus gallinarum		11.33	0.10	SUC 99		
Staphylococcus xylosus		7.50	0.08			
247472	**Low discrimination**					**247472**
					YEP	
Staphylococcus kloosii		88.60	0.35	NIT 1	99	
Staphylococcus equorum		11.40	-.03		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
247473	**Low discrimination**					**247473**
	Staphylococcus equorum	69.40	0.38	GAL 91		
	Staphylococcus cohnii subsp. urealyticum	13.47	0.31	SUC 99		
	Staphylococcus gallinarum	9.68	0.18			
247503	**Good identification**					**247503**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.29	MAN 91		
247511	**Good identification**					**247511**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.29	LAC 91		
247512	**Low discrimination**					**247512**
	Staphylococcus kloosii	68.61	0.27	NIT 1		
	Staphylococcus cohnii subsp. urealyticum	31.39	0.08	MLT 95		
247513	**Good identification**					**247513**
	Staphylococcus cohnii subsp. urealyticum	99.31	0.49			
	Staphylococcus equorum	0.69	-.03			
247531	**Low discrimination**					**247531**
	Staphylococcus equorum	81.95	0.30	SUC 99		
	Staphylococcus gallinarum	12.01	0.10	MLT 91		
	Staphylococcus xylosus	6.05	0.05			
247533	**Low discrimination**					**247533**
	Staphylococcus equorum	80.91	0.38	SUC 99		
	Staphylococcus gallinarum	11.29	0.18	MLT 91		
	Staphylococcus xylosus	6.61	0.15			
247543	**Good identification**					**247543**
	Staphylococcus cohnii subsp. urealyticum	98.34	0.44	MAN 91		
	Staphylococcus gallinarum	1.67	-.03			
247550	**Low discrimination**					**247550**
	Staphylococcus kloosii	87.26	0.53	NIT 1		
	Staphylococcus saprophyticus subsp. bovis	12.74	0.18			
247551	**Low discrimination**					**247551**
	Staphylococcus cohnii subsp. urealyticum	83.38	0.44	LAC 91		
	Staphylococcus gallinarum	5.83	0.10			
	Staphylococcus equorum	4.07	0.09			
247552	**Low discrimination**					**247552**
	Staphylococcus kloosii	85.46	0.53	NIT 1		
	Staphylococcus cohnii subsp. urealyticum	11.66	0.24			
	Staphylococcus saprophyticus subsp. bovis	2.88	0.06			
247553	**Very good identification**					**247553**
	Staphylococcus cohnii subsp. urealyticum	96.08	0.65			
	Staphylococcus gallinarum	1.63	0.18			
	Staphylococcus equorum	1.19	0.18			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
247561	**Low discrimination**						**247561**
Staphylococcus gallinarum		89.70	0.30	SUC 99			
Staphylococcus equorum		6.39	0.09	MAN 91			
Staphylococcus xylosus		3.92	0.04				
247563	**Low discrimination**						**247563**
Staphylococcus gallinarum		88.85	0.38	SUC 99			
Staphylococcus equorum		6.64	0.18	MAN 91			
Staphylococcus xylosus		4.51	0.13				
247570	**Low discrimination**						**247570**
					YEP		
Staphylococcus kloosii		59.27	0.40	NIT 1	99		
Staphylococcus gallinarum		24.00	0.10		78		
Staphylococcus equorum		16.73	0.09		1		
247571	**Low discrimination**						**247571**
Staphylococcus gallinarum		56.66	0.51	SUC 99			
Staphylococcus equorum		39.50	0.51				
Staphylococcus xylosus		3.83	0.28				
247572	**Low discrimination**						**247572**
					YEP		
Staphylococcus kloosii		37.64	0.41	NIT 1	99		
Staphylococcus gallinarum		34.44	0.18		78		
Staphylococcus equorum		25.22	0.18		1		
247573	**Low discrimination**						**247573**
Staphylococcus gallinarum		55.04	0.59	SUC 99			
Staphylococcus equorum		40.30	0.59				
Staphylococcus xylosus		4.33	0.38				
247631	**Acceptable identification**						**247631**
					OXI		
Staphylococcus equorum		94.23	0.50	GAL 91	1		
Staphylococcus sciuri		3.03	0.22	MLT 91	99		
Staphylococcus gallinarum		1.41	0.10		1		
247633	**Very good identification**						**247633**
Staphylococcus equorum		96.90	0.59	GAL 91			
Staphylococcus xylosus		1.52	0.20	MLT 91			
Staphylococcus gallinarum		1.38	0.18				
247651	**Low discrimination**						**247651**
					OXI		
Staphylococcus sciuri		50.92	0.37	URE 1	99		
Staphylococcus equorum		32.49	0.30	GLR 14	1		
Staphylococcus xylosus		5.21	0.12		1		
247652	**Low discrimination**						**247652**
					OXI	PYR	
Staphylococcus kloosii		42.00	0.27	NIT 1	1	52	
Staphylococcus saprophyticus subsp. bovis		27.77	0.06	SUC 9	1	99	
Staphylococcus sciuri		16.58	0.02		99	1	
247653	**Low discrimination**						**247653**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus equorum		49.46	0.38	GAL 91			
Staphylococcus cohnii subsp. urealyticum		29.40	0.41	XYL 99			
Staphylococcus xylosus		8.77	0.21				
247661	**Low discrimination**						**247661**
Staphylococcus gallinarum		55.41	0.30	GAL 99			
Staphylococcus equorum		38.63	0.30	MAN 91			
Staphylococcus xylosus		4.55	0.09				
247663	**Low discrimination**						**247663**
Staphylococcus gallinarum		54.72	0.38	GAL 99			
Staphylococcus equorum		40.07	0.38	MAN 91			
Staphylococcus xylosus		5.22	0.19				
247670	**Low discrimination**						**247670**
					OXI		
Staphylococcus equorum		77.36	0.30	GAL 91	1		
Staphylococcus gallinarum		11.33	0.10	MNS 99	1		
Staphylococcus sciuri		11.31	0.16		99		
247671	**Low discrimination**						**247671**
Staphylococcus equorum		84.96	0.71	GAL 91			
Staphylococcus gallinarum		12.45	0.51				
Staphylococcus xylosus		1.58	0.34				
247672	**Low discrimination**						**247672**
Staphylococcus equorum		86.20	0.38	GAL 91			
Staphylococcus gallinarum		12.02	0.18	MNS 99			
Staphylococcus xylosus		1.78	0.02				
247673	**Low discrimination**						**247673**
Staphylococcus equorum		86.14	0.79	GAL 91			
Staphylococcus gallinarum		12.02	0.59				
Staphylococcus xylosus		1.78	0.43				
247703	**Good identification**						**247703**
					NOV	YEP	
Staphylococcus hyicus		95.21	0.31	ARG 96	1	1	
Staphylococcus gallinarum		1.80	-.03	ESL 1	99	78	
Staphylococcus chromogenes		1.67	-.02		1	95	
247711	**Low discrimination**						**247711**
					OXI		
Staphylococcus equorum		48.14	0.30	XYL 99	1		
Staphylococcus sciuri		38.94	0.31	MLT 91	99		
Staphylococcus gallinarum		7.05	0.10		1		
247713	**Low discrimination**						**247713**
Staphylococcus equorum		72.96	0.38	XYL 99			
Staphylococcus gallinarum		10.18	0.18	MLT 91			
Staphylococcus xylosus		9.85	0.19				
247721	**Low discrimination**						**247721**
Staphylococcus gallinarum		56.83	0.30	MAN 91			
Staphylococcus equorum		39.62	0.30	MLT 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		3.55	0.07				
247723	**Low discrimination**						**247723**
Staphylococcus gallinarum		55.41	0.38	MAN 91			
Staphylococcus equorum		40.57	0.38	MLT 99			
Staphylococcus xylosus		4.02	0.16				
247730	**Low discrimination**						**247730**
					OXI		
Staphylococcus equorum		81.84	0.30	MLT 91	1		
Staphylococcus gallinarum		11.99	0.10	MNS 99	1		
Staphylococcus sciuri		6.17	0.10		99		
247731	**Low discrimination**						**247731**
Staphylococcus equorum		85.70	0.71	MLT 91			
Staphylococcus gallinarum		12.56	0.51				
Staphylococcus xylosus		1.21	0.31				
247732	**Low discrimination**						**247732**
Staphylococcus equorum		86.57	0.38	MLT 91			
Staphylococcus gallinarum		12.07	0.18	MNS 99			
Staphylococcus xylosus		1.36	0.00				
247733	**Low discrimination**						**247733**
Staphylococcus equorum		86.54	0.79	MLT 91			
Staphylococcus gallinarum		12.07	0.59				
Staphylococcus xylosus		1.36	0.41				
247741	**Low discrimination**						**247741**
Staphylococcus gallinarum		82.72	0.30	MAN 91			
Staphylococcus xylosus		5.97	0.08	XYL 99			
Staphylococcus equorum		5.89	0.09				
247743	**Low discrimination**						**247743**
Staphylococcus gallinarum		84.29	0.38	MAN 91			
Staphylococcus xylosus		7.07	0.18	XYL 99			
Staphylococcus equorum		6.30	0.18				
247750	**Low discrimination**						**247750**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		52.40	0.31	ESL 1	1	99	
Staphylococcus sciuri		23.32	0.24		99	1	
Staphylococcus gallinarum		9.08	0.10		1	12	
247751	**Low discrimination**						**247751**
					OXI		
Staphylococcus gallinarum		49.07	0.51	XYL 99	1		
Staphylococcus equorum		34.21	0.51		1		
Staphylococcus sciuri		10.30	0.43		99		
247753	**Low discrimination**						**247753**
Staphylococcus gallinarum		52.71	0.59	XYL 99			
Staphylococcus equorum		38.60	0.59				
Staphylococcus xylosus		6.85	0.42				
247760	**Acceptable identification**						**247760**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus gallinarum		93.35	0.30	MAN 91		
Staphylococcus equorum		6.65	0.09	MNS 99		
247761	**Good identification**					**247761**
Staphylococcus gallinarum		92.59	0.71	MAN 91		
Staphylococcus equorum		6.59	0.51			
Staphylococcus xylosus		0.78	0.30			
247762	**Acceptable identification**					**247762**
Staphylococcus gallinarum		92.21	0.38	MAN 91		
Staphylococcus equorum		6.90	0.18	MNS 99		
Staphylococcus xylosus		0.90	-.02			
247763	**Good identification**					**247763**
Staphylococcus gallinarum		92.21	0.79	MAN 91		
Staphylococcus equorum		6.90	0.59			
Staphylococcus xylosus		0.90	0.39			
247770	**Low discrimination**					**247770**
					OXI	
Staphylococcus gallinarum		57.73	0.51	MNS 99	1	
Staphylococcus equorum		40.25	0.51		1	
Staphylococcus sciuri		1.13	0.22		99	
247771	**Low discrimination**					**247771**
Staphylococcus gallinarum		58.42	0.92			
Staphylococcus equorum		40.73	0.92			
Staphylococcus xylosus		0.76	0.55			
247772	**Low discrimination**					**247772**
Staphylococcus gallinarum		57.08	0.59	MNS 99		
Staphylococcus equorum		41.80	0.59			
Staphylococcus xylosus		0.86	0.23			
247773	**Low discrimination**					**247773**
Staphylococcus gallinarum		57.23	1.00			
Staphylococcus equorum		41.90	1.00			
Staphylococcus xylosus		0.86	0.64			
250240	**Good identification**					**250240**
Staphylococcus warneri		100.0	0.27	ORN 1		
250413	**Good identification**					**250413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	ORN 1		
250440	**Good identification**					**250440**
Staphylococcus warneri		100.0	0.27	ORN 1		
250443	**Good identification**					**250443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
250451	**Good identification**					**250451**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1			
250453	**Good identification**						**250453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	ORN 1			
250513	**Good identification**						**250513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.29	ORN 1			
250553	**Good identification**						**250553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	ORN 1			
250600	**Low discrimination**						**250600**
Staphylococcus warneri		88.02	0.27	ORN 1			
Staphylococcus pasteuri		11.98	-.06	MLT 95			
250640	**Very good identification**						**250640**
Staphylococcus warneri		95.79	0.53	ORN 1			
Staphylococcus pasteuri		4.22	0.11				
250642	**Acceptable identification**						**250642**
Staphylococcus warneri		94.88	0.43	ORN 1			
Staphylococcus pasteuri		5.12	0.03				
250650	**Low discrimination**						**250650**
Staphylococcus warneri		72.82	0.50	ORN 1			
Staphylococcus pasteuri		27.18	0.27				
250652	**Low discrimination**						**250652**
					NOV		
Staphylococcus warneri		66.09	0.40	ORN 1	1		
Staphylococcus pasteuri		30.23	0.19		1		
Staphylococcus saprophyticus subsp. saprophyticus		3.69	-.06		99		
250740	**Good identification**						**250740**
Staphylococcus warneri		100.0	0.43	ORN 1			
250742	**Good identification**						**250742**
Staphylococcus warneri		100.0	0.33	ORN 1			
250750	**Good identification**						**250750**
					NOV	PYR	
Staphylococcus warneri		96.03	0.40	ORN 1	1	8	
Staphylococcus saprophyticus subsp. bovis		3.97	0.02		99	99	
250752	**Good identification**						**250752**
Staphylococcus warneri		100.0	0.30	ORN 1			
250773	**Acceptable identification**						**250773**
Staphylococcus xylosus		92.96	0.28	ORN 1			
Staphylococcus arlettae		7.04	0.14	NIT 89			
251413	**Good identification**						**251413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
251453	**Good identification**						**251453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ORN 1			
251553	**Good identification**						**251553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	ORN 1			
251773	**Low discrimination**						**251773**
					YEP		
Staphylococcus equorum		83.13	0.34	ORN 1	1		
Staphylococcus xylosus		12.95	0.15	NIT 92	40		
Staphylococcus arlettae		3.92	0.14		99		
252453	**Good identification**						**252453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	ORN 1			
252550	**Good identification**						**252550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	ORN 1			
252553	**Low discrimination**						**252553**
Staphylococcus cohnii subsp. urealyticum		80.72	0.29	ORN 1			
Staphylococcus saprophyticus subsp. bovis		11.63	0.06				
Staphylococcus xylosus		7.65	-.02				
252640	**Low discrimination**						**252640**
Staphylococcus warneri		76.16	0.46	ORN 1			
Staphylococcus pasteuri		19.15	0.19				
Staphylococcus hominis subsp. hominis		4.69	0.15				
252642	**Low discrimination**						**252642**
Staphylococcus warneri		65.57	0.36	ORN 1			
Staphylococcus pasteuri		20.20	0.11				
Staphylococcus hominis subsp. hominis		14.23	0.16				
252650	**Low discrimination**						**252650**
					NOV	PYR	
Staphylococcus pasteuri		54.88	0.35	VPT 86	1	1	
Staphylococcus warneri		25.73	0.43	ORN 1	1	8	
Staphylococcus saprophyticus subsp. bovis		19.39	0.31		99	99	
252652	**Low discrimination**						**252652**
					NOV	PYR	
Staphylococcus pasteuri		61.50	0.27	VPT 86	1	1	
Staphylococcus warneri		23.54	0.33	ORN 1	1	8	
Staphylococcus saprophyticus subsp. bovis		13.30	0.18		99	99	
252740	**Low discrimination**						**252740**
					NOV	PYR	
Staphylococcus warneri		78.05	0.35	ORN 1	1	8	
Staphylococcus hominis subsp. hominis		16.50	0.16		1	1	
Staphylococcus saprophyticus subsp. bovis		5.45	0.02		99	99	
252750	**Acceptable identification**						**252750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		90.52	0.43	ORN 1	99	99	
Staphylococcus warneri		9.48	0.32		1	8	
252751	**Low discrimination**						**252751**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		78.45	0.31	ORN 1			
Staphylococcus xylosus		21.55	0.15				
252752	**Low discrimination**						**252752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		87.74	0.31	ORN 1	99	99	
Staphylococcus warneri		12.26	0.23		1	8	
252771	**Good identification**						**252771**
Staphylococcus xylosus		97.51	0.37	ORN 1			
Staphylococcus equorum		2.49	0.06				
252773	**Good identification**						**252773**
Staphylococcus xylosus		97.75	0.47	ORN 1			
Staphylococcus equorum		2.25	0.14				
253453	**Good identification**						**253453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ORN 1			
253673	**Acceptable identification**						**253673**
Staphylococcus equorum		90.12	0.35	ORN 1			
Staphylococcus xylosus		9.88	0.14	GAL 91			
253733	**Acceptable identification**						**253733**
Staphylococcus equorum		92.30	0.35	ORN 1			
Staphylococcus xylosus		7.70	0.11	MLT 91			
253771	**Low discrimination**						**253771**
Staphylococcus equorum		89.23	0.47	ORN 1			
Staphylococcus xylosus		8.83	0.25				
Staphylococcus gallinarum		1.94	0.10				
253773	**Low discrimination**						**253773**
Staphylococcus equorum		88.47	0.55	ORN 1			
Staphylococcus xylosus		9.70	0.34				
Staphylococcus gallinarum		1.83	0.18				
254413	**Good identification**						**254413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	ORN 1			
254443	**Good identification**						**254443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1			
254450	**Good identification**						**254450**
Staphylococcus kloosii		100.0	0.47	ORN 1			
254451	**Good identification**						**254451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1			
254452	**Acceptable identification**						**254452**
Staphylococcus kloosii		91.66	0.48	ORN 1			
Staphylococcus cohnii subsp. urealyticum		8.34	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
254453	**Very good identification**					**254453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	ORN 1		
254470	**Good identification**					**254470**
Staphylococcus kloosii		100.0	0.34	ORN 1		
254472	**Good identification**					**254472**
Staphylococcus kloosii		100.0	0.35	ORN 1		
254512	**Good identification**					**254512**
Staphylococcus kloosii		100.0	0.27	ORN 1		
254513	**Good identification**					**254513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	ORN 1		
254543	**Good identification**					**254543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1		
254550	**Good identification**					**254550**
Staphylococcus kloosii		100.0	0.53	ORN 1		
254551	**Good identification**					**254551**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1		
254552	**Very good identification**					**254552**
Staphylococcus kloosii		98.03	0.53	ORN 1		
Staphylococcus cohnii subsp. urealyticum		1.97	0.07			
254553	**Good identification**					**254553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	ORN 1		
254570	**Good identification**					**254570**
Staphylococcus kloosii		100.0	0.40	ORN 1		
254572	**Good identification**					**254572**
Staphylococcus kloosii		100.0	0.41	ORN 1		
254652	**Good identification**					**254652**
Staphylococcus kloosii		100.0	0.27	ORN 1		
254750	**Good identification**					**254750**
Staphylococcus kloosii		100.0	0.32	ORN 1		
254752	**Good identification**					**254752**
Staphylococcus kloosii		100.0	0.33	ORN 1		
255413	**Good identification**					**255413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
255450	**Good identification**					**255450**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.47	ORN 1		
255452	**Good identification**					**255452**
Staphylococcus kloosii		96.58	0.48	ORN 1		
Staphylococcus cohnii subsp. urealyticum		3.42	0.06			
255453	**Good identification**					**255453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	ORN 1		
255470	**Good identification**					**255470**
Staphylococcus kloosii		100.0	0.34	ORN 1		
255472	**Good identification**					**255472**
Staphylococcus kloosii		100.0	0.35	ORN 1		
255512	**Good identification**					**255512**
Staphylococcus kloosii		100.0	0.27	ORN 1		
255550	**Good identification**					**255550**
Staphylococcus kloosii		100.0	0.53	ORN 1		
255552	**Very good identification**					**255552**
Staphylococcus kloosii		100.0	0.53	ORN 1		
255553	**Good identification**					**255553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ORN 1		
255570	**Good identification**					**255570**
Staphylococcus kloosii		100.0	0.40	ORN 1		
255572	**Good identification**					**255572**
Staphylococcus kloosii		100.0	0.41	ORN 1		
255652	**Good identification**					**255652**
Staphylococcus kloosii		100.0	0.27	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
255750	**Good identification**					**255750**
	Staphylococcus kloosii	100.0	0.32	ORN 1		
255752	**Good identification**					**255752**
	Staphylococcus kloosii	100.0	0.33	ORN 1		
255771	**Low discrimination**					**255771**
	Staphylococcus equorum	85.72	0.29	ORN 1		
	Staphylococcus gallinarum	14.28	0.10	NIT 92		
255773	**Low discrimination**					**255773**
					YEP	
	Staphylococcus equorum	80.96	0.37	ORN 1	1	
	Staphylococcus gallinarum	12.84	0.18	NIT 92	78	
	Staphylococcus arlettae	3.82	0.18		99	
256453	**Good identification**					**256453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.40	ORN 1		
256553	**Good identification**					**256553**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.32	ORN 1		
256703	**Good identification**					**256703**
					YEP	
	Staphylococcus hyicus	98.28	0.31	ARG 96	1	
	Staphylococcus chromogenes	1.72	-.02	ORN 1	95	
256750	**Good identification**					**256750**
	Staphylococcus saprophyticus subsp. bovis	100.0	0.31	ORN 1		
256773	**Low discrimination**					**256773**
	Staphylococcus xylosus	77.53	0.35	ORN 1		
	Staphylococcus gallinarum	12.98	0.18			
	Staphylococcus equorum	9.50	0.18			
257453	**Good identification**					**257453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.31	ORN 1		
257671	**Low discrimination**					**257671**
	Staphylococcus equorum	87.22	0.30	ORN 1		
	Staphylococcus gallinarum	12.78	0.10	GAL 91		
257673	**Low discrimination**					**257673**
	Staphylococcus equorum	86.20	0.38	ORN 1		
	Staphylococcus gallinarum	12.02	0.18	GAL 91		
	Staphylococcus xylosus	1.78	0.02			
257731	**Low discrimination**					**257731**
	Staphylococcus equorum	87.22	0.30	ORN 1		
	Staphylococcus gallinarum	12.78	0.10	MLT 91		
257733	**Low discrimination**					**257733**
	Staphylococcus equorum	86.57	0.38	ORN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus gallinarum		12.07	0.18	MLT 91		
Staphylococcus xylosus		1.36	0.00			
257761	**Acceptable identification**					**257761**
Staphylococcus gallinarum		93.35	0.30	ORN 1		
Staphylococcus equorum		6.65	0.09	MAN 91		
257763	**Acceptable identification**					**257763**
Staphylococcus gallinarum		92.21	0.38	ORN 1		
Staphylococcus equorum		6.90	0.18	MAN 91		
Staphylococcus xylosus		0.90	-.02			
257771	**Low discrimination**					**257771**
Staphylococcus gallinarum		58.47	0.51	ORN 1		
Staphylococcus equorum		40.77	0.51			
Staphylococcus xylosus		0.76	0.14			
257773	**Low discrimination**					**257773**
Staphylococcus gallinarum		57.23	0.59	ORN 1		
Staphylococcus equorum		41.91	0.59			
Staphylococcus xylosus		0.86	0.23			
260000	**Low discrimination**					**260000**
					OXI	
Kocuria varians		55.33	0.31	GLR 1	13	
Dermacoccus nishinomiyaensis		29.08	0.18		93	
Micrococcus sp.		15.59	0.03		89	
260013	**Good identification**					**260013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	TRE 99		
260040	**Low discrimination**					**260040**
Staphylococcus warneri		61.98	0.29	SUC 95		
Kocuria varians		38.03	0.18	TRE 95		
260043	**Good identification**					**260043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
260051	**Good identification**					**260051**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
260053	**Very good identification**					**260053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	TRE 99		
260100	**Acceptable identification**					**260100**
					OXI	
Kocuria varians		93.74	0.35	GLR 1	13	
Dermacoccus nishinomiyaensis		6.26	0.03		93	
260113	**Good identification**					**260113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	TRE 99		
260143	**Good identification**					**260143**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
260151	**Acceptable identification**						**260151**
Staphylococcus cohnii subsp. urealyticum		92.52	0.27	TRE 99			
Staphylococcus xylosus		7.48	-.05	LAC 91			
260153	**Good identification**						**260153**
Staphylococcus cohnii subsp. urealyticum		97.78	0.48	TRE 99			
Staphylococcus xylosus		2.22	0.05				
260173	**Good identification**						**260173**
Staphylococcus xylosus		96.28	0.27	NIT 89			
Staphylococcus cohnii subsp. urealyticum		3.72	0.07	SUC 95			
260200	**Low discrimination**						**260200**
Staphylococcus warneri		72.40	0.29	TRE 95			
Kocuria varians		27.60	0.14	MLT 95			
260240	**Very good identification**						**260240**
Staphylococcus warneri		100.0	0.56	TRE 95			
260242	**Good identification**						**260242**
Staphylococcus warneri		100.0	0.46	TRE 95			
260250	**Good identification**						**260250**
					NOV		
Staphylococcus warneri		98.59	0.53	TRE 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		1.41	-.06		99		
260252	**Low discrimination**						**260252**
					NOV		
Staphylococcus warneri		77.65	0.43	TRE 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		18.83	0.09		99		
Staphylococcus simulans		3.52	0.06		1		
260273	**Good identification**						**260273**
Staphylococcus xylosus		100.0	0.32	NIT 89			
260333	**Good identification**						**260333**
Staphylococcus xylosus		100.0	0.30	NIT 89			
260340	**Good identification**						**260340**
Staphylococcus warneri		97.95	0.45	TRE 95			
Kocuria varians		2.05	0.04				
260342	**Good identification**						**260342**
Staphylococcus warneri		100.0	0.35	TRE 95			
260350	**Good identification**						**260350**
					NOV	PYR	
Staphylococcus warneri		95.46	0.42	TRE 95	1	8	
Staphylococcus saprophyticus subsp. bovis		4.54	0.05		99	99	
260352	**Good identification**						**260352**
Staphylococcus warneri		100.0	0.33	TRE 95			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
260353	**Acceptable identification**					**260353**
	Staphylococcus xylosus	93.32	0.31	NIT 89		
	Staphylococcus cohnii subsp. urealyticum	6.68	0.17	TRE 96		
260363	**Good identification**					**260363**
	Staphylococcus xylosus	100.0	0.28	NIT 89		
260371	**Good identification**					**260371**
	Staphylococcus xylosus	100.0	0.44	NIT 89		
260373	**Very good identification**					**260373**
	Staphylococcus xylosus	99.54	0.53	NIT 89		
	Staphylococcus arlettae	0.46	0.14	TRE 96		
260400	**Low discrimination**					**260400**
	Staphylococcus warneri	82.62	0.29	SUC 95		
	Kocuria varians	17.38	0.08	MLT 95		
260401	**Good identification**					**260401**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.40	MAN 91		
260403	**Very good identification**					**260403**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	99.40	0.60	MAN 91	99	
	Staphylococcus simulans	0.60	0.09		1	
260411	**Very good identification**					**260411**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.60	LAC 91		
260412	**Low discrimination**					**260412**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	85.35	0.40	MNS 99	99	
	Staphylococcus simulans	6.04	0.11		1	
	Staphylococcus warneri	3.98	0.16		1	
260413	**Very good identification**					**260413**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	99.76	0.81		99	
	Staphylococcus simulans	0.24	0.21		1	
260433	**Low discrimination**					**260433**
	Staphylococcus cohnii subsp. urealyticum	89.81	0.40	XYL 1		
	Staphylococcus xylosus	10.19	0.11			
260440	**Very good identification**					**260440**
					NOV	
	Staphylococcus warneri	98.51	0.56	SUC 95	1	
	Staphylococcus cohnii subsp. urealyticum	1.49	0.14		99	
260441	**Very good identification**					**260441**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	98.66	0.55	MAN 91	99	
	Staphylococcus warneri	1.35	0.21	LAC 91	1	
260442	**Low discrimination**					**260442**
					NOV	
	Staphylococcus warneri	65.35	0.46	SUC 95	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		29.96	0.35		99		
Staphylococcus saprophyticus subsp. saprophyticus		4.68	0.01		99		
260443	**Very good identification**						**260443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.91	0.76	MAN 91	99		
Staphylococcus simulans		0.05	0.02		1		
Staphylococcus warneri		0.05	0.11		1		
260450	**Low discrimination**						**260450**
					NOV		
Staphylococcus warneri		58.51	0.53	SUC 95	1		
Staphylococcus kloosii		12.80	0.47		99		
Staphylococcus cohnii subsp. urealyticum		12.35	0.35		99		
260451	**Very good identification**						**260451**
					NOV	PYR	
Staphylococcus cohnii subsp. urealyticum		99.71	0.76	LAC 91	99	53	
Staphylococcus saprophyticus subsp. bovis		0.11	0.09		99	99	
Staphylococcus warneri		0.10	0.18		1	8	
260452	**Low discrimination**						**260452**
					NOV		
Staphylococcus cohnii subsp. urealyticum		60.43	0.55	MNS 99	99		
Staphylococcus saprophyticus subsp. saprophyticus		22.42	0.30		99		
Staphylococcus warneri		9.44	0.43		1		
260453	**Excellent identification**						**260453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.95	0.96		99		
Staphylococcus xylosus		0.02	0.13		99		
Staphylococcus simulans		0.02	0.14		1		
260463	**Low discrimination**						**260463**
Staphylococcus cohnii subsp. urealyticum		85.34	0.35	MAN 91			
Staphylococcus xylosus		14.66	0.10	XYL 1			
260470	**Low discrimination**						**260470**
					NOV		
Staphylococcus kloosii		84.41	0.34	PHS 99	99		
Staphylococcus warneri		15.59	0.12		1		
260471	**Low discrimination**						**260471**
Staphylococcus cohnii subsp. urealyticum		51.40	0.35	XYL 1			
Staphylococcus xylosus		48.60	0.25	LAC 91			
260472	**Low discrimination**						**260472**
Staphylococcus kloosii		66.72	0.35	PHS 99			
Staphylococcus cohnii subsp. urealyticum		24.28	0.14				
Staphylococcus saprophyticus subsp. saprophyticus		9.01	-.11				
260473	**Low discrimination**						**260473**
					NOV		
Staphylococcus cohnii subsp. urealyticum		78.55	0.55	XYL 1	99		
Staphylococcus xylosus		20.90	0.34		99		
Aerococcus viridans		0.54	0.22		1		
260501	**Good identification**						**260501**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
260503	**Good identification**						**260503**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.53	MAN 91			
260511	**Good identification**						**260511**
	Staphylococcus cohnii subsp. urealyticum	99.17	0.53	LAC 91			
	Staphylococcus xylosus	0.83	0.01				
260512	**Low discrimination**						**260512**
	Staphylococcus cohnii subsp. urealyticum	86.96	0.32	MNS 99			
	Staphylococcus kloosii	13.04	0.27				
260513	**Very good identification**						**260513**
					NOV		
	Staphylococcus cohnii subsp. urealyticum	99.72	0.73		99		
	Staphylococcus xylosus	0.24	0.10		99		
	Staphylococcus simulans	0.05	-.01		1		
260533	**Low discrimination**						**260533**
	Staphylococcus xylosus	72.81	0.32	NIT 89			
	Staphylococcus cohnii subsp. urealyticum	27.19	0.32	SUC 95			
260540	**Good identification**						**260540**
					NOV		
	Staphylococcus warneri	97.99	0.45	SUC 95	1		
	Staphylococcus cohnii subsp. urealyticum	2.01	0.06		99		
260541	**Good identification**						**260541**
					NOV		
	Staphylococcus cohnii subsp. urealyticum	97.78	0.47	MAN 91	99		
	Staphylococcus xylosus	1.24	-.01	LAC 91	99		
	Staphylococcus warneri	0.98	0.10		1		
260542	**Low discrimination**						**260542**
					NOV		
	Staphylococcus warneri	61.64	0.35	SUC 95	1		
	Staphylococcus cohnii subsp. urealyticum	38.36	0.27		99		
260543	**Very good identification**						**260543**
	Staphylococcus cohnii subsp. urealyticum	99.65	0.68	MAN 91			
	Staphylococcus xylosus	0.36	0.09				
260550	**Low discrimination**						**260550**
					NOV	PYR	
	Staphylococcus kloosii	34.38	0.53	PHS 99	99	52	
	Staphylococcus saprophyticus subsp. bovis	30.12	0.34		99	99	
	Staphylococcus warneri	25.57	0.42		1	8	
260551	**Very good identification**						**260551**
	Staphylococcus cohnii subsp. urealyticum	97.09	0.68	LAC 91			
	Staphylococcus xylosus	1.90	0.24				
	Staphylococcus saprophyticus subsp. bovis	1.01	0.22				
260552	**Low discrimination**						**260552**
					NOV		
	Staphylococcus cohnii subsp. urealyticum	55.71	0.48	MNS 99	99		
	Staphylococcus kloosii	28.02	0.53		99		
	Staphylococcus warneri	6.41	0.33		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
260553	**Very good identification**					**260553**
Staphylococcus cohnii subsp. urealyticum		99.41	0.89			
Staphylococcus xylosus		0.55	0.33			
Staphylococcus saprophyticus subsp. bovis		0.03	0.09			
260563	**Low discrimination**					**260563**
Staphylococcus xylosus		80.21	0.30	NIT 89		
Staphylococcus cohnii subsp. urealyticum		19.80	0.27	SUC 95		
260570	**Low discrimination**					**260570**
Staphylococcus kloosii		83.90	0.40	PHS 99		
Staphylococcus xylosus		16.11	0.05			
260571	**Good identification**					**260571**
Staphylococcus xylosus		95.71	0.46	NIT 89		
Staphylococcus cohnii subsp. urealyticum		4.29	0.27	SUC 95		
260572	**Low discrimination**					**260572**
Staphylococcus kloosii		58.09	0.41	PHS 99		
Staphylococcus xylosus		29.30	0.14			
Staphylococcus arlettae		7.95	0.12			
260573	**Low discrimination**					**260573**
Staphylococcus xylosus		85.80	0.55	NIT 89		
Staphylococcus cohnii subsp. urealyticum		13.67	0.48	SUC 95		
Staphylococcus arlettae		0.31	0.14			
260600	**Very good identification**					**260600**
Staphylococcus warneri		99.45	0.56	MLT 95		
Staphylococcus pasteuri		0.55	-.06			
260602	**Low discrimination**					**260602**
Staphylococcus warneri		74.22	0.46	MLT 95		
Staphylococcus simulans		25.78	0.27			
260603	**Low discrimination**					**260603**
					NOV	
Staphylococcus simulans		81.26	0.37	ARG 92	1	
Staphylococcus cohnii subsp. urealyticum		17.32	0.29	NIT 96	99	
Staphylococcus warneri		1.43	0.11		1	
260610	**Acceptable identification**					**260610**
					NOV	
Staphylococcus warneri		94.26	0.53	MLT 95	1	
Staphylococcus pasteuri		4.40	0.11		1	
Staphylococcus saprophyticus subsp. saprophyticus		1.35	-.06		99	
260611	**Low discrimination**					**260611**
					NOV	
Staphylococcus cohnii subsp. urealyticum		63.05	0.29	SUC 3	99	
Staphylococcus xylosus		13.72	0.06	LAC 91	99	
Staphylococcus simulans		11.95	0.09		1	
260612	**Low discrimination**					**260612**
					NOV	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus simulans		58.97	0.39	ARG 92	1			
Staphylococcus warneri		30.74	0.43	NIT 96	1			
Staphylococcus saprophyticus subsp. saprophyticus		7.46	0.09		99			
260613	**Low discrimination**							**260613**
					NOV			
Staphylococcus simulans		63.63	0.50	ARG 92	1			
Staphylococcus cohnii subsp. urealyticum		34.27	0.50	NIT 96	99			
Staphylococcus xylosus		2.10	0.16		99			
260631	**Good identification**							**260631**
Staphylococcus xylosus		100.0	0.28	NIT 89				
260633	**Good identification**							**260633**
					NOV			
Staphylococcus xylosus		96.06	0.38	NIT 89	99			
Staphylococcus simulans		2.56	0.09	GAL 91	1			
Staphylococcus cohnii subsp. urealyticum		1.38	0.09	MLT 93	99			
260640	**Very good identification**							**260640**
					NOV			
Staphylococcus warneri		99.60	0.82		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.22	0.07		99			
Staphylococcus pasteuri		0.18	0.11		1			
260641	**Low discrimination**							**260641**
					OXI	NOV	PYR	
Staphylococcus warneri		80.57	0.47	MNS 2	1	1	8	
Staphylococcus cohnii subsp. urealyticum		9.62	0.24		1	99	53	
Kocuria kristinae		6.65	0.10		99	1	93	
260642	**Very good identification**							**260642**
					NOV			
Staphylococcus warneri		95.37	0.72		1			
Staphylococcus saprophyticus subsp. saprophyticus		3.64	0.22		99			
Staphylococcus simulans		0.78	0.19		1			
260643	**Low discrimination**							**260643**
					NOV			
Staphylococcus cohnii subsp. urealyticum		56.88	0.45	SUC 3	99			
Staphylococcus simulans		21.16	0.30	MAN 91	1			
Staphylococcus warneri		15.71	0.37		1			
260650	**Good identification**							**260650**
					NOV	PYR		
Staphylococcus warneri		90.08	0.79		1	8		
Staphylococcus saprophyticus subsp. saprophyticus		6.70	0.35		99	7		
Staphylococcus saprophyticus subsp. bovis		1.76	0.34		99	99		
260651	**Low discrimination**							**260651**
					NOV			
Staphylococcus cohnii subsp. urealyticum		43.65	0.45	SUC 3	99			
Staphylococcus warneri		26.19	0.44	LAC 91	1			
Staphylococcus xylosus		22.27	0.29		99			
260652	**Low discrimination**							**260652**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		53.60	0.50	GLR 2	99			
Staphylococcus warneri		42.41	0.69		1			
Staphylococcus simulans		1.92	0.32		1			
260653	**Low discrimination**							**260653**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus cohnii subsp. urealyticum		75.26	0.65	SUC 3	99		
Staphylococcus simulans		11.08	0.43		1		
Staphylococcus xylosus		10.81	0.39		99		
260660	**Good identification**						**260660**
Staphylococcus warneri		100.0	0.41	XYL 1			
260661	**Good identification**						**260661**
Staphylococcus xylosus		100.0	0.27	NIT 89			
260662	**Good identification**						**260662**
Staphylococcus warneri		100.0	0.31	XYL 1			
260663	**Good identification**						**260663**
Staphylococcus xylosus		100.0	0.36	NIT 89			
260670	**Low discrimination**						**260670**
					NOV		
Staphylococcus warneri		78.50	0.38	XYL 1	1		
Staphylococcus xylosus		15.66	0.10		99		
Staphylococcus saprophyticus subsp. saprophyticus		5.84	-.06		99		
260671	**Good identification**						**260671**
Staphylococcus xylosus		99.63	0.51	NIT 89			
Staphylococcus equorum		0.37	0.03	GAL 91			
260672	**Low discrimination**						**260672**
Staphylococcus arlettae		52.01	0.37	URE 1			
Staphylococcus xylosus		20.61	0.20				
Staphylococcus saprophyticus subsp. saprophyticus		15.29	0.09				
260673	**Very good identification**						**260673**
Staphylococcus xylosus		95.65	0.61	NIT 89			
Staphylococcus arlettae		3.23	0.40	GAL 91			
Staphylococcus cohnii subsp. urealyticum		0.59	0.24				
260700	**Good identification**						**260700**
Staphylococcus warneri		100.0	0.45	MLT 95			
260702	**Acceptable identification**						**260702**
Staphylococcus warneri		91.27	0.35	MLT 95			
Staphylococcus simulans		8.73	0.05				
260710	**Good identification**						**260710**
					NOV	PYR	
Staphylococcus warneri		95.46	0.42	MLT 95	1	8	
Staphylococcus saprophyticus subsp. bovis		4.54	0.05		99	99	
260711	**Low discrimination**						**260711**
Staphylococcus xylosus		83.70	0.27	NIT 89			
Staphylococcus cohnii subsp. urealyticum		16.31	0.22	XYL 92			
260712	**Low discrimination**						**260712**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		62.63	0.33	MLT 95	1		
Staphylococcus simulans		33.08	0.17		1		
Staphylococcus xylosus		4.29	-.05		99		
260713	**Low discrimination**						**260713**
					NOV		
Staphylococcus xylosus		49.98	0.36	NIT 89	99		
Staphylococcus cohnii subsp. urealyticum		34.59	0.42	XYL 92	99		
Staphylococcus simulans		13.03	0.28	MLT 93	1		
260723	**Good identification**						**260723**
Staphylococcus xylosus		100.0	0.34	NIT 89			
260731	**Good identification**						**260731**
Staphylococcus xylosus		99.52	0.49	NIT 89			
Staphylococcus equorum		0.48	0.03	MLT 93			
260733	**Very good identification**						**260733**
					YEP		
Staphylococcus xylosus		99.32	0.58	NIT 89	40		
Staphylococcus equorum		0.43	0.11	MLT 93	1		
Staphylococcus arlettae		0.25	0.14		99		
260740	**Very good identification**						**260740**
					NOV	PYR	
Staphylococcus warneri		99.82	0.72		1	8	
Staphylococcus saprophyticus subsp. bovis		0.18	0.05		99	99	
260742	**Very good identification**						**260742**
					NOV		
Staphylococcus warneri		98.96	0.62		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.04	0.00		99		
260743	**Low discrimination**						**260743**
					NOV		
Staphylococcus xylosus		61.86	0.35	NIT 89	99		
Staphylococcus cohnii subsp. urealyticum		28.29	0.37	MAN 94	99		
Staphylococcus warneri		5.76	0.27	XYL 92	1		
260750	**Low discrimination**						**260750**
					NOV	PYR	
Staphylococcus warneri		78.06	0.69		1	8	
Staphylococcus saprophyticus subsp. bovis		19.36	0.47		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		1.60	0.13		99	7	
260751	**Low discrimination**						**260751**
Staphylococcus xylosus		81.04	0.50	NIT 89			
Staphylococcus saprophyticus subsp. bovis		9.04	0.34	XYL 92			
Staphylococcus cohnii subsp. urealyticum		6.73	0.37				
260752	**Low discrimination**						**260752**
					NOV	PYR	
Staphylococcus warneri		61.01	0.59		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		21.23	0.29		99	7	
Staphylococcus saprophyticus subsp. bovis		11.35	0.34		99	99	
260753	**Low discrimination**						**260753**
Staphylococcus xylosus		75.34	0.60	NIT 89			
Staphylococcus cohnii subsp. urealyticum		22.24	0.58	XYL 92			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus saprophyticus subsp. bovis		0.74	0.22			
260760	**Low discrimination**					**260760**
					NOV	
Staphylococcus warneri		77.20	0.30	XYL 1	1	
Staphylococcus xylosus		22.80	0.06		99	
260761	**Good identification**					**260761**
Staphylococcus xylosus		100.0	0.47	NIT 89		
260763	**Very good identification**					**260763**
Staphylococcus xylosus		99.70	0.57	NIT 89		
Staphylococcus arlettae		0.30	0.14	MAN 94		
260770	**Low discrimination**					**260770**
					NOV	
Staphylococcus xylosus		79.17	0.31	NIT 89	99	
Staphylococcus warneri		12.39	0.28	MNS 99	1	
Staphylococcus arlettae		3.22	0.12		99	
260771	**Very good identification**					**260771**
					YEP	
Staphylococcus xylosus		99.58	0.72	NIT 89	40	
Staphylococcus equorum		0.37	0.23		1	
Staphylococcus arlettae		0.05	0.14		99	
260772	**Low discrimination**					**260772**
					NOV	
Staphylococcus arlettae		57.89	0.53	URE 1	99	
Staphylococcus xylosus		40.93	0.40		99	
Staphylococcus warneri		0.75	0.18		1	
260773	**Very good identification**					**260773**
					YEP	
Staphylococcus xylosus		97.79	0.81	NIT 89	40	
Staphylococcus arlettae		1.85	0.55		99	
Staphylococcus equorum		0.32	0.32		1	
261013	**Good identification**					**261013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
261053	**Good identification**					**261053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	TRE 99		
261153	**Good identification**					**261153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	TRE 99		
261371	**Low discrimination**					**261371**
Staphylococcus xylosus		73.94	0.31	NIT 89		
Staphylococcus equorum		26.06	0.23	TRE 96		
261373	**Low discrimination**					**261373**
					YEP	
Staphylococcus xylosus		74.81	0.41	NIT 89	40	
Staphylococcus equorum		23.82	0.32	TRE 96	1	
Staphylococcus arlettae		1.37	0.14		99	
261401	**Good identification**					**261401**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	MAN 91			
261403	**Good identification**						**261403**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.23	0.52	MAN 91	99		
Aerococcus viridans		0.77	0.19		1		
261411	**Good identification**						**261411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	LAC 91			
261412	**Low discrimination**						**261412**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		84.57	0.31	MNS 99	99	64	
Aerococcus viridans		8.22	0.22		1	1	
Staphylococcus kloosii		7.20	0.21		99	99	
261413	**Very good identification**						**261413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.77	0.72		99		
Aerococcus viridans		0.23	0.29		1		
261433	**Low discrimination**						**261433**
					NOV		
Staphylococcus cohnii subsp. urealyticum		76.93	0.31	XYL 1	99		
Aerococcus viridans		17.46	0.29		1		
Staphylococcus xylosus		5.61	-.01		99		
261441	**Good identification**						**261441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	MAN 91			
261443	**Very good identification**						**261443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.46	0.67	MAN 91	99		
Aerococcus viridans		0.54	0.32		1		
261450	**Low discrimination**						**261450**
					OXI		
Staphylococcus kloosii		58.80	0.47	PHS 99	1		
Staphylococcus cohnii subsp. urealyticum		22.06	0.26		1		
Macrococcus equipersicus		16.42	0.18		99		
261451	**Very good identification**						**261451**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.82	0.67	LAC 91	99		
Aerococcus viridans		0.18	0.22		1		
261452	**Low discrimination**						**261452**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		73.32	0.47	MNS 99	99	64	
Staphylococcus kloosii		20.94	0.48		99	99	
Aerococcus viridans		5.03	0.34		1	1	
261453	**Very good identification**						**261453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.82	0.88		99		
Aerococcus viridans		0.16	0.42		1		
Staphylococcus xylosus		0.02	0.00		99		
261470	**Good identification**						**261470**
Staphylococcus kloosii		100.0	0.34	PHS 99			
261472	**Low discrimination**						**261472**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus kloosii		47.55	0.35	PHS 99	99	99	
Aerococcus viridans		45.72	0.34		1	1	
Staphylococcus cohnii subsp. urealyticum		6.73	0.06		99	64	
261473	**Low discrimination**						**261473**
					NOV		
Staphylococcus cohnii subsp. urealyticum		72.76	0.47	XYL 1	99		
Staphylococcus xylosus		12.45	0.22		99		
Aerococcus viridans		11.65	0.42		1		
261503	**Good identification**						**261503**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.23	0.44	MAN 91	99		
Aerococcus viridans		1.77	0.19		1		
261511	**Good identification**						**261511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	LAC 91			
261513	**Very good identification**						**261513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.32	0.65		99		
Aerococcus viridans		0.53	0.29		1		
Staphylococcus xylosus		0.15	-.02		99		
261541	**Good identification**						**261541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	MAN 91			
261543	**Very good identification**						**261543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.52	0.60	MAN 91	99		
Aerococcus viridans		1.25	0.32		1		
Staphylococcus xylosus		0.23	-.04		99		
261550	**Low discrimination**						**261550**
					OXI		
Staphylococcus kloosii		87.37	0.53	PHS 99	1		
Staphylococcus cohnii subsp. urealyticum		7.24	0.19		1		
Macrococcus equipersicus		5.39	0.10		99		
261551	**Very good identification**						**261551**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.34	0.60	LAC 91	99		
Staphylococcus xylosus		1.24	0.11		99		
Aerococcus viridans		0.42	0.22		1		
261552	**Low discrimination**						**261552**
					NOV	YEP	
Staphylococcus kloosii		52.71	0.53	PHS 99	99	99	
Staphylococcus cohnii subsp. urealyticum		40.76	0.39		99	64	
Aerococcus viridans		6.53	0.34		1	1	
261553	**Very good identification**						**261553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.27	0.80		99		
Aerococcus viridans		0.38	0.42		1		
Staphylococcus xylosus		0.35	0.21		99		
261570	**Good identification**						**261570**
Staphylococcus kloosii		100.0	0.40	PHS 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
261571	**Low discrimination**						**261571**
Staphylococcus xylosus		72.57	0.33	NIT 89			
Staphylococcus equorum		20.24	0.23	SUC 95			
Staphylococcus cohnii subsp. urealyticum		5.06	0.19				
261572	**Low discrimination**						**261572**
					NOV	YEP	
Staphylococcus kloosii		56.87	0.41	PHS 99	99	99	
Aerococcus viridans		28.17	0.34		1	1	
Staphylococcus arlettae		7.79	0.12		99	99	
261573	**Low discrimination**						**261573**
Staphylococcus xylosus		62.23	0.43	NIT 89			
Staphylococcus equorum		15.68	0.32	SUC 95			
Staphylococcus cohnii subsp. urealyticum		15.42	0.39				
261613	**Low discrimination**						**261613**
					NOV		
Staphylococcus cohnii subsp. urealyticum		86.08	0.41	SUC 3	99		
Aerococcus viridans		6.38	0.29		1		
Staphylococcus simulans		4.15	0.09		1		
261633	**Low discrimination**						**261633**
					NOV	PYR	
Staphylococcus equorum		59.72	0.31	NIT 92	99	5	
Staphylococcus xylosus		34.61	0.25	GAL 91	99	65	
Aerococcus viridans		5.66	0.29	MLT 91	1	95	
261640	**Good identification**						**261640**
Staphylococcus warneri		100.0	0.41	ESL 1			
261642	**Low discrimination**						**261642**
					PYR		
Staphylococcus warneri		82.31	0.31	ESL 1	8		
Aerococcus viridans		17.69	0.24		95		
261643	**Low discrimination**						**261643**
					NOV		
Staphylococcus cohnii subsp. urealyticum		80.92	0.36	SUC 3	99		
Aerococcus viridans		14.26	0.32	MAN 91	1		
Staphylococcus xylosus		4.82	0.02		99		
261650	**Low discrimination**						**261650**
					NOV		
Staphylococcus warneri		84.24	0.38	ESL 1	1		
Staphylococcus kloosii		9.50	0.26		99		
Staphylococcus saprophyticus subsp. saprophyticus		6.26	-.06		99		
261651	**Low discrimination**						**261651**
					NOV		
Staphylococcus cohnii subsp. urealyticum		69.35	0.36	SUC 3	99		
Staphylococcus xylosus		22.74	0.17	LAC 91	99		
Aerococcus viridans		4.08	0.22		1		
261653	**Low discrimination**						**261653**
					NOV		
Staphylococcus cohnii subsp. urealyticum		86.02	0.57	SUC 3	99		
Staphylococcus xylosus		7.94	0.26		99		
Aerococcus viridans		4.50	0.42		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
261671	**Low discrimination**							**261671**
					NOV	PYR		
Staphylococcus equorum		58.87	0.44	NIT 92	99	5		
Staphylococcus xylosus		40.50	0.39	GAL 91	99	65		
Aerococcus viridans		0.63	0.22		1	95		
261672	**Low discrimination**							**261672**
					NOV	YEP	PYR	
Staphylococcus arlettae		69.63	0.37	URE 1	99	99	1	
Aerococcus viridans		14.42	0.34		1	1	95	
Staphylococcus equorum		9.06	0.11		99	1	5	
261673	**Low discrimination**							**261673**
					YEP			
Staphylococcus equorum		52.38	0.52	NIT 92	1			
Staphylococcus xylosus		39.89	0.48	GAL 91	40			
Staphylococcus arlettae		5.39	0.40		99			
261713	**Low discrimination**							**261713**
Staphylococcus cohnii subsp. urealyticum		43.71	0.34	SUC 3				
Staphylococcus xylosus		40.60	0.24					
Staphylococcus equorum		8.14	0.11					
261731	**Low discrimination**							**261731**
Staphylococcus equorum		65.63	0.44	NIT 92				
Staphylococcus xylosus		34.37	0.36	MLT 91				
261733	**Low discrimination**							**261733**
					NOV	PYR		
Staphylococcus equorum		62.70	0.52	NIT 92	99	5		
Staphylococcus xylosus		36.34	0.46	MLT 91	99	65		
Aerococcus viridans		0.59	0.29		1	95		
261740	**Good identification**							**261740**
Staphylococcus warneri		100.0	0.30	ESL 1				
261750	**Low discrimination**							**261750**
					NOV	PYR		
Staphylococcus warneri		48.08	0.28	ESL 1	1	8		
Staphylococcus kloosii		33.32	0.32		99	52		
Staphylococcus saprophyticus subsp. bovis		11.92	0.05		99	99		
261751	**Low discrimination**							**261751**
Staphylococcus xylosus		76.00	0.38	NIT 89				
Staphylococcus equorum		12.83	0.23	XYL 92				
Staphylococcus cohnii subsp. urealyticum		9.82	0.29					
261752	**Low discrimination**							**261752**
					NOV	YEP		
Aerococcus viridans		32.86	0.34	URE 1	1	1		
Staphylococcus kloosii		26.24	0.33		99	99		
Staphylococcus xylosus		13.82	0.06		99	40		
261753	**Low discrimination**							**261753**
Staphylococcus xylosus		59.65	0.47	NIT 89				
Staphylococcus cohnii subsp. urealyticum		27.39	0.49	XYL 92				
Staphylococcus equorum		9.10	0.32					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
261761	**Low discrimination**						**261761**
Staphylococcus xylosus		81.30	0.35	NIT 89			
Staphylococcus equorum		18.70	0.23	MAN 94			
261763	**Low discrimination**						**261763**
					NOV	PYR	
Staphylococcus xylosus		80.02	0.44	NIT 89	99	65	
Staphylococcus equorum		16.63	0.32	MAN 94	99	5	
Aerococcus viridans		2.04	0.32		1	95	
261771	**Low discrimination**						**261771**
Staphylococcus equorum		59.07	0.64	NIT 92			
Staphylococcus xylosus		40.64	0.60				
Staphylococcus gallinarum		0.14	0.07				
261772	**Low discrimination**						**261772**
					YEP		
Staphylococcus arlettae		68.84	0.53	URE 1	99		
Staphylococcus equorum		15.98	0.32		1		
Staphylococcus xylosus		12.17	0.28		40		
261773	**Low discrimination**						**261773**
					YEP		
Staphylococcus equorum		54.78	0.73	NIT 92	1		
Staphylococcus xylosus		41.72	0.69		40		
Staphylococcus arlettae		3.16	0.55		99		
262000	**Low discrimination**						**262000**
					OXI		
Kocuria varians		89.16	0.46	GLR 1	13		
Kocuria rosea		5.44	0.12		35		
Dermacoccus nishinomiyaensis		5.40	0.13		93		
262002	**Good identification**						**262002**
Kocuria varians		97.08	0.34	GLR 1			
Staphylococcus simulans		2.92	0.00				
262020	**Acceptable identification**						**262020**
Kocuria varians		93.07	0.36	GLR 1			
Kocuria rosea		6.93	0.04				
262040	**Low discrimination**						**262040**
Kocuria varians		87.48	0.32	GLR 1			
Staphylococcus warneri		12.52	0.22				
262053	**Low discrimination**						**262053**
					NOV		
Staphylococcus cohnii subsp. urealyticum		84.85	0.40	TRE 99	99		
Staphylococcus simulans		11.41	0.16		1		
Staphylococcus xylosus		3.75	0.03		99		
262100	**Good identification**						**262100**
					OXI		
Kocuria varians		99.24	0.49	GLR 1	13		
Dermacoccus nishinomiyaensis		0.76	-.02		93		
262101	**Good identification**						**262101**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		100.0	0.27	GLR 1			
262102	**Good identification**						**262102**
Kocuria varians		100.0	0.38	GLR 1			
262120	**Good identification**						**262120**
Kocuria varians		100.0	0.40	GLR 1			
262122	**Good identification**						**262122**
Kocuria varians		100.0	0.28	GLR 1			
262140	**Good identification**						**262140**
Kocuria varians		97.08	0.36	GLR 1			
Staphylococcus warneri		2.92	0.11				
262150	**Good identification**						**262150**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.12	0.34	TRE 99	99	99	
Staphylococcus warneri		1.88	0.08		1	8	
262171	**Good identification**						**262171**
Staphylococcus xylosus		100.0	0.36	SUC 95			
262173	**Good identification**						**262173**
Staphylococcus xylosus		100.0	0.45	SUC 95			
262200	**Low discrimination**						**262200**
Kocuria varians		81.28	0.28	GLR 1			
Staphylococcus warneri		18.72	0.22	SUC 12			
262202	**Low discrimination**						**262202**
Staphylococcus simulans		80.62	0.29	ARG 92			
Kocuria varians		15.23	0.16	TRE 95			
Staphylococcus warneri		4.15	0.12				
262203	**Good identification**						**262203**
Staphylococcus simulans		100.0	0.40	ARG 92			
262212	**Good identification**						**262212**
Staphylococcus simulans		99.08	0.41	ARG 92			
Staphylococcus warneri		0.92	0.09	TRE 95			
262213	**Good identification**						**262213**
					NOV		
Staphylococcus simulans		99.13	0.52	ARG 92	1		
Staphylococcus xylosus		0.87	0.06	TRE 95	99		
262233	**Acceptable identification**						**262233**
					NOV		
Staphylococcus xylosus		90.93	0.28	GAL 91	99		
Staphylococcus simulans		9.07	0.11	TRE 96	1		
262240	**Acceptable identification**						**262240**
Staphylococcus warneri		93.89	0.48	TRE 95			
Kocuria varians		4.71	0.14				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		1.40	0.05				
262242	**Low discrimination**						**262242**
Staphylococcus warneri		64.31	0.38	TRE 95			
Staphylococcus simulans		29.58	0.22				
Staphylococcus hominis subsp. hominis		3.38	0.06				
262243	**Acceptable identification**						**262243**
					NOV		
Staphylococcus simulans		93.77	0.32	ARG 92	1		
Staphylococcus xylosus		6.23	0.05	TRE 95	99		
262250	**Low discrimination**						**262250**
					NOV	PYR	
Staphylococcus warneri		53.54	0.45	TRE 95	1	8	
Staphylococcus saprophyticus subsp. bovis		46.46	0.34		99	99	
262252	**Low discrimination**						**262252**
					NOV	PYR	
Staphylococcus simulans		60.61	0.34	ARG 92	1	70	
Staphylococcus warneri		23.86	0.35	TRE 95	1	8	
Staphylococcus saprophyticus subsp. bovis		15.53	0.22		99	99	
262253	**Low discrimination**						**262253**
					NOV		
Staphylococcus simulans		77.58	0.45	ARG 92	1		
Staphylococcus xylosus		20.20	0.29	TRE 95	99		
Staphylococcus saprophyticus subsp. bovis		1.48	0.09		99		
262271	**Good identification**						**262271**
Staphylococcus xylosus		100.0	0.42	GAL 91			
262273	**Good identification**						**262273**
					NOV		
Staphylococcus xylosus		99.66	0.51	GAL 91	99		
Staphylococcus simulans		0.34	0.04	TRE 96	1		
262300	**Good identification**						**262300**
Kocuria varians		95.38	0.31	GLR 1			
Staphylococcus warneri		4.62	0.11	SUC 12			
262313	**Low discrimination**						**262313**
					COA	NOV	
Staphylococcus xylosus		47.17	0.27	TRE 96	1	99	
Staphylococcus simulans		46.08	0.30	XYL 92	1	1	
Staphylococcus intermedius		6.75	0.11	MLT 93	99	1	
262331	**Good identification**						**262331**
Staphylococcus xylosus		100.0	0.39	TRE 96			
262333	**Good identification**						**262333**
Staphylococcus xylosus		100.0	0.49	TRE 96			
262340	**Low discrimination**						**262340**
					NOV		
Staphylococcus warneri		68.93	0.38	TRE 95	1		
Kocuria varians		16.42	0.18		1		
Staphylococcus hominis subsp. novobiosepticus		5.58	-.02		99		
262342	**Low discrimination**						**262342**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		61.00	0.28	TRE 95			
Kocuria varians		12.30	0.06				
Staphylococcus hominis subsp. hominis		11.01	0.07				
262350	**Acceptable identification**						**262350**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		91.03	0.47	TRE 99	99	99	
Staphylococcus warneri		8.28	0.35		1	8	
Staphylococcus xylosus		0.69	-.01		99	65	
262351	**Low discrimination**						**262351**
					COA	NOV	
Staphylococcus xylosus		74.71	0.40	TRE 96	1	99	
Staphylococcus saprophyticus subsp. bovis		24.73	0.34	XYL 92	1	99	
Staphylococcus intermedius		0.56	-.02		99	1	
262352	**Low discrimination**						**262352**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		77.34	0.34	TRE 99	99	99	
Staphylococcus warneri		9.38	0.25		1	8	
Staphylococcus xylosus		6.72	0.09		99	65	
262353	**Acceptable identification**						**262353**
					NOV		
Staphylococcus xylosus		92.74	0.50	TRE 96	99		
Staphylococcus simulans		3.06	0.23	XYL 92	1		
Staphylococcus saprophyticus subsp. bovis		2.70	0.22		99		
262361	**Good identification**						**262361**
Staphylococcus xylosus		100.0	0.38	TRE 96			
262363	**Good identification**						**262363**
Staphylococcus xylosus		100.0	0.47	TRE 96			
262371	**Very good identification**						**262371**
Staphylococcus xylosus		99.87	0.62	TRE 96			
Staphylococcus equorum		0.13	0.04				
262372	**Good identification**						**262372**
Staphylococcus xylosus		100.0	0.31	TRE 96			
262373	**Very good identification**						**262373**
Staphylococcus xylosus		99.89	0.72	TRE 96			
Staphylococcus equorum		0.11	0.12				
262402	**Low discrimination**						**262402**
Staphylococcus simulans		83.57	0.27	ARG 92			
Kocuria varians		11.01	0.11	SUC 96			
Staphylococcus warneri		5.43	0.12				
262403	**Low discrimination**						**262403**
					NOV		
Staphylococcus cohnii subsp. urealyticum		54.87	0.45	MAN 91	99		
Staphylococcus simulans		45.13	0.37		1		
262411	**Acceptable identification**						**262411**
					NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		92.15	0.45	LAC 91	99		
Staphylococcus saprophyticus subsp. bovis		3.29	0.09		99		
Staphylococcus simulans		3.06	0.09		1		
262412	**Low discrimination**						**262412**
					NOV		
Staphylococcus simulans		86.80	0.39	ARG 92	1		
Staphylococcus cohnii subsp. urealyticum		9.02	0.24	SUC 96	99		
Staphylococcus saprophyticus subsp. bovis		3.16	0.09		99		
262413	**Low discrimination**						**262413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		75.19	0.65		99		
Staphylococcus simulans		24.47	0.50		1		
Staphylococcus xylosus		0.34	0.08		99		
262433	**Low discrimination**						**262433**
					NOV		
Staphylococcus xylosus		79.70	0.30	GAL 91	99		
Staphylococcus cohnii subsp. urealyticum		15.32	0.24	SUC 95	99		
Staphylococcus simulans		4.98	0.09	MLT 93	1		
262440	**Low discrimination**						**262440**
					NOV	PYR	
Staphylococcus warneri		82.77	0.48	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		13.00	0.22		99	99	
Kocuria varians		2.29	0.09		1	10	
262441	**Low discrimination**						**262441**
					NOV	PYR	
Staphylococcus cohnii subsp. urealyticum		89.17	0.40	MAN 91	99	53	
Staphylococcus saprophyticus subsp. bovis		5.68	0.09	LAC 91	99	99	
Staphylococcus warneri		2.95	0.13		1	8	
262442	**Low discrimination**						**262442**
					NOV		
Staphylococcus warneri		59.86	0.38	SUC 95	1		
Staphylococcus simulans		21.79	0.19		1		
Staphylococcus cohnii subsp. urealyticum		11.30	0.19		99		
262443	**Good identification**						**262443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.27	0.60	MAN 91	99		
Staphylococcus simulans		6.08	0.30		1		
Staphylococcus xylosus		0.65	0.07		99		
262450	**Very good identification**						**262450**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		95.18	0.63		99	99	
Staphylococcus warneri		4.43	0.45		1	8	
Staphylococcus cohnii subsp. urealyticum		0.39	0.19		99	53	
262451	**Low discrimination**						**262451**
Staphylococcus cohnii subsp. urealyticum		59.98	0.60	LAC 91			
Staphylococcus saprophyticus subsp. bovis		37.44	0.50				
Staphylococcus xylosus		2.28	0.22				
262452	**Low discrimination**						**262452**
					NOV		
Staphylococcus saprophyticus subsp. bovis		73.85	0.50		99		
Staphylococcus cohnii subsp. urealyticum		12.09	0.40		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		9.22	0.32		1		
262453	**Very good identification**						**262453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		95.06	0.81		99		
Staphylococcus simulans		2.45	0.43		1		
Staphylococcus saprophyticus subsp. bovis		1.47	0.38		99		
262463	**Low discrimination**						**262463**
Staphylococcus xylosus		88.73	0.28	GAL 91			
Staphylococcus cohnii subsp. urealyticum		11.27	0.19	SUC 95			
262471	**Good identification**						**262471**
Staphylococcus xylosus		96.39	0.44	GAL 91			
Staphylococcus cohnii subsp. urealyticum		2.22	0.19	SUC 95			
Staphylococcus saprophyticus subsp. bovis		1.39	0.09				
262473	**Good identification**						**262473**
					NOV		
Staphylococcus xylosus		92.24	0.53	GAL 91	99		
Staphylococcus cohnii subsp. urealyticum		7.56	0.40	SUC 95	99		
Staphylococcus simulans		0.20	0.02		1		
262503	**Low discrimination**						**262503**
					NOV		
Staphylococcus cohnii subsp. urealyticum		80.88	0.37	MAN 91	99		
Staphylococcus simulans		13.50	0.16		1		
Staphylococcus xylosus		5.63	0.04		99		
262510	**Good identification**						**262510**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.12	0.34	MLT 99	99	99	
Staphylococcus warneri		1.88	0.08		1	8	
262511	**Low discrimination**						**262511**
Staphylococcus cohnii subsp. urealyticum		58.26	0.37	LAC 91			
Staphylococcus xylosus		22.32	0.19				
Staphylococcus saprophyticus subsp. bovis		19.42	0.22				
262513	**Low discrimination**						**262513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		84.60	0.58		99		
Staphylococcus xylosus		9.12	0.29		99		
Staphylococcus simulans		5.59	0.28		1		
262531	**Good identification**						**262531**
Staphylococcus xylosus		100.0	0.41	SUC 95			
262533	**Good identification**						**262533**
Staphylococcus xylosus		99.19	0.51	SUC 95			
Staphylococcus cohnii subsp. urealyticum		0.81	0.17	MLT 93			
262540	**Low discrimination**						**262540**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		63.75	0.34	MAN 99	99	99	
Staphylococcus warneri		32.04	0.38		1	8	
Kocuria varians		4.21	0.12		1	10	
262541	**Low discrimination**						**262541**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		45.99	0.32	MAN 91			
Staphylococcus saprophyticus subsp. bovis		27.36	0.22	LAC 91			
Staphylococcus xylosus		26.66	0.18				
262543	**Low discrimination**						**262543**
Staphylococcus cohnii subsp. urealyticum		83.96	0.53	MAN 91			
Staphylococcus xylosus		13.70	0.27				
Staphylococcus saprophyticus subsp. bovis		1.24	0.09				
262550	**Very good identification**						**262550**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		99.55	0.75		99	99	
Staphylococcus warneri		0.37	0.35		1	8	
Staphylococcus cohnii subsp. urealyticum		0.04	0.12		99	53	
262551	**Low discrimination**						**262551**
Staphylococcus saprophyticus subsp. bovis		75.42	0.63				
Staphylococcus cohnii subsp. urealyticum		12.95	0.53				
Staphylococcus xylosus		11.63	0.43				
262552	**Very good identification**						**262552**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		97.13	0.63		99	99	
Staphylococcus cohnii subsp. urealyticum		1.70	0.32		99	53	
Staphylococcus warneri		0.48	0.25		1	8	
262553	**Low discrimination**						**262553**
Staphylococcus cohnii subsp. urealyticum		71.32	0.73				
Staphylococcus xylosus		18.03	0.52				
Staphylococcus saprophyticus subsp. bovis		10.27	0.50				
262561	**Good identification**						**262561**
Staphylococcus xylosus		100.0	0.40	SUC 95			
262563	**Good identification**						**262563**
					COA	NOV	
Staphylococcus xylosus		96.98	0.49	SUC 95	1	99	
Staphylococcus lutrae		2.50	0.12	MAN 94	99	1	
Staphylococcus cohnii subsp. urealyticum		0.52	0.12		1	99	
262570	**Low discrimination**						**262570**
Staphylococcus saprophyticus subsp. bovis		69.29	0.34	XYL 1			
Staphylococcus xylosus		30.72	0.23				
262571	**Very good identification**						**262571**
Staphylococcus xylosus		99.24	0.64	SUC 95			
Staphylococcus saprophyticus subsp. bovis		0.57	0.22				
Staphylococcus cohnii subsp. urealyticum		0.10	0.12				
262572	**Low discrimination**						**262572**
Staphylococcus xylosus		83.46	0.33	SUC 95			
Staphylococcus saprophyticus subsp. bovis		16.54	0.22	MNS 99			
262573	**Very good identification**						**262573**
					COA	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		99.20	0.74	SUC 95	1	99	
Staphylococcus cohnii subsp. urealyticum		0.35	0.32		1	99	
Staphylococcus lutrae		0.32	0.18		99	1	
262600	**Low discrimination**						**262600**
					YEP		
Staphylococcus warneri		89.76	0.48	MLT 95	43		
Staphylococcus simulans		5.88	0.14		1		
Staphylococcus pasteuri		2.82	0.02		86		
262602	**Good identification**						**262602**
					YEP		
Staphylococcus simulans		94.93	0.55	ARG 92	1		
Staphylococcus warneri		4.88	0.38		43		
Staphylococcus pasteuri		0.19	-.06		86		
262603	**Very good identification**						**262603**
					NOV		
Staphylococcus simulans		99.21	0.66	ARG 92	1		
Staphylococcus hyicus		0.35	0.08		1		
Staphylococcus xylosus		0.28	0.10		99		
262610	**Low discrimination**						**262610**
					NOV	PYR	
Staphylococcus warneri		40.06	0.45	MLT 95	1	8	
Staphylococcus saprophyticus subsp. bovis		34.77	0.34		99	99	
Staphylococcus simulans		14.49	0.26		1	70	
262611	**Low discrimination**						**262611**
					NOV		
Staphylococcus simulans		61.67	0.37	ARG 92	1		
Staphylococcus xylosus		23.86	0.25	LAC 99	99		
Staphylococcus saprophyticus subsp. bovis		11.05	0.22		99		
262612	**Very good identification**						**262612**
					NOV	PYR	
Staphylococcus simulans		98.19	0.67	ARG 92	1	70	
Staphylococcus warneri		0.91	0.35		1	8	
Staphylococcus saprophyticus subsp. bovis		0.60	0.22		99	99	
262613	**Very good identification**						**262613**
					NOV		
Staphylococcus simulans		98.47	0.78	ARG 92	1		
Staphylococcus xylosus		1.10	0.34		99		
Staphylococcus cohnii subsp. urealyticum		0.39	0.34		99		
262623	**Low discrimination**						**262623**
					NOV		
Staphylococcus xylosus		76.37	0.32	GAL 91	99		
Staphylococcus simulans		23.63	0.25	MAN 94	1		
262631	**Good identification**						**262631**
Staphylococcus xylosus		99.32	0.47	GAL 91			
Staphylococcus equorum		0.68	0.04	MLT 93			
262633	**Good identification**						**262633**
					NOV		
Staphylococcus xylosus		92.15	0.56	GAL 91	99		
Staphylococcus simulans		7.28	0.37	MLT 93	1		
Staphylococcus equorum		0.57	0.12		99		
262640	**Very good identification**						**262640**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Staphylococcus warneri		95.38	0.74		1	8	
Staphylococcus saprophyticus subsp. bovis		3.16	0.34		99	99	
Staphylococcus pasteuri		0.97	0.19		1	1	
262641	**Low discrimination**						**262641**
					NOV	PYR	
Staphylococcus warneri		40.81	0.39	MNS 2	1	8	
Staphylococcus xylosus		30.28	0.24		99	65	
Staphylococcus saprophyticus subsp. bovis		16.54	0.22		99	99	
262642	**Low discrimination**						**262642**
					NOV	PYR	
Staphylococcus warneri		66.41	0.64		1	8	
Staphylococcus simulans		30.54	0.48		1	70	
Staphylococcus saprophyticus subsp. bovis		1.65	0.22		99	99	
262643	**Low discrimination**						**262643**
					NOV		
Staphylococcus simulans		89.21	0.59	ARG 92	1		
Staphylococcus xylosus		7.49	0.33		99		
Staphylococcus cohnii subsp. urealyticum		1.76	0.29		99		
262650	**Low discrimination**						**262650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		80.44	0.75		99	99	
Staphylococcus warneri		17.79	0.71		1	8	
Staphylococcus pasteuri		1.53	0.35		1	1	
262651	**Low discrimination**						**262651**
					NOV		
Staphylococcus saprophyticus subsp. bovis		74.32	0.63		99		
Staphylococcus xylosus		21.53	0.48		99		
Staphylococcus simulans		1.88	0.30		1		
262652	**Low discrimination**						**262652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		47.77	0.63		99	99	
Staphylococcus simulans		35.78	0.60		1	70	
Staphylococcus warneri		14.09	0.62		1	8	
262653	**Low discrimination**						**262653**
					NOV		
Staphylococcus simulans		67.53	0.71	ARG 92	1		
Staphylococcus xylosus		22.21	0.58		99		
Staphylococcus saprophyticus subsp. bovis		6.73	0.50		99		
262660	**Low discrimination**						**262660**
					NOV		
Staphylococcus warneri		85.17	0.33	XYL 1	1		
Staphylococcus xylosus		14.83	0.04		99		
262661	**Good identification**						**262661**
Staphylococcus xylosus		100.0	0.45	GAL 91			
262663	**Very good identification**						**262663**
					NOV		
Staphylococcus xylosus		98.97	0.55	GAL 91	99		
Staphylococcus simulans		1.04	0.18	MAN 94	1		
262670	**Low discrimination**						**262670**
					NOV	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		48.68	0.34	XYL 1	99	99	
Staphylococcus xylosus		40.55	0.29		99	65	
Staphylococcus warneri		10.77	0.30		1	8	
262671	**Very good identification**						**262671**
					PYR		
Staphylococcus xylosus		99.18	0.70	GAL 91	65		
Staphylococcus equorum		0.52	0.24		5		
Staphylococcus saprophyticus subsp. bovis		0.30	0.22		99		
262672	**Low discrimination**						**262672**
					NOV		
Staphylococcus xylosus		82.27	0.38	GAL 91	99		
Staphylococcus saprophyticus subsp. bovis		8.68	0.22	MNS 99	99		
Staphylococcus simulans		6.50	0.19		1		
262673	**Very good identification**						**262673**
					NOV		
Staphylococcus xylosus		99.23	0.79	GAL 91	99		
Staphylococcus equorum		0.47	0.33		99		
Staphylococcus simulans		0.27	0.30		1		
262700	**Low discrimination**						**262700**
					NOV	PYR	
Staphylococcus warneri		86.05	0.38	MLT 95	1	8	
Kocuria varians		7.03	0.08		1	10	
Staphylococcus saprophyticus subsp. bovis		6.92	0.05		99	99	
262702	**Low discrimination**						**262702**
Staphylococcus simulans		84.27	0.33	ARG 92			
Staphylococcus warneri		15.74	0.28	GAL 8			
262703	**Low discrimination**						**262703**
					NOV		
Staphylococcus simulans		51.01	0.44	ARG 92	1		
Staphylococcus hyicus		24.03	0.30	GAL 8	1		
Staphylococcus xylosus		16.84	0.30		99		
262710	**Low discrimination**						**262710**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		89.78	0.47	MLT 99	99	99	
Staphylococcus warneri		8.17	0.35		1	8	
Staphylococcus xylosus		1.24	0.05		99	65	
262711	**Low discrimination**						**262711**
					COA	NOV	
Staphylococcus xylosus		80.23	0.46	XYL 92	1	99	
Staphylococcus saprophyticus subsp. bovis		14.70	0.34	MLT 93	1	99	
Staphylococcus intermedius		2.94	0.18		99	1	
262712	**Low discrimination**						**262712**
					NOV		
Staphylococcus simulans		73.73	0.46	ARG 92	1		
Staphylococcus saprophyticus subsp. bovis		20.55	0.34	GAL 8	99		
Staphylococcus xylosus		3.22	0.14		99		
262713	**Low discrimination**						**262713**
					COA	NOV	
Staphylococcus xylosus		53.02	0.55	XYL 92	1	99	
Staphylococcus simulans		41.00	0.56	MLT 93	1	1	
Staphylococcus intermedius		4.20	0.34		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
262721	**Good identification**						**262721**
Staphylococcus xylosus		100.0	0.43	MAN 94			
262723	**Good identification**						**262723**
					NOV		
Staphylococcus xylosus		99.74	0.52	MAN 94	99		
Staphylococcus simulans		0.27	0.03	MLT 93	1		
262730	**Acceptable identification**						**262730**
Staphylococcus xylosus		94.01	0.26	MLT 93			
Staphylococcus saprophyticus subsp. bovis		5.99	0.05	MNS 99			
262731	**Very good identification**						**262731**
Staphylococcus xylosus		99.32	0.68	MLT 93			
Staphylococcus equorum		0.68	0.24				
262732	**Good identification**						**262732**
					NOV		
Staphylococcus xylosus		98.03	0.36	MLT 93	99		
Staphylococcus simulans		1.97	0.05	MNS 99	1		
262733	**Very good identification**						**262733**
					NOV		
Staphylococcus xylosus		99.32	0.77	MLT 93	99		
Staphylococcus equorum		0.62	0.33		99		
Staphylococcus simulans		0.07	0.15		1		
262740	**Low discrimination**						**262740**
					NOV	PYR	
Staphylococcus warneri		69.81	0.64		1	8	
Staphylococcus saprophyticus subsp. bovis		29.25	0.47		99	99	
Staphylococcus hominis subsp. hominis		0.60	0.16		1	1	
262741	**Low discrimination**						**262741**
					NOV	PYR	
Staphylococcus xylosus		77.99	0.44	MAN 94	99	65	
Staphylococcus saprophyticus subsp. bovis		16.85	0.34	XYL 92	99	99	
Staphylococcus warneri		3.28	0.29		1	8	
262742	**Low discrimination**						**262742**
					NOV	PYR	
Staphylococcus warneri		66.11	0.54		1	8	
Staphylococcus saprophyticus subsp. bovis		20.77	0.34		99	99	
Staphylococcus simulans		8.37	0.26		1	70	
262743	**Low discrimination**						**262743**
					COA	NOV	
Staphylococcus xylosus		84.17	0.54	MAN 94	1	99	
Staphylococcus simulans		8.62	0.37	XYL 92	1	1	
Staphylococcus intermedius		4.36	0.29		99	1	
262750	**Very good identification**						**262750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.11	0.88		99	99	
Staphylococcus warneri		1.71	0.61		1	8	
Staphylococcus xylosus		0.18	0.28		99	65	
262751	**Low discrimination**						**262751**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		57.50	0.75		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		42.11	0.69		1	99	
Staphylococcus intermedius		0.17	0.22		99	1	
262752	**Good identification**						**262752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		94.29	0.75		99	99	
Staphylococcus warneri		2.20	0.51		1	8	
Staphylococcus xylosus		1.99	0.37		99	65	
262753	**Low discrimination**						**262753**
					NOV		
Staphylococcus xylosus		86.04	0.78	XYL 92	99		
Staphylococcus saprophyticus subsp. bovis		10.32	0.63		99		
Staphylococcus simulans		2.25	0.49		1		
262761	**Very good identification**						**262761**
Staphylococcus xylosus		99.92	0.66	MAN 94			
Staphylococcus equorum		0.08	0.04				
262762	**Good identification**						**262762**
					NOV		
Staphylococcus xylosus		97.94	0.34	MAN 94	99		
Staphylococcus warneri		2.06	0.13	MNS 99	1		
262763	**Very good identification**						**262763**
					COA	NOV	
Staphylococcus xylosus		99.78	0.75	MAN 94	1	99	
Staphylococcus lutrae		0.14	0.12		99	1	
Staphylococcus equorum		0.07	0.12		1	99	
262770	**Low discrimination**						**262770**
					NOV	PYR	
Staphylococcus xylosus		67.19	0.50	MNS 99	99	65	
Staphylococcus saprophyticus subsp. bovis		31.91	0.47		99	99	
Staphylococcus warneri		0.56	0.20		1	8	
262771	**Very good identification**						**262771**
					PYR		
Staphylococcus xylosus		99.35	0.91		65		
Staphylococcus equorum		0.52	0.45		5		
Staphylococcus saprophyticus subsp. bovis		0.12	0.34		99		
262772	**Very good identification**						**262772**
					PYR		
Staphylococcus xylosus		95.31	0.59	MNS 99	65		
Staphylococcus saprophyticus subsp. bovis		3.98	0.34		99		
Staphylococcus equorum		0.45	0.12		5		
262773	**Excellent identification**						**262773**
					COA	NOV	
Staphylococcus xylosus		99.49	1.00		1	99	
Staphylococcus equorum		0.47	0.54		1	99	
Staphylococcus lutrae		0.02	0.18		99	1	
263053	**Good identification**						**263053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99			
263173	**Acceptable identification**						**263173**
Staphylococcus xylosus		92.01	0.33	SUC 95			
Staphylococcus equorum		7.99	0.12	TRE 96			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
263271	**Low discrimination**					**263271**
Staphylococcus xylosus		66.63	0.29	GAL 91		
Staphylococcus equorum		33.37	0.24	TRE 96		
263273	**Low discrimination**					**263273**
Staphylococcus xylosus		68.85	0.39	GAL 91		
Staphylococcus equorum		31.15	0.33	TRE 96		
263331	**Low discrimination**					**263331**
Staphylococcus xylosus		60.31	0.27	TRE 96		
Staphylococcus equorum		39.69	0.24	MLT 93		
263333	**Low discrimination**					**263333**
Staphylococcus xylosus		62.71	0.36	TRE 96		
Staphylococcus equorum		37.29	0.33	MLT 93		
263351	**Acceptable identification**					**263351**
Staphylococcus xylosus		94.50	0.28	TRE 96		
Staphylococcus equorum		5.50	0.04	XYL 92		
263353	**Good identification**					**263353**
Staphylococcus xylosus		95.01	0.37	TRE 96		
Staphylococcus equorum		4.99	0.12	XYL 92		
263363	**Acceptable identification**					**263363**
Staphylococcus xylosus		92.17	0.35	TRE 96		
Staphylococcus equorum		6.60	0.12	MAN 94		
Staphylococcus gallinarum		1.23	-.05			
263371	**Low discrimination**					**263371**
Staphylococcus xylosus		66.19	0.50	TRE 96		
Staphylococcus equorum		33.15	0.45			
Staphylococcus gallinarum		0.66	0.07			
263373	**Low discrimination**					**263373**
Staphylococcus xylosus		68.44	0.59	TRE 96		
Staphylococcus equorum		30.97	0.54			
Staphylococcus gallinarum		0.59	0.15			
263403	**Good identification**					**263403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91		
263411	**Good identification**					**263411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	LAC 91		
263413	**Very good identification**					**263413**
					NOV	
Staphylococcus cohnii subsp. urealyticum		98.87	0.57		99	
Staphylococcus simulans		0.84	0.09		1	
Staphylococcus xylosus		0.29	-.04		99	
263441	**Good identification**					**263441**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	MAN 91		
263443	**Good identification**					**263443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	MAN 91		
263451	**Good identification**					**263451**
Staphylococcus cohnii subsp. urealyticum		96.09	0.52	LAC 91		
Staphylococcus xylosus		2.35	0.09			
Staphylococcus saprophyticus subsp. bovis		1.56	0.09			
263452	**Low discrimination**					**263452**
Staphylococcus cohnii subsp. urealyticum		86.30	0.31	MNS 99		
Staphylococcus saprophyticus subsp. bovis		13.70	0.09			
263453	**Very good identification**					**263453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.25	0.72		99	
Staphylococcus xylosus		0.68	0.19		99	
Staphylococcus simulans		0.07	0.02		1	
263471	**Low discrimination**					**263471**
Staphylococcus xylosus		69.81	0.31	GAL 91		
Staphylococcus equorum		27.68	0.24	SUC 95		
Staphylococcus cohnii subsp. urealyticum		2.51	0.11			
263473	**Low discrimination**					**263473**
Staphylococcus xylosus		67.31	0.41	GAL 91		
Staphylococcus equorum		24.11	0.33	SUC 95		
Staphylococcus cohnii subsp. urealyticum		8.58	0.31			
263503	**Good identification**					**263503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.29	MAN 91		
263511	**Low discrimination**					**263511**
Staphylococcus cohnii subsp. urealyticum		80.24	0.29	LAC 91		
Staphylococcus xylosus		19.76	0.07			
263513	**Acceptable identification**					**263513**
Staphylococcus cohnii subsp. urealyticum		93.52	0.49			
Staphylococcus xylosus		6.48	0.16			
263531	**Low discrimination**					**263531**
Staphylococcus xylosus		65.75	0.29	SUC 95		
Staphylococcus equorum		34.25	0.24	MLT 93		
263533	**Low discrimination**					**263533**
Staphylococcus xylosus		67.41	0.38	SUC 95		
Staphylococcus equorum		31.73	0.33	MLT 93		
Staphylococcus cohnii subsp. urealyticum		0.85	0.08			
263543	**Acceptable identification**					**263543**
Staphylococcus cohnii subsp. urealyticum		90.51	0.44	MAN 91		
Staphylococcus xylosus		9.49	0.15			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
263550	**Good identification**						**263550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ESL 1			
263551	**Low discrimination**						**263551**
Staphylococcus cohnii subsp. urealyticum		56.97	0.44	LAC 91			
Staphylococcus xylosus		32.90	0.30				
Staphylococcus saprophyticus subsp. bovis		8.62	0.22				
263553	**Low discrimination**						**263553**
Staphylococcus cohnii subsp. urealyticum		85.25	0.65				
Staphylococcus xylosus		13.86	0.40				
Staphylococcus equorum		0.58	0.12				
263561	**Acceptable identification**						**263561**
Staphylococcus xylosus		94.10	0.27	SUC 95			
Staphylococcus equorum		5.90	0.04	MAN 94			
263563	**Acceptable identification**						**263563**
Staphylococcus xylosus		93.70	0.37	SUC 95			
Staphylococcus equorum		5.31	0.12	MAN 94			
Staphylococcus gallinarum		0.99	-.05				
263571	**Low discrimination**						**263571**
Staphylococcus xylosus		71.20	0.52	SUC 95			
Staphylococcus equorum		28.23	0.45				
Staphylococcus gallinarum		0.57	0.07				
263573	**Low discrimination**						**263573**
Staphylococcus xylosus		72.97	0.61	SUC 95			
Staphylococcus equorum		26.14	0.54				
Staphylococcus gallinarum		0.50	0.15				
263613	**Low discrimination**						**263613**
					NOV		
Staphylococcus simulans		66.40	0.37	ARG 92	1		
Staphylococcus xylosus		18.28	0.22	ESL 1	99		
Staphylococcus cohnii subsp. urealyticum		10.12	0.26		99		
263631	**Low discrimination**						**263631**
Staphylococcus equorum		73.08	0.45	GAL 91			
Staphylococcus xylosus		26.92	0.34	MLT 91			
263633	**Low discrimination**						**263633**
Staphylococcus equorum		71.04	0.53	GAL 91			
Staphylococcus xylosus		28.96	0.44	MLT 91			
263640	**Good identification**						**263640**
Staphylococcus warneri		100.0	0.33	ESL 1			
263650	**Low discrimination**						**263650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		81.89	0.34	ESL 1	99	99	
Staphylococcus warneri		18.11	0.30		1	8	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
263651	**Low discrimination**					**263651**
					PYR	
Staphylococcus xylosus		69.16	0.36	GAL 91	65	
Staphylococcus equorum		16.60	0.24	XYL 92	5	
Staphylococcus saprophyticus subsp. bovis		9.64	0.22		99	
263653	**Low discrimination**					**263653**
Staphylococcus xylosus		62.96	0.45	GAL 91		
Staphylococcus cohnii subsp. urealyticum		14.88	0.41	XYL 92		
Staphylococcus equorum		13.65	0.33			
263661	**Low discrimination**					**263661**
Staphylococcus xylosus		75.36	0.33	GAL 91		
Staphylococcus equorum		24.64	0.24	MAN 94		
263663	**Low discrimination**					**263663**
Staphylococcus xylosus		76.86	0.42	GAL 91		
Staphylococcus equorum		22.70	0.33	MAN 94		
Staphylococcus gallinarum		0.43	-.05			
263671	**Low discrimination**					**263671**
Staphylococcus equorum		67.29	0.66	GAL 91		
Staphylococcus xylosus		32.57	0.58			
Staphylococcus gallinarum		0.14	0.07			
263672	**Low discrimination**					**263672**
Staphylococcus equorum		65.12	0.33	GAL 91		
Staphylococcus xylosus		34.89	0.26	MNS 99		
263673	**Low discrimination**					**263673**
Staphylococcus equorum		65.03	0.74	GAL 91		
Staphylococcus xylosus		34.84	0.67			
Staphylococcus gallinarum		0.13	0.15			
263711	**Low discrimination**					**263711**
Staphylococcus xylosus		76.03	0.33	XYL 92		
Staphylococcus equorum		23.97	0.24	MLT 93		
263713	**Low discrimination**					**263713**
					NOV	
Staphylococcus xylosus		73.79	0.43	XYL 92	99	
Staphylococcus equorum		21.02	0.33	MLT 93	99	
Staphylococcus simulans		2.31	0.15		1	
263721	**Low discrimination**					**263721**
Staphylococcus xylosus		69.95	0.31	MAN 94		
Staphylococcus equorum		30.05	0.24	MLT 93		
263723	**Low discrimination**					**263723**
Staphylococcus xylosus		71.66	0.40	MAN 94		
Staphylococcus equorum		27.81	0.33	MLT 93		
Staphylococcus gallinarum		0.53	-.05			
263731	**Low discrimination**					**263731**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus equorum		72.97	0.66	MLT 91			
Staphylococcus xylosus		26.88	0.55				
Staphylococcus gallinarum		0.15	0.07				
263732	**Low discrimination**						**263732**
Staphylococcus equorum		71.04	0.33	MLT 91			
Staphylococcus xylosus		28.96	0.23	MNS 99			
263733	**Low discrimination**						**263733**
Staphylococcus equorum		70.91	0.74	MLT 91			
Staphylococcus xylosus		28.91	0.65				
Staphylococcus gallinarum		0.14	0.15				
263741	**Good identification**						**263741**
Staphylococcus xylosus		96.34	0.32	MAN 94			
Staphylococcus equorum		3.66	0.04	XYL 92			
263743	**Acceptable identification**						**263743**
Staphylococcus xylosus		94.68	0.41	MAN 94			
Staphylococcus equorum		3.25	0.12	XYL 92			
Staphylococcus cohnii subsp. urealyticum		1.47	0.13				
263750	**Acceptable identification**						**263750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		93.09	0.47	ESL 1	99	99	
Staphylococcus xylosus		4.26	0.15		99	65	
Staphylococcus warneri		1.63	0.20		1	8	
263751	**Low discrimination**						**263751**
					PYR		
Staphylococcus xylosus		76.65	0.56	XYL 92	65		
Staphylococcus equorum		18.39	0.45		5		
Staphylococcus saprophyticus subsp. bovis		4.23	0.34		99		
263752	**Low discrimination**						**263752**
					PYR		
Staphylococcus saprophyticus subsp. bovis		60.34	0.34	ESL 1	99		
Staphylococcus xylosus		31.44	0.25		65		
Staphylococcus equorum		6.82	0.12		5		
263753	**Low discrimination**						**263753**
Staphylococcus xylosus		80.69	0.66	XYL 92			
Staphylococcus equorum		17.50	0.54				
Staphylococcus cohnii subsp. urealyticum		0.81	0.34				
263761	**Low discrimination**						**263761**
Staphylococcus xylosus		71.89	0.54	MAN 94			
Staphylococcus equorum		23.50	0.45				
Staphylococcus gallinarum		4.61	0.28				
263763	**Low discrimination**						**263763**
Staphylococcus xylosus		74.04	0.63	MAN 94			
Staphylococcus equorum		21.87	0.54				
Staphylococcus gallinarum		4.09	0.36				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
263770	**Low discrimination**					**263770**
Staphylococcus equorum		66.08	0.45	MNS 99		
Staphylococcus xylosus		31.98	0.37			
Staphylococcus gallinarum		1.32	0.07			
263771	**Low discrimination**					**263771**
Staphylococcus equorum		66.48	0.86			
Staphylococcus xylosus		32.18	0.78			
Staphylococcus gallinarum		1.33	0.48			
263772	**Low discrimination**					**263772**
Staphylococcus equorum		64.16	0.54	MNS 99		
Staphylococcus xylosus		34.37	0.47			
Staphylococcus gallinarum		1.22	0.15			
263773	**Low discrimination**					**263773**
Staphylococcus equorum		64.31	0.95			
Staphylococcus xylosus		34.45	0.88			
Staphylococcus gallinarum		1.23	0.56			
264013	**Good identification**					**264013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	TRE 99		
264043	**Good identification**					**264043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	TRE 99		
264050	**Good identification**					**264050**
Staphylococcus kloosii		100.0	0.47	TRE 99		
264051	**Good identification**					**264051**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	TRE 99		
264052	**Low discrimination**					**264052**
Staphylococcus kloosii		87.99	0.48	TRE 99		
Staphylococcus cohnii subsp. urealyticum		12.01	0.18			
264053	**Very good identification**					**264053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	TRE 99		
264070	**Good identification**					**264070**
Staphylococcus kloosii		100.0	0.34	TRE 99		
264072	**Good identification**					**264072**
Staphylococcus kloosii		100.0	0.35	TRE 99		
264112	**Good identification**					**264112**
Staphylococcus kloosii		100.0	0.27	TRE 99		
264113	**Good identification**					**264113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	TRE 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
264143	**Good identification**					**264143**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
264150	**Good identification**					**264150**
Staphylococcus kloosii		100.0	0.53	TRE 99		
264151	**Good identification**					**264151**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
264152	**Very good identification**					**264152**
Staphylococcus kloosii		97.08	0.53	TRE 99		
Staphylococcus cohnii subsp. urealyticum		2.92	0.10			
264153	**Good identification**					**264153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	TRE 99		
264170	**Good identification**					**264170**
Staphylococcus kloosii		100.0	0.40	TRE 99		
264172	**Good identification**					**264172**
Staphylococcus kloosii		100.0	0.41	TRE 99		
264252	**Good identification**					**264252**
Staphylococcus kloosii		100.0	0.27	SUC 9		
264253	**Low discrimination**					**264253**
					NOV	
Staphylococcus cohnii subsp. urealyticum		78.01	0.28	SUC 3	99	
Staphylococcus simulans		13.30	0.07	TRE 99	1	
Staphylococcus xylosus		8.69	-.01		99	
264350	**Good identification**					**264350**
Staphylococcus kloosii		100.0	0.32	SUC 9		
264352	**Good identification**					**264352**
Staphylococcus kloosii		100.0	0.33	SUC 9		
264371	**Good identification**					**264371**
Staphylococcus xylosus		100.0	0.32	NIT 89		
264373	**Good identification**					**264373**
Staphylococcus xylosus		97.62	0.42	NIT 89		
Staphylococcus arlettae		2.38	0.18	TRE 96		
264401	**Good identification**					**264401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91		
264403	**Very good identification**					**264403**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.87	0.64	MAN 91	99	
Staphylococcus simulans		0.13	-.01		1	
264410	**Very good identification**					**264410**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		95.33	0.62	MLT 95			
Staphylococcus cohnii subsp. urealyticum		4.67	0.23				
264411	**Very good identification**						**264411**
Staphylococcus cohnii subsp. urealyticum		99.79	0.64	LAC 91			
Staphylococcus kloosii		0.21	0.20				
264412	**Low discrimination**						**264412**
					NOV	YEP	
Staphylococcus kloosii		68.27	0.62	MLT 95	99	99	
Staphylococcus cohnii subsp. urealyticum		31.24	0.43		99	64	
Staphylococcus simulans		0.49	0.01		1	1	
264413	**Very good identification**						**264413**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		99.93	0.85		99	64	
Staphylococcus simulans		0.05	0.12		1	1	
Staphylococcus kloosii		0.02	0.21		99	99	
264430	**Good identification**						**264430**
Staphylococcus kloosii		100.0	0.49	MLT 95			
264432	**Good identification**						**264432**
Staphylococcus kloosii		100.0	0.50	MLT 95			
264433	**Good identification**						**264433**
Staphylococcus cohnii subsp. urealyticum		97.91	0.43	XYL 1			
Staphylococcus xylosus		2.09	0.00				
264440	**Low discrimination**						**264440**
					NOV		
Staphylococcus kloosii		82.86	0.47	MAN 99	99		
Staphylococcus cohnii subsp. urealyticum		11.86	0.18		99		
Staphylococcus warneri		5.28	0.14		1		
264441	**Very good identification**						**264441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	MAN 91			
264442	**Low discrimination**						**264442**
Staphylococcus cohnii subsp. urealyticum		57.19	0.38	MAN 91			
Staphylococcus kloosii		42.81	0.48	MNS 99			
264443	**Very good identification**						**264443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.79	MAN 91			
264450	**Very good identification**						**264450**
Staphylococcus kloosii		98.41	0.88				
Staphylococcus cohnii subsp. urealyticum		1.44	0.38				
Staphylococcus saprophyticus subsp. bovis		0.11	0.09				
264451	**Very good identification**						**264451**
Staphylococcus cohnii subsp. urealyticum		99.31	0.79	LAC 91			
Staphylococcus kloosii		0.69	0.47				
264452	**Low discrimination**						**264452**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		87.97	0.89				
Staphylococcus cohnii subsp. urealyticum		12.00	0.59				
Staphylococcus saprophyticus subsp. saprophyticus		0.03	-.11				
264453	**Excellent identification**						**264453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		99.92	1.00		99	64	
Staphylococcus kloosii		0.08	0.48		99	99	
Staphylococcus simulans		0.00	0.04		1	1	
264460	**Good identification**						**264460**
Staphylococcus kloosii		100.0	0.34	MAN 99			
264462	**Good identification**						**264462**
Staphylococcus kloosii		100.0	0.35	MAN 99			
264463	**Good identification**						**264463**
Staphylococcus cohnii subsp. urealyticum		96.87	0.38	MAN 91			
Staphylococcus xylosus		3.13	-.02	XYL 1			
264470	**Very good identification**						**264470**
Staphylococcus kloosii		100.0	0.75				
264471	**Low discrimination**						**264471**
Staphylococcus cohnii subsp. urealyticum		74.05	0.38	XYL 1			
Staphylococcus xylosus		13.16	0.14	LAC 91			
Staphylococcus kloosii		12.79	0.34				
264472	**Very good identification**						**264472**
Staphylococcus kloosii		99.45	0.76				
Staphylococcus cohnii subsp. urealyticum		0.55	0.18				
264473	**Good identification**						**264473**
Staphylococcus cohnii subsp. urealyticum		93.59	0.59	XYL 1			
Staphylococcus xylosus		4.68	0.23				
Staphylococcus kloosii		1.73	0.35				
264501	**Good identification**						**264501**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91			
264502	**Low discrimination**						**264502**
Staphylococcus kloosii		50.28	0.27	MAN 99			
Staphylococcus cohnii subsp. urealyticum		49.72	0.15	MLT 95			
264503	**Very good identification**						**264503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.56	MAN 91			
264510	**Very good identification**						**264510**
Staphylococcus kloosii		98.93	0.67	MLT 95			
Staphylococcus cohnii subsp. urealyticum		1.07	0.15				
264511	**Very good identification**						**264511**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		99.07	0.56	LAC 91			
Staphylococcus kloosii		0.93	0.26				
264512	**Good identification**						**264512**
Staphylococcus kloosii		90.83	0.68	MLT 95			
Staphylococcus cohnii subsp. urealyticum		9.17	0.36				
264513	**Very good identification**						**264513**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		99.81	0.77		1	99	
Staphylococcus kloosii		0.10	0.27		1	99	
Staphylococcus intermedius		0.05	0.01		99	1	
264530	**Very good identification**						**264530**
Staphylococcus kloosii		100.0	0.55	MLT 95			
264532	**Very good identification**						**264532**
Staphylococcus kloosii		100.0	0.56	MLT 95			
264533	**Low discrimination**						**264533**
Staphylococcus cohnii subsp. urealyticum		66.51	0.36	XYL 1			
Staphylococcus xylosus		33.49	0.21				
264540	**Good identification**						**264540**
Staphylococcus kloosii		96.94	0.53	MAN 99			
Staphylococcus cohnii subsp. urealyticum		3.06	0.10				
264541	**Good identification**						**264541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	MAN 91			
264542	**Low discrimination**						**264542**
Staphylococcus kloosii		77.22	0.53	MAN 99			
Staphylococcus cohnii subsp. urealyticum		22.78	0.31				
264543	**Very good identification**						**264543**
Staphylococcus cohnii subsp. urealyticum		99.93	0.72	MAN 91			
Staphylococcus xylosus		0.07	-.03				
264550	**Excellent identification**						**264550**
Staphylococcus kloosii		99.46	0.94				
Staphylococcus cohnii subsp. urealyticum		0.32	0.31				
Staphylococcus saprophyticus subsp. bovis		0.22	0.22				
264551	**Very good identification**						**264551**
Staphylococcus cohnii subsp. urealyticum		96.43	0.72	LAC 91			
Staphylococcus kloosii		3.05	0.53				
Staphylococcus xylosus		0.36	0.13				
264552	**Very good identification**						**264552**
Staphylococcus kloosii		97.03	0.95				
Staphylococcus cohnii subsp. urealyticum		2.92	0.51				
Staphylococcus saprophyticus subsp. bovis		0.05	0.09				
264553	**Very good identification**						**264553**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		99.55	0.92				
Staphylococcus kloosii		0.34	0.53				
Staphylococcus xylosus		0.10	0.22				
264560	**Good identification**						**264560**
Staphylococcus kloosii		100.0	0.40	MAN 99			
264562	**Good identification**						**264562**
Staphylococcus kloosii		100.0	0.41	MAN 99			
264563	**Low discrimination**						**264563**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		43.00	0.31	MAN 91	1	99	
Staphylococcus xylosus		32.76	0.19	XYL 1	1	99	
Staphylococcus lutrae		24.23	0.12		99	1	
264570	**Very good identification**						**264570**
Staphylococcus kloosii		100.0	0.81				
264571	**Low discrimination**						**264571**
Staphylococcus xylosus		70.18	0.34	NIT 89			
Staphylococcus cohnii subsp. urealyticum		16.73	0.31	SUC 95			
Staphylococcus kloosii		13.09	0.40				
264572	**Very good identification**						**264572**
Staphylococcus kloosii		99.53	0.82				
Staphylococcus arlettae		0.21	0.15				
Staphylococcus xylosus		0.14	0.03				
264573	**Low discrimination**						**264573**
					COA	NOV	
Staphylococcus xylosus		49.28	0.44	NIT 89	1	99	
Staphylococcus cohnii subsp. urealyticum		41.75	0.51	SUC 95	1	99	
Staphylococcus lutrae		4.52	0.18		99	1	
264603	**Low discrimination**						**264603**
					NOV		
Staphylococcus simulans		51.05	0.28	ARG 92	1		
Staphylococcus cohnii subsp. urealyticum		48.95	0.33	NIT 96	99		
264610	**Acceptable identification**						**264610**
					NOV		
Staphylococcus kloosii		91.78	0.41	SUC 9	99		
Staphylococcus warneri		8.22	0.12	MLT 95	1		
264611	**Good identification**						**264611**
					NOV		
Staphylococcus cohnii subsp. urealyticum		95.96	0.33	SUC 3	99		
Staphylococcus simulans		4.04	-.01	LAC 91	1		
264612	**Low discrimination**						**264612**
					NOV	YEP	
Staphylococcus simulans		60.43	0.29	ARG 92	1	1	
Staphylococcus kloosii		34.62	0.42	NIT 96	99	99	
Staphylococcus cohnii subsp. urealyticum		4.95	0.12		99	64	
264613	**Low discrimination**						**264613**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		69.90	0.53	SUC 3	99		
Staphylococcus simulans		28.84	0.40		1		
Staphylococcus xylosus		0.81	0.04		99		
264630	**Good identification**						**264630**
Staphylococcus kloosii		100.0	0.28	SUC 9			
264632	**Good identification**						**264632**
Staphylococcus kloosii		100.0	0.29	SUC 9			
264640	**Acceptable identification**						**264640**
					NOV		
Staphylococcus warneri		92.45	0.41	PHS 1	1		
Staphylococcus kloosii		7.55	0.26		99		
264641	**Good identification**						**264641**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	SUC 3			
264642	**Low discrimination**						**264642**
					NOV	YEP	
Staphylococcus warneri		60.71	0.31	PHS 1	1	43	
Staphylococcus simulans		16.45	0.10		1	1	
Staphylococcus kloosii		16.11	0.27		99	99	
264643	**Acceptable identification**						**264643**
					NOV		
Staphylococcus cohnii subsp. urealyticum		90.90	0.48	SUC 3	99		
Staphylococcus simulans		7.52	0.20	MAN 91	1		
Staphylococcus xylosus		1.58	0.03		99		
264650	**Low discrimination**						**264650**
					NOV	PYR	
Staphylococcus kloosii		87.39	0.67	SUC 9	99	52	
Staphylococcus warneri		7.83	0.38		1	8	
Staphylococcus saprophyticus subsp. bovis		3.79	0.22		99	99	
264651	**Low discrimination**						**264651**
Staphylococcus cohnii subsp. urealyticum		87.55	0.48	SUC 3			
Staphylococcus xylosus		8.40	0.18	LAC 91			
Staphylococcus saprophyticus subsp. bovis		2.10	0.09				
264652	**Low discrimination**						**264652**
					NOV	YEP	
Staphylococcus kloosii		86.48	0.68	SUC 9	99	99	
Staphylococcus cohnii subsp. urealyticum		3.69	0.28		99	64	
Staphylococcus simulans		3.57	0.22		1	1	
264653	**Good identification**						**264653**
					NOV		
Staphylococcus cohnii subsp. urealyticum		94.00	0.69	SUC 3	99		
Staphylococcus simulans		3.08	0.33		1		
Staphylococcus xylosus		2.54	0.28		99		
264670	**Very good identification**						**264670**
Staphylococcus kloosii		98.02	0.55	SUC 9			
Staphylococcus xylosus		1.98	-.01				
264671	**Good identification**						**264671**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus xylosus		97.21	0.40	NIT 89				
Staphylococcus equorum		1.90	0.06	GAL 91				
Staphylococcus cohnii subsp. urealyticum		0.89	0.07					
264672	**Low discrimination**							**264672**
Staphylococcus kloosii		56.68	0.56	SUC 9				
Staphylococcus arlettae		40.32	0.41					
Staphylococcus xylosus		3.00	0.08					
264673	**Low discrimination**							**264673**
Staphylococcus xylosus		81.31	0.49	NIT 89				
Staphylococcus arlettae		14.61	0.43	GAL 91				
Staphylococcus cohnii subsp. urealyticum		2.65	0.28					
264703	**Low discrimination**							**264703**
					NOV	PYR	COA	
Staphylococcus intermedius		71.67	0.33	GLR 3	1	90	99	
Staphylococcus cohnii subsp. urealyticum		12.76	0.25	NIT 93	99	53	1	
Staphylococcus hyicus		10.64	0.11		1	1	24	
264710	**Good identification**							**264710**
Staphylococcus kloosii		100.0	0.47	SUC 9				
264712	**Low discrimination**							**264712**
					COA	NOV	YEP	
Staphylococcus kloosii		86.41	0.48	SUC 9	1	99	99	
Staphylococcus simulans		6.76	0.07	MLT 95	1	1	1	
Staphylococcus intermedius		4.10	0.01		99	1	1	
264713	**Low discrimination**							**264713**
					COA	NOV		
Staphylococcus intermedius		52.22	0.42	GLR 3	99	1		
Staphylococcus cohnii subsp. urealyticum		34.76	0.46	NIT 93	1	99		
Staphylococcus xylosus		9.44	0.25		1	99		
264730	**Low discrimination**							**264730**
Staphylococcus kloosii		86.94	0.34	SUC 9				
Staphylococcus xylosus		13.06	-.04	MLT 95				
264731	**Good identification**							**264731**
Staphylococcus xylosus		97.50	0.38	NIT 89				
Staphylococcus equorum		2.50	0.06	MLT 93				
264732	**Low discrimination**							**264732**
Staphylococcus kloosii		55.76	0.35	SUC 9				
Staphylococcus arlettae		22.22	0.15	MLT 95				
Staphylococcus xylosus		22.01	0.06					
264733	**Good identification**							**264733**
					YEP			
Staphylococcus xylosus		95.71	0.47	NIT 89	40			
Staphylococcus equorum		2.22	0.15	MLT 93	1			
Staphylococcus arlettae		1.29	0.18		99			
264740	**Low discrimination**							**264740**
					NOV			
Staphylococcus warneri		66.59	0.30	PHS 1	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		33.41	0.32		99		
264742	**Low discrimination**						**264742**
					NOV		
Staphylococcus kloosii		61.99	0.33	SUC 9	99		
Staphylococcus warneri		38.01	0.21	MAN 99	1		
264743	**Low discrimination**						**264743**
					COA	NOV	
Staphylococcus intermedius		51.31	0.37	GLR 3	99	1	
Staphylococcus cohnii subsp. urealyticum		34.50	0.41	NIT 93	1	99	
Staphylococcus xylosus		14.18	0.24		1	99	
264750	**Good identification**						**264750**
					NOV	PYR	
Staphylococcus kloosii		90.39	0.73	SUC 9	99	52	
Staphylococcus saprophyticus subsp. bovis		8.09	0.34		99	99	
Staphylococcus warneri		1.32	0.28		1	8	
264751	**Low discrimination**						**264751**
					COA	NOV	
Staphylococcus xylosus		53.34	0.39	NIT 89	1	99	
Staphylococcus cohnii subsp. urealyticum		23.57	0.41	XYL 92	1	99	
Staphylococcus intermedius		15.43	0.30		99	1	
264752	**Very good identification**						**264752**
Staphylococcus kloosii		95.11	0.74	SUC 9			
Staphylococcus saprophyticus subsp. bovis		1.96	0.22				
Staphylococcus cohnii subsp. urealyticum		0.90	0.20				
264753	**Low discrimination**						**264753**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		48.42	0.61	SUC 3	1	99	
Staphylococcus xylosus		30.85	0.48		1	99	
Staphylococcus intermedius		19.26	0.46		99	1	
264761	**Good identification**						**264761**
Staphylococcus xylosus		100.0	0.36	NIT 89			
264763	**Acceptable identification**						**264763**
					COA	NOV	
Staphylococcus xylosus		94.44	0.45	NIT 89	1	99	
Staphylococcus lutrae		3.68	0.12	MAN 94	99	1	
Staphylococcus arlettae		1.51	0.18		1	99	
264770	**Low discrimination**						**264770**
Staphylococcus kloosii		88.67	0.61	SUC 9			
Staphylococcus xylosus		9.32	0.20				
Staphylococcus arlettae		2.02	0.15				
264771	**Very good identification**						**264771**
Staphylococcus xylosus		97.43	0.61	NIT 89			
Staphylococcus equorum		1.90	0.27				
Staphylococcus gallinarum		0.29	0.07				
264772	**Low discrimination**						**264772**
Staphylococcus arlettae		61.94	0.56	URE 1			
Staphylococcus kloosii		29.83	0.61				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		8.23	0.29			
264773	**Low discrimination**					**264773**
					YEP	
Staphylococcus xylosus		88.65	0.70	NIT 89	40	
Staphylococcus arlettae		8.93	0.59		99	
Staphylococcus equorum		1.56	0.35		1	
265013	**Good identification**					**265013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
265043	**Good identification**					**265043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	TRE 99		
265050	**Good identification**					**265050**
Staphylococcus kloosii		100.0	0.47	TRE 99		
265051	**Good identification**					**265051**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	TRE 99		
265052	**Acceptable identification**					**265052**
Staphylococcus kloosii		94.96	0.48	TRE 99		
Staphylococcus cohnii subsp. urealyticum		5.04	0.09			
265053	**Good identification**					**265053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	TRE 99		
265070	**Good identification**					**265070**
Staphylococcus kloosii		100.0	0.34	TRE 99		
265072	**Good identification**					**265072**
Staphylococcus kloosii		100.0	0.35	TRE 99		
265112	**Good identification**					**265112**
Staphylococcus kloosii		100.0	0.27	TRE 99		
265113	**Good identification**					**265113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99		
265150	**Good identification**					**265150**
Staphylococcus kloosii		100.0	0.53	TRE 99		
265152	**Very good identification**					**265152**
Staphylococcus kloosii		100.0	0.53	TRE 99		
265153	**Good identification**					**265153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	TRE 99		
265170	**Good identification**					**265170**
Staphylococcus kloosii		100.0	0.40	TRE 99		
265172	**Good identification**					**265172**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.41	TRE 99		
265252	**Good identification**					**265252**
Staphylococcus kloosii		100.0	0.27	SUC 9		
265350	**Good identification**					**265350**
Staphylococcus kloosii		100.0	0.32	SUC 9		
265352	**Good identification**					**265352**
Staphylococcus kloosii		100.0	0.33	SUC 9		
265371	**Low discrimination**					**265371**
Staphylococcus equorum		59.27	0.27	NIT 92		
Staphylococcus xylosus		31.63	0.20	TRE 99		
Staphylococcus gallinarum		9.11	0.07			
265373	**Low discrimination**					**265373**
Staphylococcus equorum		55.72	0.35	NIT 92		
Staphylococcus xylosus		32.91	0.29	TRE 99		
Staphylococcus gallinarum		8.15	0.15			
265401	**Good identification**					**265401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91		
265403	**Very good identification**					**265403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	MAN 91		
265410	**Very good identification**					**265410**
Staphylococcus kloosii		98.13	0.62	MLT 95		
Staphylococcus cohnii subsp. urealyticum		1.87	0.14			
265411	**Very good identification**					**265411**
Staphylococcus cohnii subsp. urealyticum		99.47	0.55	LAC 91		
Staphylococcus kloosii		0.53	0.20			
265412	**Low discrimination**					**265412**
Staphylococcus kloosii		84.90	0.62	MLT 95		
Staphylococcus cohnii subsp. urealyticum		15.11	0.35			
265413	**Very good identification**					**265413**
Staphylococcus cohnii subsp. urealyticum		99.94	0.76			
Staphylococcus kloosii		0.06	0.21			
265430	**Good identification**					**265430**
Staphylococcus kloosii		100.0	0.49	MLT 95		
265432	**Good identification**					**265432**
Staphylococcus kloosii		100.0	0.50	MLT 95		
265433	**Good identification**					**265433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	XYL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
265440	**Acceptable identification**					**265440**
	Staphylococcus kloosii	94.73	0.47	MAN 99		
	Staphylococcus cohnii subsp. urealyticum	5.27	0.09			
265441	**Good identification**					**265441**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.50	MAN 91		
265442	**Low discrimination**					**265442**
	Staphylococcus kloosii	65.81	0.48	MAN 99		
	Staphylococcus cohnii subsp. urealyticum	34.19	0.30			
265443	**Very good identification**					**265443**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.71	MAN 91		
265450	**Very good identification**					**265450**
	Staphylococcus kloosii	99.44	0.88			
	Staphylococcus cohnii subsp. urealyticum	0.57	0.30			
265451	**Very good identification**					**265451**
	Staphylococcus cohnii subsp. urealyticum	98.24	0.71	LAC 91		
	Staphylococcus kloosii	1.76	0.47			
265452	**Good identification**					**265452**
	Staphylococcus kloosii	94.96	0.89			
	Staphylococcus cohnii subsp. urealyticum	5.04	0.51			
265453	**Very good identification**					**265453**
	Staphylococcus cohnii subsp. urealyticum	99.81	0.92			
	Staphylococcus kloosii	0.19	0.48			
265460	**Good identification**					**265460**
	Staphylococcus kloosii	100.0	0.34	MAN 99		
265462	**Good identification**					**265462**
	Staphylococcus kloosii	100.0	0.35	MAN 99		
265463	**Good identification**					**265463**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.30	MAN 91		
265470	**Very good identification**					**265470**
	Staphylococcus kloosii	100.0	0.75			
265471	**Low discrimination**					**265471**
					YEP	
	Staphylococcus cohnii subsp. urealyticum	57.87	0.30	XYL 1	64	
	Staphylococcus kloosii	25.70	0.34	LAC 91	99	
	Staphylococcus equorum	9.81	0.06		1	
265472	**Very good identification**					**265472**
	Staphylococcus kloosii	99.79	0.76			
	Staphylococcus cohnii subsp. urealyticum	0.21	0.09			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
265473	**Low discrimination**					**265473**
					YEP	
Staphylococcus cohnii subsp. urealyticum		89.06	0.51	XYL 1	64	
Staphylococcus kloosii		4.24	0.35		99	
Staphylococcus equorum		3.84	0.15		1	
265501	**Good identification**					**265501**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91		
265502	**Low discrimination**					**265502**
Staphylococcus kloosii		72.22	0.27	MAN 99		
Staphylococcus cohnii subsp. urealyticum		27.78	0.07	MLT 95		
265503	**Good identification**					**265503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	MAN 91		
265510	**Very good identification**					**265510**
Staphylococcus kloosii		99.58	0.67	MLT 95		
Staphylococcus cohnii subsp. urealyticum		0.42	0.07			
265511	**Good identification**					**265511**
Staphylococcus cohnii subsp. urealyticum		97.63	0.48	LAC 91		
Staphylococcus kloosii		2.37	0.26			
265512	**Very good identification**					**265512**
Staphylococcus kloosii		96.22	0.68	MLT 95		
Staphylococcus cohnii subsp. urealyticum		3.78	0.27			
265513	**Very good identification**					**265513**
Staphylococcus cohnii subsp. urealyticum		99.74	0.69			
Staphylococcus kloosii		0.26	0.27			
265530	**Very good identification**					**265530**
Staphylococcus kloosii		100.0	0.55	MLT 95		
265532	**Very good identification**					**265532**
Staphylococcus kloosii		100.0	0.56	MLT 95		
265533	**Low discrimination**					**265533**
Staphylococcus cohnii subsp. urealyticum		52.81	0.27	XYL 1		
Staphylococcus equorum		30.10	0.15			
Staphylococcus xylosus		17.09	0.08			
265540	**Good identification**					**265540**
Staphylococcus kloosii		100.0	0.53	MAN 99		
265541	**Good identification**					**265541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91		
265542	**Low discrimination**					**265542**
Staphylococcus kloosii		89.71	0.53	MAN 99		
Staphylococcus cohnii subsp. urealyticum		10.29	0.22			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
265543	**Very good identification**					**265543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.63	MAN 91		
265550	**Excellent identification**					**265550**
Staphylococcus kloosii		99.88	0.94			
Staphylococcus cohnii subsp. urealyticum		0.13	0.22			
265551	**Good identification**					**265551**
Staphylococcus cohnii subsp. urealyticum		92.28	0.63	LAC 91		
Staphylococcus kloosii		7.50	0.53			
Staphylococcus xylosus		0.22	0.00			
265552	**Very good identification**					**265552**
Staphylococcus kloosii		98.84	0.95			
Staphylococcus cohnii subsp. urealyticum		1.16	0.43			
265553	**Very good identification**					**265553**
Staphylococcus cohnii subsp. urealyticum		99.07	0.84			
Staphylococcus kloosii		0.86	0.53			
Staphylococcus xylosus		0.07	0.09			
265560	**Good identification**					**265560**
Staphylococcus kloosii		100.0	0.40	MAN 99		
265562	**Good identification**					**265562**
Staphylococcus kloosii		100.0	0.41	MAN 99		
265570	**Very good identification**					**265570**
Staphylococcus kloosii		100.0	0.81			
265571	**Low discrimination**					**265571**
					YEP	
Staphylococcus equorum		38.75	0.27	NIT 92	1	
Staphylococcus xylosus		26.12	0.22	SUC 99	40	
Staphylococcus kloosii		19.49	0.40		99	
265572	**Very good identification**					**265572**
Staphylococcus kloosii		99.79	0.82			
Staphylococcus arlettae		0.21	0.15			
265573	**Low discrimination**					**265573**
Staphylococcus equorum		31.80	0.35	NIT 92		
Staphylococcus cohnii subsp. urealyticum		31.26	0.43	SUC 99		
Staphylococcus xylosus		23.72	0.31			
265610	**Good identification**					**265610**
Staphylococcus kloosii		100.0	0.41	SUC 9		
265612	**Good identification**					**265612**
Staphylococcus kloosii		100.0	0.42	SUC 9		
265613	**Good identification**					**265613**
					NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		98.94	0.45	SUC 3	99	
Staphylococcus simulans		1.06	-.01		1	
265630	**Good identification**					**265630**
Staphylococcus kloosii		100.0	0.28	SUC 9		
265631	**Acceptable identification**					**265631**
Staphylococcus equorum		91.04	0.27	NIT 92		
Staphylococcus xylosus		8.96	0.04	GAL 91		
265632	**Good identification**					**265632**
Staphylococcus kloosii		100.0	0.29	SUC 9		
265633	**Acceptable identification**					**265633**
Staphylococcus equorum		90.17	0.35	NIT 92		
Staphylococcus xylosus		9.83	0.14	GAL 91		
265642	**Good identification**					**265642**
Staphylococcus kloosii		100.0	0.27	SUC 9		
265643	**Good identification**					**265643**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	SUC 3		
265650	**Very good identification**					**265650**
Staphylococcus kloosii		100.0	0.67	SUC 9		
265651	**Low discrimination**					**265651**
Staphylococcus cohnii subsp. urealyticum		85.15	0.40	SUC 3		
Staphylococcus xylosus		5.25	0.06	LAC 91		
Staphylococcus kloosii		4.89	0.26			
265652	**Very good identification**					**265652**
Staphylococcus kloosii		98.37	0.68	SUC 9		
Staphylococcus cohnii subsp. urealyticum		1.63	0.19			
265653	**Very good identification**					**265653**
Staphylococcus cohnii subsp. urealyticum		96.38	0.61	SUC 3		
Staphylococcus xylosus		1.67	0.15			
Staphylococcus equorum		1.36	0.15			
265670	**Very good identification**					**265670**
					YEP	
Staphylococcus kloosii		96.25	0.55	SUC 9	99	
Staphylococcus equorum		3.75	0.06		1	
265671	**Low discrimination**					**265671**
Staphylococcus equorum		87.33	0.47	NIT 92		
Staphylococcus xylosus		11.30	0.28	GAL 91		
Staphylococcus gallinarum		1.37	0.07			
265672	**Low discrimination**					**265672**
					YEP	
Staphylococcus kloosii		55.03	0.56	SUC 9	99	
Staphylococcus arlettae		39.15	0.41		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus equorum		5.09	0.15		1	
265673	**Low discrimination**					**265673**
					YEP	
Staphylococcus equorum		78.85	0.56	NIT 92	1	
Staphylococcus xylosus		11.29	0.37	GAL 91	40	
Staphylococcus arlettae		8.12	0.43		99	
265710	**Good identification**					**265710**
Staphylococcus kloosii		100.0	0.47	SUC 9		
265712	**Good identification**					**265712**
Staphylococcus kloosii		100.0	0.48	SUC 9		
265713	**Low discrimination**					**265713**
Staphylococcus cohnii subsp. urealyticum		71.44	0.37	SUC 3		
Staphylococcus equorum		13.30	0.15			
Staphylococcus xylosus		12.48	0.13			
265730	**Low discrimination**					**265730**
					YEP	
Staphylococcus kloosii		72.38	0.34	SUC 9	99	
Staphylococcus equorum		27.62	0.06	MLT 95	1	
265731	**Low discrimination**					**265731**
Staphylococcus equorum		89.76	0.47	NIT 92		
Staphylococcus xylosus		8.84	0.25	MLT 91		
Staphylococcus gallinarum		1.41	0.07			
265732	**Low discrimination**					**265732**
					YEP	
Staphylococcus kloosii		43.40	0.35	SUC 9	99	
Staphylococcus equorum		39.31	0.15	MLT 95	1	
Staphylococcus arlettae		17.30	0.15		99	
265733	**Low discrimination**					**265733**
Staphylococcus equorum		88.51	0.56	NIT 92		
Staphylococcus xylosus		9.65	0.35	MLT 91		
Staphylococcus gallinarum		1.32	0.15			
265740	**Good identification**					**265740**
Staphylococcus kloosii		100.0	0.32	SUC 9		
265742	**Good identification**					**265742**
Staphylococcus kloosii		100.0	0.33	SUC 9		
265743	**Low discrimination**					**265743**
Staphylococcus cohnii subsp. urealyticum		76.17	0.32	SUC 3		
Staphylococcus xylosus		20.13	0.11	MAN 91		
Staphylococcus gallinarum		3.70	-.05			
265750	**Very good identification**					**265750**
Staphylococcus kloosii		100.0	0.73	SUC 9		
265752	**Very good identification**					**265752**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		99.12	0.74	SUC 9			
Staphylococcus arlettae		0.52	0.15				
Staphylococcus cohnii subsp. urealyticum		0.36	0.12				
265753	**Low discrimination**						**265753**
Staphylococcus cohnii subsp. urealyticum		54.12	0.53	SUC 3			
Staphylococcus xylosus		22.16	0.36				
Staphylococcus equorum		17.98	0.35				
265761	**Low discrimination**						**265761**
Staphylococcus gallinarum		45.28	0.28	NIT 99			
Staphylococcus equorum		30.11	0.27	MAN 91			
Staphylococcus xylosus		24.61	0.24				
265763	**Low discrimination**						**265763**
Staphylococcus gallinarum		42.19	0.36	NIT 99			
Staphylococcus equorum		29.46	0.35	MAN 91			
Staphylococcus xylosus		26.65	0.33				
265770	**Low discrimination**						**265770**
					YEP		
Staphylococcus kloosii		77.93	0.61	SUC 9	99		
Staphylococcus equorum		15.82	0.27		1		
Staphylococcus gallinarum		2.43	0.07		78		
265771	**Low discrimination**						**265771**
Staphylococcus equorum		77.82	0.68	NIT 92			
Staphylococcus gallinarum		11.96	0.48				
Staphylococcus xylosus		10.07	0.48				
265772	**Low discrimination**						**265772**
					YEP		
Staphylococcus arlettae		56.15	0.56	URE 1	99		
Staphylococcus kloosii		27.04	0.61		99		
Staphylococcus equorum		13.03	0.35		1		
265773	**Low discrimination**						**265773**
Staphylococcus equorum		74.20	0.76	NIT 92			
Staphylococcus gallinarum		10.85	0.56				
Staphylococcus xylosus		10.63	0.58				
266013	**Low discrimination**						**266013**
					NOV		
Staphylococcus cohnii subsp. urealyticum		72.64	0.28	TRE 99	99		
Staphylococcus simulans		27.37	0.14		1		
266053	**Good identification**						**266053**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.10	0.43	TRE 99	99		
Staphylococcus simulans		2.90	0.07		1		
266153	**Low discrimination**						**266153**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		79.00	0.36	TRE 99	1	99	
Staphylococcus xylosus		15.49	0.12		1	99	
Staphylococcus intermedius		5.51	0.05		99	1	
266173	**Low discrimination**						**266173**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	NOV		
Staphylococcus xylosus		78.63	0.34	SUC 95	1	99		
Staphylococcus lutrae		21.37	0.18	TRE 96	99	1		
266203	**Low discrimination**							**266203**
Staphylococcus simulans		83.12	0.30	ARG 92				
Staphylococcus hyicus		16.88	0.08	TRE 95				
266212	**Good identification**							**266212**
Staphylococcus simulans		100.0	0.31	ARG 92				
266213	**Good identification**							**266213**
					COA			
Staphylococcus simulans		98.77	0.42	ARG 92	1			
Staphylococcus intermedius		1.23	0.01	TRE 95	99			
266253	**Low discrimination**							**266253**
					NOV			
Staphylococcus simulans		76.62	0.35	ARG 92	1			
Staphylococcus xylosus		16.88	0.18	TRE 95	99			
Staphylococcus cohnii subsp. urealyticum		3.31	0.12		99			
266271	**Good identification**							**266271**
Staphylococcus xylosus		100.0	0.30	GAL 91				
266273	**Good identification**							**266273**
Staphylococcus xylosus		100.0	0.40	GAL 91				
266303	**Low discrimination**							**266303**
					YEP	PYR	COA	
Staphylococcus hyicus		52.98	0.30	ARG 96	1	1	24	
Staphylococcus intermedius		41.47	0.33	BGA 10	1	90	99	
Staphylococcus chromogenes		3.57	0.10	TRE 92	95	45	1	
266313	**Low discrimination**							**266313**
					COA	NOV		
Staphylococcus intermedius		86.73	0.42	GLR 3	99	1		
Staphylococcus simulans		6.10	0.20	TRE 93	1	1		
Staphylococcus xylosus		5.29	0.15		1	99		
266331	**Good identification**							**266331**
Staphylococcus xylosus		100.0	0.28	TRE 96				
266333	**Good identification**							**266333**
					COA	NOV		
Staphylococcus xylosus		98.58	0.37	TRE 96	1	99		
Staphylococcus intermedius		1.42	0.01	MLT 93	99	1		
266343	**Low discrimination**							**266343**
					COA	NOV	PYR	
Staphylococcus intermedius		80.29	0.37	GLR 3	99	1	90	
Staphylococcus xylosus		7.48	0.14	TRE 93	1	99	65	
Staphylococcus epidermidis		6.47	0.17		1	1	1	
266350	**Good identification**							**266350**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	TRE 99				
266351	**Low discrimination**							**266351**
					COA	NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		46.48	0.29	TRE 96	1	99	
Staphylococcus intermedius		39.88	0.30	XYL 92	99	1	
Staphylococcus saprophyticus subsp. bovis		13.64	0.22		1	99	
266353	**Low discrimination**						**266353**
					COA	NOV	
Staphylococcus intermedius		63.07	0.46	GLR 3	99	1	
Staphylococcus xylosus		34.06	0.38	TRE 93	1	99	
Staphylococcus simulans		1.33	0.13		1	1	
266363	**Low discrimination**						**266363**
					COA	NOV	
Staphylococcus xylosus		88.71	0.36	TRE 96	1	99	
Staphylococcus lutrae		10.24	0.12	MAN 94	99	1	
Staphylococcus gallinarum		1.05	-.05		1	99	
266371	**Good identification**						**266371**
Staphylococcus xylosus		98.46	0.51	TRE 96			
Staphylococcus gallinarum		0.88	0.07				
Staphylococcus equorum		0.66	0.08				
266373	**Very good identification**						**266373**
					COA	NOV	
Staphylococcus xylosus		97.12	0.60	TRE 96	1	99	
Staphylococcus lutrae		1.39	0.18		99	1	
Staphylococcus gallinarum		0.74	0.15		1	99	
266401	**Good identification**						**266401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	MAN 91			
266403	**Low discrimination**						**266403**
					NOV		
Staphylococcus cohnii subsp. urealyticum		82.35	0.48	MAN 91	99		
Staphylococcus simulans		15.05	0.28		1		
Staphylococcus felis		2.60	0.03		1		
266411	**Good identification**						**266411**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.27	0.48	LAC 91	99		
Staphylococcus simulans		0.73	-.01		1		
266412	**Low discrimination**						**266412**
					NOV	YEP	
Staphylococcus simulans		65.52	0.29	ARG 92	1	1	
Staphylococcus cohnii subsp. urealyticum		30.64	0.28	SUC 96	99	64	
Staphylococcus kloosii		3.83	0.21		99	99	
266413	**Good identification**						**266413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		91.40	0.69		99		
Staphylococcus simulans		6.61	0.40		1		
Staphylococcus felis		1.91	0.20		1		
266433	**Low discrimination**						**266433**
					NOV		
Staphylococcus cohnii subsp. urealyticum		48.76	0.28	XYL 1	99		
Staphylococcus xylosus		47.71	0.19		99		
Staphylococcus simulans		3.53	-.01		1		
266441	**Good identification**						**266441**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91			
266443	**Very good identification**						**266443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.45	0.64	MAN 91	99		
Staphylococcus simulans		1.43	0.20		1		
Staphylococcus xylosus		0.13	-.05		99		
266450	**Low discrimination**						**266450**
Staphylococcus saprophyticus subsp. bovis		89.34	0.50				
Staphylococcus kloosii		8.49	0.47				
Staphylococcus cohnii subsp. urealyticum		2.17	0.23				
266451	**Low discrimination**						**266451**
Staphylococcus cohnii subsp. urealyticum		90.00	0.64	LAC 91			
Staphylococcus saprophyticus subsp. bovis		9.36	0.38				
Staphylococcus xylosus		0.64	0.11				
266452	**Low discrimination**						**266452**
Staphylococcus saprophyticus subsp. bovis		39.06	0.38				
Staphylococcus cohnii subsp. urealyticum		38.35	0.43				
Staphylococcus kloosii		16.09	0.48				
266453	**Very good identification**						**266453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.98	0.85		99		
Staphylococcus simulans		0.57	0.33		1		
Staphylococcus saprophyticus subsp. bovis		0.26	0.26		99		
266470	**Low discrimination**						**266470**
Staphylococcus kloosii		70.17	0.34	NIT 1			
Staphylococcus saprophyticus subsp. bovis		29.83	0.09				
266471	**Low discrimination**						**266471**
Staphylococcus xylosus		89.07	0.32	GAL 91			
Staphylococcus cohnii subsp. urealyticum		10.93	0.23	SUC 95			
266472	**Low discrimination**						**266472**
Staphylococcus kloosii		81.90	0.35	NIT 1			
Staphylococcus xylosus		18.10	0.01				
266473	**Low discrimination**						**266473**
					COA	NOV	
Staphylococcus xylosus		64.50	0.42	GAL 91	1	99	
Staphylococcus cohnii subsp. urealyticum		28.12	0.43	SUC 95	1	99	
Staphylococcus lutrae		7.39	0.18		99	1	
266503	**Low discrimination**						**266503**
					NOV		
Staphylococcus cohnii subsp. urealyticum		64.03	0.41	MAN 91	99		
Staphylococcus felis		15.78	0.14		1		
Staphylococcus hyicus		9.36	0.11		1		
266511	**Low discrimination**						**266511**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		84.33	0.41	LAC 91	1	99	
Staphylococcus xylosus		6.08	0.08		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		4.91	0.08		99	1		
266513	**Low discrimination**							**266513**
					NOV			
Staphylococcus cohnii subsp. urealyticum		78.40	0.61		99			
Staphylococcus felis		12.79	0.31		1			
Staphylococcus condimenti		3.30	0.14		1			
266531	**Good identification**							**266531**
Staphylococcus xylosus		100.0	0.30	SUC 95				
266533	**Low discrimination**							**266533**
					COA	NOV		
Staphylococcus xylosus		83.77	0.39	SUC 95	1	99		
Staphylococcus lutrae		12.61	0.18	MLT 93	99	1		
Staphylococcus cohnii subsp. urealyticum		3.63	0.20		1	99		
266541	**Low discrimination**							**266541**
Staphylococcus cohnii subsp. urealyticum		79.01	0.36	MAN 91				
Staphylococcus xylosus		8.61	0.07	LAC 91				
Staphylococcus saprophyticus subsp. bovis		7.83	0.09					
266543	**Good identification**							**266543**
					COA	PYR	NOV	
Staphylococcus cohnii subsp. urealyticum		91.08	0.56	MAN 91	1	53	99	
Staphylococcus intermedius		3.19	0.19		99	90	1	
Staphylococcus lutrae		2.94	0.12		99	10	1	
266550	**Very good identification**							**266550**
Staphylococcus saprophyticus subsp. bovis		95.35	0.63					
Staphylococcus kloosii		4.40	0.53					
Staphylococcus cohnii subsp. urealyticum		0.25	0.15					
266551	**Low discrimination**							**266551**
Staphylococcus cohnii subsp. urealyticum		46.40	0.56	LAC 91				
Staphylococcus saprophyticus subsp. bovis		45.05	0.50					
Staphylococcus xylosus		7.84	0.31					
266552	**Low discrimination**							**266552**
Staphylococcus saprophyticus subsp. bovis		76.12	0.50					
Staphylococcus kloosii		15.22	0.53					
Staphylococcus cohnii subsp. urealyticum		8.01	0.36					
266553	**Good identification**							**266553**
Staphylococcus cohnii subsp. urealyticum		91.89	0.77					
Staphylococcus xylosus		4.37	0.41					
Staphylococcus saprophyticus subsp. bovis		2.21	0.38					
266561	**Low discrimination**							**266561**
					COA	NOV		
Staphylococcus xylosus		79.35	0.28	SUC 95	1	99		
Staphylococcus lutrae		20.65	0.12	MAN 94	99	1		
266563	**Low discrimination**							**266563**
					COA	NOV		
Staphylococcus lutrae		89.71	0.53	GLR 1	99	1		
Staphylococcus xylosus		9.91	0.38		1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		0.28	0.15		1	99	
266570	**Low discrimination**						**266570**
Staphylococcus kloosii		43.22	0.40	NIT 1			
Staphylococcus saprophyticus subsp. bovis		37.85	0.22				
Staphylococcus xylosus		18.93	0.12				
266571	**Good identification**						**266571**
					COA	NOV	
Staphylococcus xylosus		94.80	0.53	SUC 95	1	99	
Staphylococcus lutrae		3.06	0.18		99	1	
Staphylococcus gallinarum		0.67	0.07		1	99	
266573	**Low discrimination**						**266573**
					COA	NOV	
Staphylococcus lutrae		52.14	0.59	GLR 1	99	1	
Staphylococcus xylosus		46.50	0.62		1	99	
Staphylococcus cohnii subsp. urealyticum		0.86	0.36		1	99	
266602	**Good identification**						**266602**
Staphylococcus simulans		100.0	0.45	ARG 92			
266603	**Low discrimination**						**266603**
Staphylococcus simulans		85.44	0.56	ARG 92			
Staphylococcus hyicus		10.50	0.30				
Staphylococcus felis		1.25	0.09				
266611	**Low discrimination**						**266611**
					NOV		
Staphylococcus simulans		56.28	0.27	ARG 92	1		
Staphylococcus xylosus		18.42	0.14	LAC 99	99		
Staphylococcus cohnii subsp. urealyticum		9.82	0.17		99		
266612	**Very good identification**						**266612**
					NOV		
Staphylococcus simulans		99.55	0.58	ARG 92	1		
Staphylococcus saprophyticus subsp. bovis		0.45	0.09		99		
266613	**Good identification**						**266613**
					NOV		
Staphylococcus simulans		94.24	0.68	ARG 92	1		
Staphylococcus felis		2.30	0.26		1		
Staphylococcus cohnii subsp. urealyticum		1.68	0.38		99		
266631	**Good identification**						**266631**
Staphylococcus xylosus		96.48	0.36	GAL 91			
Staphylococcus equorum		3.52	0.07	MLT 93			
266633	**Low discrimination**						**266633**
					NOV		
Staphylococcus xylosus		88.79	0.45	GAL 91	99		
Staphylococcus simulans		8.29	0.27	MLT 93	1		
Staphylococcus equorum		2.92	0.16		99		
266640	**Low discrimination**						**266640**
					NOV	PYR	
Staphylococcus warneri		54.99	0.33	PHS 1	1	8	
Staphylococcus saprophyticus subsp. bovis		45.01	0.22		99	99	
266642	**Acceptable identification**						**266642**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Staphylococcus simulans		90.39	0.38	ARG 92	1	70	
Staphylococcus warneri		5.96	0.23		1	8	
Staphylococcus saprophyticus subsp. bovis		3.66	0.09		99	99	
266643	**Low discrimination**						**266643**
					NOV		
Staphylococcus simulans		81.52	0.49	ARG 92	1		
Staphylococcus cohnii subsp. urealyticum		7.25	0.33		99		
Staphylococcus xylosus		5.79	0.22		99		
266650	**Very good identification**						**266650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.56	0.63		99	99	
Staphylococcus warneri		0.88	0.30		1	8	
Staphylococcus simulans		0.25	0.09		1	70	
266651	**Low discrimination**						**266651**
Staphylococcus saprophyticus subsp. bovis		68.93	0.50				
Staphylococcus xylosus		22.53	0.37				
Staphylococcus cohnii subsp. urealyticum		5.12	0.33				
266652	**Low discrimination**						**266652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		49.07	0.50		99	99	
Staphylococcus simulans		49.01	0.50		1	70	
Staphylococcus warneri		0.59	0.20		1	8	
266653	**Low discrimination**						**266653**
					NOV		
Staphylococcus simulans		62.17	0.61	ARG 92	1		
Staphylococcus xylosus		17.30	0.46		99		
Staphylococcus cohnii subsp. urealyticum		13.97	0.53		99		
266661	**Good identification**						**266661**
Staphylococcus xylosus		100.0	0.34	GAL 91			
266663	**Acceptable identification**						**266663**
					COA	NOV	
Staphylococcus xylosus		93.81	0.43	GAL 91	1	99	
Staphylococcus lutrae		4.56	0.12	MAN 94	99	1	
Staphylococcus simulans		1.16	0.08		1	1	
266671	**Very good identification**						**266671**
Staphylococcus xylosus		96.70	0.59	GAL 91			
Staphylococcus equorum		2.68	0.28				
Staphylococcus gallinarum		0.36	0.07				
266672	**Low discrimination**						**266672**
					NOV		
Staphylococcus xylosus		84.26	0.27	GAL 91	99		
Staphylococcus saprophyticus subsp. bovis		7.88	0.09	MNS 99	99		
Staphylococcus simulans		7.87	0.09		1		
266673	**Very good identification**						**266673**
					COA	NOV	
Staphylococcus xylosus		96.32	0.68	GAL 91	1	99	
Staphylococcus equorum		2.42	0.36		1	99	
Staphylococcus lutrae		0.58	0.18		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
266701	**Low discrimination**							**266701**
					YEP	PYR	COA	
Staphylococcus intermedius		59.43	0.40	GLR 3	1	90	99	
Staphylococcus hyicus		35.10	0.30	LAC 86	1	1	24	
Staphylococcus chromogenes		3.03	0.12		95	45	1	
266703	**Low discrimination**							**266703**
					YEP	PYR	COA	
Staphylococcus hyicus		47.91	0.52	ARG 96	1	1	24	
Staphylococcus intermedius		43.33	0.56	BGA 10	1	90	99	
Staphylococcus chromogenes		3.23	0.32		95	45	1	
266710	**Acceptable identification**							**266710**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		93.86	0.34	MLT 99	1	99		
Staphylococcus intermedius		6.14	0.08		99	1		
266711	**Low discrimination**							**266711**
					COA	NOV		
Staphylococcus intermedius		76.98	0.49	GLR 3	99	1		
Staphylococcus xylosus		18.30	0.34	LAC 86	1	99		
Staphylococcus saprophyticus subsp. bovis		2.97	0.22		1	99		
266712	**Low discrimination**							**266712**
					COA	NOV		
Staphylococcus simulans		61.97	0.36	ARG 92	1	1		
Staphylococcus intermedius		20.82	0.24	GAL 8	99	1		
Staphylococcus saprophyticus subsp. bovis		12.95	0.22		1	99		
266713	**Low discrimination**							**266713**
					COA	NOV		
Staphylococcus intermedius		74.30	0.65	GLR 3	99	1		
Staphylococcus xylosus		8.18	0.44		1	99		
Staphylococcus simulans		7.48	0.47		1	1		
266721	**Good identification**							**266721**
					COA	NOV		
Staphylococcus xylosus		97.90	0.32	MAN 94	1	99		
Staphylococcus intermedius		2.10	-.01	MLT 93	99	1		
266723	**Low discrimination**							**266723**
					NOV	COA		
Staphylococcus xylosus		85.63	0.41	MAN 94	99	1		
Staphylococcus lutrae		5.47	0.12	MLT 93	1	99		
Staphylococcus hyicus		4.38	0.11		1	24		
266731	**Very good identification**							**266731**
Staphylococcus xylosus		95.69	0.56	MLT 93				
Staphylococcus equorum		3.49	0.28					
Staphylococcus gallinarum		0.47	0.07					
266733	**Good identification**							**266733**
					COA	NOV		
Staphylococcus xylosus		94.82	0.66	MLT 93	1	99		
Staphylococcus equorum		3.12	0.36		1	99		
Staphylococcus intermedius		0.76	0.24		99	1		
266740	**Low discrimination**							**266740**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		88.28	0.34	MAN 99	1	99	99	
Staphylococcus warneri		8.51	0.23		1	1	8	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		3.20	0.03		99	1	90	
266741	**Low discrimination**							**266741**
					COA	NOV		
Staphylococcus intermedius		67.92	0.44	GLR 3	99	1		
Staphylococcus xylosus		24.68	0.33	LAC 86	1	99		
Staphylococcus saprophyticus subsp. bovis		4.73	0.22		1	99		
266743	**Low discrimination**							**266743**
					COA	NOV	YEP	
Staphylococcus intermedius		78.87	0.60	GLR 3	99	1	1	
Staphylococcus xylosus		13.28	0.42		1	99	40	
Staphylococcus chromogenes		3.92	0.32		1	1	95	
266750	**Very good identification**							**266750**
Staphylococcus saprophyticus subsp. bovis		99.51	0.75					
Staphylococcus xylosus		0.21	0.16					
Staphylococcus kloosii		0.11	0.32					
266751	**Low discrimination**							**266751**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		44.76	0.63		1	99		
Staphylococcus xylosus		36.98	0.58		1	99		
Staphylococcus intermedius		17.57	0.53		99	1		
266752	**Good identification**							**266752**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		92.89	0.63		1	99		
Staphylococcus intermedius		2.26	0.28		99	1		
Staphylococcus xylosus		2.21	0.26		1	99		
266753	**Low discrimination**							**266753**
					COA	NOV		
Staphylococcus intermedius		46.26	0.69	GLR 3	99	1		
Staphylococcus xylosus		45.13	0.67		1	99		
Staphylococcus saprophyticus subsp. bovis		4.80	0.50		1	99		
266761	**Good identification**							**266761**
					COA	NOV		
Staphylococcus xylosus		92.82	0.55	MAN 94	1	99		
Staphylococcus gallinarum		5.28	0.28		1	99		
Staphylococcus lutrae		1.27	0.12		99	1		
266763	**Low discrimination**							**266763**
					COA	NOV		
Staphylococcus xylosus		65.17	0.64	MAN 94	1	99		
Staphylococcus lutrae		31.04	0.53		99	1		
Staphylococcus gallinarum		3.19	0.36		1	99		
266770	**Low discrimination**							**266770**
					PYR			
Staphylococcus xylosus		66.79	0.38	MNS 99	65			
Staphylococcus saprophyticus subsp. bovis		28.11	0.34		99			
Staphylococcus gallinarum		2.45	0.07		12			
266771	**Good identification**							**266771**
Staphylococcus xylosus		93.67	0.79					
Staphylococcus gallinarum		3.44	0.48					
Staphylococcus equorum		2.60	0.49					
266772	**Low discrimination**							**266772**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COA	NOV	
Staphylococcus xylosus		85.72	0.48	MNS 99	1	99	
Staphylococcus lutrae		5.06	0.18		99	1	
Staphylococcus saprophyticus subsp. bovis		3.17	0.22		1	99	
266773	**Low discrimination**						**266773**
					COA	NOV	
Staphylococcus xylosus		89.54	0.89		1	99	
Staphylococcus lutrae		5.29	0.59		99	1	
Staphylococcus gallinarum		2.83	0.56		1	99	
267053	**Good identification**						**267053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99			
267153	**Low discrimination**						**267153**
Staphylococcus cohnii subsp. urealyticum		88.81	0.27	TRE 99			
Staphylococcus xylosus		11.20	0.00				
267271	**Low discrimination**						**267271**
Staphylococcus equorum		66.20	0.28	GAL 91			
Staphylococcus xylosus		24.86	0.18	TRE 99			
Staphylococcus gallinarum		8.94	0.07				
267273	**Low discrimination**						**267273**
Staphylococcus equorum		64.76	0.36	GAL 91			
Staphylococcus xylosus		26.91	0.27	TRE 99			
Staphylococcus gallinarum		8.33	0.15				
267331	**Low discrimination**						**267331**
Staphylococcus equorum		70.38	0.28	TRE 99			
Staphylococcus xylosus		20.11	0.15	MLT 91			
Staphylococcus gallinarum		9.51	0.07				
267333	**Low discrimination**						**267333**
Staphylococcus equorum		69.21	0.36	TRE 99			
Staphylococcus xylosus		21.89	0.25	MLT 91			
Staphylococcus gallinarum		8.90	0.15				
267361	**Low discrimination**						**267361**
Staphylococcus gallinarum		79.34	0.28	TRE 99			
Staphylococcus xylosus		14.54	0.14	MAN 91			
Staphylococcus equorum		6.13	0.08				
267363	**Low discrimination**						**267363**
Staphylococcus gallinarum		77.27	0.36	TRE 99			
Staphylococcus xylosus		16.46	0.23	MAN 91			
Staphylococcus equorum		6.27	0.16				
267371	**Low discrimination**						**267371**
Staphylococcus gallinarum		49.02	0.48	TRE 99			
Staphylococcus equorum		37.07	0.49				
Staphylococcus xylosus		13.92	0.39				
267373	**Low discrimination**						**267373**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus gallinarum		47.08	0.56	TRE 99		
Staphylococcus equorum		37.39	0.57			
Staphylococcus xylosus		15.54	0.48			
267403	**Good identification**					**267403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	MAN 91		
267411	**Good identification**					**267411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	LAC 91		
267413	**Very good identification**					**267413**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.81	0.61		99	
Staphylococcus simulans		0.19	-.01		1	
267441	**Good identification**					**267441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91		
267443	**Very good identification**					**267443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	MAN 91		
267450	**Low discrimination**					**267450**
Staphylococcus kloosii		82.95	0.47	NIT 1		
Staphylococcus saprophyticus subsp. bovis		8.82	0.09			
Staphylococcus cohnii subsp. urealyticum		8.24	0.14			
267451	**Very good identification**					**267451**
Staphylococcus cohnii subsp. urealyticum		99.54	0.55	LAC 91		
Staphylococcus xylosus		0.46	-.02			
267452	**Low discrimination**					**267452**
Staphylococcus kloosii		51.89	0.48	NIT 1		
Staphylococcus cohnii subsp. urealyticum		48.11	0.35			
267453	**Very good identification**					**267453**
Staphylococcus cohnii subsp. urealyticum		99.87	0.76			
Staphylococcus xylosus		0.13	0.08			
267470	**Good identification**					**267470**
Staphylococcus kloosii		100.0	0.34	NIT 1		
267471	**Low discrimination**					**267471**
Staphylococcus equorum		58.83	0.28	GAL 91		
Staphylococcus xylosus		27.90	0.20	SUC 99		
Staphylococcus gallinarum		7.95	0.07			
267472	**Good identification**					**267472**
Staphylococcus kloosii		100.0	0.35	NIT 1		
267473	**Low discrimination**					**267473**
Staphylococcus equorum		49.77	0.36	GAL 91		
Staphylococcus xylosus		26.12	0.29	SUC 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		17.71	0.35			
267503	**Good identification**					**267503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91		
267511	**Good identification**					**267511**
Staphylococcus cohnii subsp. urealyticum		95.57	0.32	LAC 91		
Staphylococcus xylosus		4.43	-.04			
267512	**Low discrimination**					**267512**
Staphylococcus kloosii		59.30	0.27	NIT 1		
Staphylococcus cohnii subsp. urealyticum		40.70	0.12	MLT 95		
267513	**Very good identification**					**267513**
Staphylococcus cohnii subsp. urealyticum		98.71	0.53			
Staphylococcus xylosus		1.29	0.05			
267531	**Low discrimination**					**267531**
Staphylococcus equorum		66.85	0.28	SUC 99		
Staphylococcus xylosus		24.13	0.18	MLT 91		
Staphylococcus gallinarum		9.03	0.07			
267533	**Low discrimination**					**267533**
Staphylococcus equorum		64.31	0.36	SUC 99		
Staphylococcus xylosus		25.69	0.27	MLT 91		
Staphylococcus gallinarum		8.27	0.15			
267541	**Good identification**					**267541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91		
267543	**Good identification**					**267543**
Staphylococcus cohnii subsp. urealyticum		97.25	0.48	MAN 91		
Staphylococcus xylosus		1.92	0.04			
Staphylococcus gallinarum		0.83	-.05			
267550	**Low discrimination**					**267550**
Staphylococcus kloosii		80.59	0.53	NIT 1		
Staphylococcus saprophyticus subsp. bovis		17.65	0.22			
Staphylococcus cohnii subsp. urealyticum		1.77	0.07			
267551	**Low discrimination**					**267551**
Staphylococcus cohnii subsp. urealyticum		83.64	0.48	LAC 91		
Staphylococcus xylosus		9.08	0.19			
Staphylococcus gallinarum		2.94	0.07			
267552	**Low discrimination**					**267552**
Staphylococcus kloosii		79.67	0.53	NIT 1		
Staphylococcus cohnii subsp. urealyticum		16.31	0.27			
Staphylococcus saprophyticus subsp. bovis		4.03	0.09			
267553	**Very good identification**					**267553**
Staphylococcus cohnii subsp. urealyticum		95.62	0.69			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		2.92	0.28				
Staphylococcus gallinarum		0.81	0.15				
267561	**Low discrimination**						**267561**
Staphylococcus gallinarum		76.41	0.28	SUC 99			
Staphylococcus xylosus		17.68	0.16	MAN 91			
Staphylococcus equorum		5.90	0.08				
267563	**Low discrimination**						**267563**
					COA	NOV	
Staphylococcus gallinarum		68.47	0.36	SUC 99	1	99	
Staphylococcus xylosus		18.42	0.25	MAN 91	1	99	
Staphylococcus lutrae		6.74	0.12		99	1	
267570	**Low discrimination**						**267570**
					YEP		
Staphylococcus kloosii		60.76	0.40	NIT 1	99		
Staphylococcus gallinarum		18.56	0.07		78		
Staphylococcus equorum		14.03	0.08		1		
267571	**Low discrimination**						**267571**
Staphylococcus gallinarum		47.22	0.48	SUC 99			
Staphylococcus equorum		35.71	0.49				
Staphylococcus xylosus		16.93	0.41				
267572	**Low discrimination**						**267572**
					YEP		
Staphylococcus kloosii		39.59	0.41	NIT 1	99		
Staphylococcus gallinarum		27.32	0.15		78		
Staphylococcus equorum		21.70	0.16		1		
267573	**Low discrimination**						**267573**
Staphylococcus gallinarum		44.61	0.56	SUC 99			
Staphylococcus equorum		35.42	0.57				
Staphylococcus xylosus		18.59	0.50				
267613	**Low discrimination**						**267613**
					NOV		
Staphylococcus simulans		44.02	0.27	ARG 92	1		
Staphylococcus cohnii subsp. urealyticum		30.20	0.30	ESL 1	99		
Staphylococcus equorum		15.53	0.16		99		
267631	**Acceptable identification**						**267631**
Staphylococcus equorum		92.33	0.48	GAL 91			
Staphylococcus xylosus		6.40	0.23	MLT 91			
Staphylococcus gallinarum		1.27	0.07				
267633	**Good identification**						**267633**
Staphylococcus equorum		91.76	0.57	GAL 91			
Staphylococcus xylosus		7.04	0.33	MLT 91			
Staphylococcus gallinarum		1.21	0.15				
267651	**Low discrimination**						**267651**
Staphylococcus equorum		43.62	0.28	GAL 91			
Staphylococcus xylosus		34.18	0.24	XYL 99			
Staphylococcus cohnii subsp. urealyticum		12.09	0.24				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
267652	**Low discrimination**						**267652**
					NOV	YEP	
Staphylococcus kloosii		33.53	0.27	NIT 1	99	99	
Staphylococcus saprophyticus subsp. bovis		33.26	0.09	SUC 9	99	80	
Staphylococcus simulans		33.21	0.09		1	1	
267653	**Low discrimination**						**267653**
Staphylococcus cohnii subsp. urealyticum		33.93	0.45	SUC 3			
Staphylococcus equorum		31.14	0.36				
Staphylococcus xylosus		27.00	0.34				
267661	**Low discrimination**						**267661**
Staphylococcus gallinarum		45.64	0.28	GAL 99			
Staphylococcus equorum		34.51	0.28	MAN 91			
Staphylococcus xylosus		19.85	0.22				
267663	**Low discrimination**						**267663**
Staphylococcus gallinarum		43.48	0.36	GAL 99			
Staphylococcus equorum		34.53	0.36	MAN 91			
Staphylococcus xylosus		21.98	0.31				
267670	**Low discrimination**						**267670**
Staphylococcus equorum		81.56	0.28	GAL 91			
Staphylococcus gallinarum		11.02	0.07	MNS 99			
Staphylococcus xylosus		7.42	0.05				
267671	**Low discrimination**						**267671**
Staphylococcus equorum		81.56	0.69	GAL 91			
Staphylococcus gallinarum		11.02	0.48				
Staphylococcus xylosus		7.42	0.46				
267672	**Low discrimination**						**267672**
Staphylococcus equorum		81.34	0.36	GAL 91			
Staphylococcus gallinarum		10.46	0.15	MNS 99			
Staphylococcus xylosus		8.20	0.15				
267673	**Low discrimination**						**267673**
Staphylococcus equorum		81.34	0.78	GAL 91			
Staphylococcus gallinarum		10.46	0.56				
Staphylococcus xylosus		8.20	0.56				
267711	**Low discrimination**						**267711**
Staphylococcus equorum		54.56	0.28	XYL 99			
Staphylococcus xylosus		32.54	0.22	MLT 91			
Staphylococcus gallinarum		7.37	0.07				
267713	**Low discrimination**						**267713**
					COA	NOV	
Staphylococcus equorum		46.29	0.36	XYL 99	1	99	
Staphylococcus xylosus		30.55	0.31	MLT 91	1	99	
Staphylococcus intermedius		11.21	0.24		99	1	
267721	**Low discrimination**						**267721**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus gallinarum		47.91	0.28	MAN 91			
Staphylococcus equorum		36.23	0.28	MLT 99			
Staphylococcus xylosus		15.86	0.19				
267723	**Low discrimination**						**267723**
Staphylococcus gallinarum		45.89	0.36	MAN 91			
Staphylococcus equorum		36.45	0.36	MLT 99			
Staphylococcus xylosus		17.66	0.29				
267730	**Low discrimination**						**267730**
Staphylococcus equorum		83.03	0.28	MLT 91			
Staphylococcus gallinarum		11.22	0.07	MNS 99			
Staphylococcus xylosus		5.75	0.03				
267731	**Low discrimination**						**267731**
Staphylococcus equorum		83.03	0.69	MLT 91			
Staphylococcus gallinarum		11.22	0.48				
Staphylococcus xylosus		5.75	0.44				
267732	**Low discrimination**						**267732**
Staphylococcus equorum		82.97	0.36	MLT 91			
Staphylococcus gallinarum		10.67	0.15	MNS 99			
Staphylococcus xylosus		6.36	0.12				
267733	**Low discrimination**						**267733**
Staphylococcus equorum		82.97	0.78	MLT 91			
Staphylococcus gallinarum		10.67	0.56				
Staphylococcus xylosus		6.36	0.53				
267741	**Low discrimination**						**267741**
Staphylococcus gallinarum		66.57	0.28	MAN 91			
Staphylococcus xylosus		25.46	0.21	XYL 99			
Staphylococcus equorum		5.14	0.08				
267743	**Low discrimination**						**267743**
					COA	NOV	
Staphylococcus gallinarum		59.92	0.36	MAN 91	1	99	
Staphylococcus xylosus		26.63	0.30	XYL 99	1	99	
Staphylococcus intermedius		6.39	0.19		99	1	
267750	**Low discrimination**						**267750**
					PYR		
Staphylococcus saprophyticus subsp. bovis		75.89	0.34	ESL 1	99		
Staphylococcus kloosii		8.57	0.32		52		
Staphylococcus gallinarum		6.62	0.07		12		
267751	**Low discrimination**						**267751**
Staphylococcus gallinarum		41.66	0.48	XYL 99			
Staphylococcus equorum		31.50	0.49				
Staphylococcus xylosus		24.68	0.45				
267753	**Low discrimination**						**267753**
Staphylococcus gallinarum		39.18	0.56	XYL 99			
Staphylococcus equorum		31.11	0.57				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		26.98	0.55			
267760	**Low discrimination**					**267760**
Staphylococcus gallinarum		89.16	0.28	MAN 91		
Staphylococcus equorum		6.89	0.08	MNS 99		
Staphylococcus xylosus		3.96	0.01			
267761	**Low discrimination**					**267761**
Staphylococcus gallinarum		89.16	0.69	MAN 91		
Staphylococcus equorum		6.89	0.49			
Staphylococcus xylosus		3.96	0.42			
267762	**Low discrimination**					**267762**
Staphylococcus gallinarum		88.28	0.36	MAN 91		
Staphylococcus equorum		7.16	0.16	MNS 99		
Staphylococcus xylosus		4.56	0.11			
267763	**Low discrimination**					**267763**
Staphylococcus gallinarum		88.20	0.77	MAN 91		
Staphylococcus equorum		7.15	0.57			
Staphylococcus xylosus		4.55	0.52			
267770	**Low discrimination**					**267770**
Staphylococcus gallinarum		54.70	0.48	MNS 99		
Staphylococcus equorum		41.36	0.49			
Staphylococcus xylosus		3.76	0.26			
267771	**Low discrimination**					**267771**
Staphylococcus gallinarum		54.80	0.90			
Staphylococcus equorum		41.43	0.90			
Staphylococcus xylosus		3.77	0.67			
267772	**Low discrimination**					**267772**
Staphylococcus gallinarum		53.23	0.56	MNS 99		
Staphylococcus equorum		42.27	0.57			
Staphylococcus xylosus		4.26	0.35			
267773	**Low discrimination**					**267773**
Staphylococcus gallinarum		53.35	0.98			
Staphylococcus equorum		42.37	0.98			
Staphylococcus xylosus		4.27	0.76			
270413	**Good identification**					**270413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	ORN 1		
270443	**Good identification**					**270443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1		
270451	**Good identification**					**270451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1		
270453	**Very good identification**					**270453**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	ORN 1			
270513	**Good identification**						**270513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	ORN 1			
270543	**Good identification**						**270543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1			
270551	**Good identification**						**270551**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1			
270553	**Good identification**						**270553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	ORN 1			
270640	**Good identification**						**270640**
Staphylococcus warneri		100.0	0.41	ORN 1			
270642	**Good identification**						**270642**
Staphylococcus warneri		100.0	0.31	ORN 1			
270650	**Acceptable identification**						**270650**
					NOV		
Staphylococcus warneri		93.08	0.38	ORN 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		6.92	-.06		99		
270740	**Good identification**						**270740**
Staphylococcus warneri		100.0	0.30	ORN 1			
270750	**Low discrimination**						**270750**
					NOV	PYR	
Staphylococcus warneri		80.13	0.28	ORN 1	1	8	
Staphylococcus saprophyticus subsp. bovis		19.87	0.05		99	99	
270771	**Good identification**						**270771**
Staphylococcus xylosus		100.0	0.31	ORN 1			
270773	**Good identification**						**270773**
Staphylococcus xylosus		98.14	0.40	ORN 1			
Staphylococcus arlettae		1.86	0.14	NIT 89			
271413	**Good identification**						**271413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1			
271453	**Good identification**						**271453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	ORN 1			
271553	**Good identification**						**271553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ORN 1			
271773	**Low discrimination**						**271773**
					YEP		
Staphylococcus equorum		54.97	0.32	ORN 1	1		
Staphylococcus xylosus		41.86	0.28	NIT 92	40		
Staphylococcus arlettae		3.17	0.14		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
272373	**Good identification**						**272373**
	Staphylococcus xylosus	100.0	0.31	ORN 1			
272453	**Good identification**						**272453**
					NOV		
	Staphylococcus cohnii subsp. urealyticum	97.49	0.40	ORN 1	99		
	Staphylococcus simulans	2.52	0.02		1		
272550	**Good identification**						**272550**
	Staphylococcus saprophyticus subsp. bovis	100.0	0.34	ORN 1			
272553	**Low discrimination**						**272553**
	Staphylococcus cohnii subsp. urealyticum	71.59	0.32	ORN 1			
	Staphylococcus xylosus	18.10	0.11				
	Staphylococcus saprophyticus subsp. bovis	10.31	0.09				
272573	**Good identification**						**272573**
	Staphylococcus xylosus	100.0	0.33	ORN 1			
272613	**Good identification**						**272613**
	Staphylococcus simulans	100.0	0.37	ARG 92			
272640	**Good identification**						**272640**
	Staphylococcus warneri	100.0	0.33	ORN 1			
272650	**Low discrimination**						**272650**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	81.89	0.34	ORN 1	99	99	
	Staphylococcus warneri	18.11	0.30		1	8	
272653	**Low discrimination**						**272653**
					NOV		
	Staphylococcus simulans	67.64	0.30	ARG 92	1		
	Staphylococcus xylosus	22.24	0.16	ORN 1	99		
	Staphylococcus saprophyticus subsp. bovis	6.74	0.09		99		
272671	**Good identification**						**272671**
	Staphylococcus xylosus	100.0	0.29	ORN 1			
272673	**Good identification**						**272673**
	Staphylococcus xylosus	100.0	0.38	ORN 1			
272731	**Good identification**						**272731**
	Staphylococcus xylosus	100.0	0.26	ORN 1			
272733	**Good identification**						**272733**
	Staphylococcus xylosus	100.0	0.36	ORN 1			
272750	**Good identification**						**272750**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	98.28	0.47	ORN 1	99	99	
	Staphylococcus warneri	1.72	0.20		1	8	
272751	**Low discrimination**						**272751**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		57.72	0.34	ORN 1			
Staphylococcus xylosus		42.28	0.28				
272752	**Good identification**						**272752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		95.76	0.34	ORN 1	99	99	
Staphylococcus warneri		2.23	0.10		1	8	
Staphylococcus xylosus		2.02	-.04		99	65	
272753	**Low discrimination**						**272753**
					NOV		
Staphylococcus xylosus		87.25	0.37	ORN 1	99		
Staphylococcus saprophyticus subsp. bovis		10.46	0.22	XYL 92	99		
Staphylococcus simulans		2.28	0.08		1		
272763	**Good identification**						**272763**
Staphylococcus xylosus		100.0	0.34	ORN 1			
272771	**Good identification**						**272771**
Staphylococcus xylosus		99.48	0.50	ORN 1			
Staphylococcus equorum		0.52	0.04				
272773	**Very good identification**						**272773**
Staphylococcus xylosus		99.53	0.59	ORN 1			
Staphylococcus equorum		0.47	0.12				
273453	**Good identification**						**273453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1			
273673	**Low discrimination**						**273673**
Staphylococcus equorum		65.12	0.33	ORN 1			
Staphylococcus xylosus		34.89	0.26	GAL 91			
273733	**Low discrimination**						**273733**
Staphylococcus equorum		71.04	0.33	ORN 1			
Staphylococcus xylosus		28.96	0.23	MLT 91			
273771	**Low discrimination**						**273771**
Staphylococcus equorum		66.49	0.45	ORN 1			
Staphylococcus xylosus		32.18	0.37				
Staphylococcus gallinarum		1.33	0.07				
273773	**Low discrimination**						**273773**
Staphylococcus equorum		64.32	0.54	ORN 1			
Staphylococcus xylosus		34.46	0.47				
Staphylococcus gallinarum		1.23	0.15				
274413	**Good identification**						**274413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ORN 1			
274443	**Good identification**						**274443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	ORN 1			
274450	**Good identification**						**274450**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.47	ORN 1		
274451	**Good identification**					**274451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	ORN 1		
274452	**Low discrimination**					**274452**
Staphylococcus kloosii		87.99	0.48	ORN 1		
Staphylococcus cohnii subsp. urealyticum		12.01	0.18			
274453	**Very good identification**					**274453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	ORN 1		
274470	**Good identification**					**274470**
Staphylococcus kloosii		100.0	0.34	ORN 1		
274472	**Good identification**					**274472**
Staphylococcus kloosii		100.0	0.35	ORN 1		
274512	**Good identification**					**274512**
Staphylococcus kloosii		100.0	0.27	ORN 1		
274513	**Good identification**					**274513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	ORN 1		
274543	**Good identification**					**274543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
274550	**Good identification**					**274550**
Staphylococcus kloosii		100.0	0.53	ORN 1		
274551	**Good identification**					**274551**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
274552	**Very good identification**					**274552**
Staphylococcus kloosii		97.08	0.53	ORN 1		
Staphylococcus cohnii subsp. urealyticum		2.92	0.10			
274553	**Good identification**					**274553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	ORN 1		
274570	**Good identification**					**274570**
Staphylococcus kloosii		100.0	0.40	ORN 1		
274572	**Good identification**					**274572**
Staphylococcus kloosii		100.0	0.41	ORN 1		
274652	**Good identification**					**274652**
Staphylococcus kloosii		100.0	0.27	ORN 1		
274653	**Good identification**					**274653**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
274750	**Good identification**					**274750**
Staphylococcus kloosii		100.0	0.32	ORN 1		
274752	**Good identification**					**274752**
Staphylococcus kloosii		100.0	0.33	ORN 1		
274773	**Acceptable identification**					**274773**
Staphylococcus xylosus		90.85	0.29	ORN 1		
Staphylococcus arlettae		9.15	0.18	NIT 89		
275413	**Good identification**					**275413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1		
275443	**Good identification**					**275443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
275450	**Good identification**					**275450**
Staphylococcus kloosii		100.0	0.47	ORN 1		
275451	**Good identification**					**275451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
275452	**Acceptable identification**					**275452**
Staphylococcus kloosii		94.96	0.48	ORN 1		
Staphylococcus cohnii subsp. urealyticum		5.04	0.09			
275453	**Good identification**					**275453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	ORN 1		
275470	**Good identification**					**275470**
Staphylococcus kloosii		100.0	0.34	ORN 1		
275472	**Good identification**					**275472**
Staphylococcus kloosii		100.0	0.35	ORN 1		
275512	**Good identification**					**275512**
Staphylococcus kloosii		100.0	0.27	ORN 1		
275513	**Good identification**					**275513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1		
275550	**Good identification**					**275550**
Staphylococcus kloosii		100.0	0.53	ORN 1		
275552	**Very good identification**					**275552**
Staphylococcus kloosii		100.0	0.53	ORN 1		
275553	**Good identification**					**275553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ORN 1		
275570	**Good identification**					**275570**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.40	ORN 1			
275572	**Good identification**						**275572**
Staphylococcus kloosii		100.0	0.41	ORN 1			
275652	**Good identification**						**275652**
Staphylococcus kloosii		100.0	0.27	ORN 1			
275750	**Good identification**						**275750**
Staphylococcus kloosii		100.0	0.32	ORN 1			
275752	**Good identification**						**275752**
Staphylococcus kloosii		100.0	0.33	ORN 1			
275771	**Low discrimination**						**275771**
Staphylococcus equorum		77.94	0.27	ORN 1			
Staphylococcus gallinarum		11.97	0.07	NIT 92			
Staphylococcus xylosus		10.08	0.07				
275773	**Low discrimination**						**275773**
Staphylococcus equorum		74.23	0.35	ORN 1			
Staphylococcus gallinarum		10.86	0.15	NIT 92			
Staphylococcus xylosus		10.63	0.17				
276413	**Acceptable identification**						**276413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.26	0.28	ORN 1	99		
Staphylococcus simulans		6.74	-.01		1		
276453	**Good identification**						**276453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ORN 1			
276553	**Good identification**						**276553**
Staphylococcus cohnii subsp. urealyticum		95.46	0.36	ORN 1			
Staphylococcus xylosus		4.54	-.01				
276613	**Good identification**						**276613**
Staphylococcus simulans		100.0	0.27	ARG 92			
276673	**Good identification**						**276673**
Staphylococcus xylosus		100.0	0.27	ORN 1			
276750	**Good identification**						**276750**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
276753	**Low discrimination**						**276753**
					COA	NOV	
Staphylococcus intermedius		47.34	0.28	ORN 1	99	1	
Staphylococcus xylosus		46.17	0.26	GLR 3	1	99	
Staphylococcus saprophyticus subsp. bovis		4.91	0.09		1	99	
276771	**Acceptable identification**						**276771**
Staphylococcus xylosus		93.95	0.38	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus gallinarum		3.45	0.07				
Staphylococcus equorum		2.61	0.08				
276773	**Low discrimination**						**276773**
					COA	NOV	
Staphylococcus xylosus		89.63	0.48	ORN	1	99	
Staphylococcus lutrae		5.29	0.18		99	1	
Staphylococcus gallinarum		2.83	0.15		1	99	
277453	**Good identification**						**277453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN			
277553	**Good identification**						**277553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN			
277671	**Low discrimination**						**277671**
Staphylococcus equorum		81.56	0.28	ORN			
Staphylococcus gallinarum		11.02	0.07	GAL	91		
Staphylococcus xylosus		7.42	0.05				
277673	**Low discrimination**						**277673**
Staphylococcus equorum		81.34	0.36	ORN			
Staphylococcus gallinarum		10.46	0.15	GAL	91		
Staphylococcus xylosus		8.20	0.15				
277731	**Low discrimination**						**277731**
Staphylococcus equorum		83.03	0.28	ORN			
Staphylococcus gallinarum		11.22	0.07	MLT	91		
Staphylococcus xylosus		5.75	0.03				
277733	**Low discrimination**						**277733**
Staphylococcus equorum		82.97	0.36	ORN			
Staphylococcus gallinarum		10.67	0.15	MLT	91		
Staphylococcus xylosus		6.36	0.12				
277761	**Low discrimination**						**277761**
Staphylococcus gallinarum		89.16	0.28	ORN			
Staphylococcus equorum		6.89	0.08	MAN	91		
Staphylococcus xylosus		3.96	0.01				
277763	**Low discrimination**						**277763**
Staphylococcus gallinarum		88.28	0.36	ORN			
Staphylococcus equorum		7.16	0.16	MAN	91		
Staphylococcus xylosus		4.56	0.11				
277771	**Low discrimination**						**277771**
Staphylococcus gallinarum		54.80	0.48	ORN			
Staphylococcus equorum		41.43	0.49				
Staphylococcus xylosus		3.77	0.26				
277773	**Low discrimination**						**277773**
Staphylococcus gallinarum		53.36	0.56	ORN			
Staphylococcus equorum		42.37	0.57				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus xylosus		4.27	0.35					
300000	**Low discrimination**							**300000**
					OXI			
Dermacoccus nishinomiyaensis		66.63	0.74	VPT 5	93			
Micrococcus sp.		21.00	0.54		89			
Kocuria varians		11.88	0.66		13			
300001	**Low discrimination**							**300001**
					OXI			
Kocuria varians		47.20	0.44	VPT 14	13			
Dermacoccus nishinomiyaensis		30.75	0.33	MNS 8	93			
Micrococcus sp.		19.57	0.19		89			
300002	**Low discrimination**							**300002**
					OXI			
Kocuria varians		79.11	0.55	VPT 14	13			
Dermacoccus nishinomiyaensis		15.89	0.33		93			
Micrococcus sp.		5.01	0.13		89			
300003	**Good identification**							**300003**
Kocuria varians		100.0	0.33	VPT 14				
300010	**Low discrimination**							**300010**
					YEP			
Macrococcus equipersicus		43.14	0.43	VPT 1	99			
Dermacoccus nishinomiyaensis		22.80	0.33		1			
Macrococcus bovicus		20.89	0.35		99			
300011	**Low discrimination**							**300011**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		89.51	0.27	URE 1	1	1	1	
Macrococcus equipersicus		3.78	0.02	NIT 89	99	99	99	
Staphylococcus cohnii subsp. cohnii		3.64	0.07		1	99	12	
300020	**Low discrimination**							**300020**
					OXI			
Kocuria varians		79.75	0.57	VPT 14	13			
Dermacoccus nishinomiyaensis		12.86	0.33		93			
Micrococcus sp.		4.05	0.13		89			
300021	**Good identification**							**300021**
Kocuria varians		97.52	0.35	VPT 14				
Kocuria rosea		2.48	-.07	MNS 8				
300022	**Good identification**							**300022**
Kocuria varians		100.0	0.45	VPT 14				
300040	**Low discrimination**							**300040**
					OXI			
Kocuria varians		44.94	0.52	VPT 14	13			
Micrococcus sp.		27.23	0.31		89			
Dermacoccus nishinomiyaensis		11.60	0.33		93			
300041	**Low discrimination**							**300041**
					NOV	OXI	PYR	
Kocuria varians		43.89	0.31	VPT 14	1	13	10	
Staphylococcus hominis subsp. novobiosepticus		29.20	0.17	MNS 8	99	1	1	
Kocuria kristinae		14.70	0.13		1	99	93	
300042	**Low discrimination**							**300042**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Kocuria varians		64.37	0.41	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		28.44	0.24		99		
Staphylococcus warneri		3.01	0.21		1		
300050	**Low discrimination**						**300050**
					OXI		
Macrococcus equipersicus		67.34	0.52	VPT 1	99		
Macrococcus bovicus		28.14	0.43		99		
Staphylococcus saprophyticus subsp. saprophyticus		1.52	0.07		1		
300060	**Good identification**						**300060**
					OXI		
Kocuria varians		98.29	0.43	VPT 14	13		
Micrococcus sp.		1.71	-.10		89		
300062	**Good identification**						**300062**
Kocuria varians		100.0	0.32	VPT 14			
300100	**Low discrimination**						**300100**
					OXI		
Kocuria varians		57.92	0.70	VPT 14	13		
Dermacoccus nishinomiyaensis		41.24	0.59		93		
Micrococcus sp.		0.69	0.13		89		
300101	**Acceptable identification**						**300101**
					OXI		
Kocuria varians		92.36	0.48	VPT 14	13		
Dermacoccus nishinomiyaensis		7.64	0.18	MNS 8	93		
300102	**Very good identification**						**300102**
					OXI	NOV	
Kocuria varians		97.33	0.58	VPT 14	13	1	
Dermacoccus nishinomiyaensis		2.48	0.18		93	80	
Staphylococcus hominis subsp. novobiosepticus		0.19	-.07		1	99	
300103	**Good identification**						**300103**
Kocuria varians		100.0	0.36	VPT 14			
300110	**Low discrimination**						**300110**
					NOV	OXI	
Macrococcus equipersicus		48.81	0.36	VPT 1	99	99	
Macrococcus bovicus		23.63	0.28		99	99	
Kocuria varians		16.10	0.29		1	13	
300120	**Very good identification**						**300120**
					OXI		
Kocuria varians		97.99	0.60	VPT 14	13		
Dermacoccus nishinomiyaensis		2.01	0.18		93		
300121	**Good identification**						**300121**
Kocuria varians		100.0	0.38	VPT 14			
300122	**Good identification**						**300122**
Kocuria varians		100.0	0.49	VPT 14			
300123	**Good identification**						**300123**
Kocuria varians		100.0	0.27	VPT 14			
300140	**Low discrimination**						**300140**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV		
Kocuria varians		65.32	0.56	VPT 14	13	1		
Staphylococcus hominis subsp. novobiosepticus		30.23	0.39		1	99		
Dermacoccus nishinomiyaensis		2.14	0.18		93	80		
300141	**Low discrimination**							**300141**
					NOV	OXI	PYR	
Staphylococcus hominis subsp. novobiosepticus		54.66	0.27	NIT 90	99	1	1	
Kocuria varians		41.08	0.34	SUC 99	1	13	10	
Kocuria kristinae		3.06	0.03		1	99	93	
300142	**Low discrimination**							**300142**
					NOV			
Kocuria varians		52.03	0.45	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		45.97	0.34		99			
Staphylococcus epidermidis		0.80	0.05		1			
300150	**Low discrimination**							**300150**
					NOV	OXI		
Macrococcus equipersicus		67.89	0.45	VPT 1	99	99		
Macrococcus bovicus		28.37	0.35		99	99		
Kocuria varians		1.82	0.15		1	13		
300160	**Good identification**							**300160**
					NOV			
Kocuria varians		98.69	0.47	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		1.31	-.02		99			
300162	**Good identification**							**300162**
					NOV			
Kocuria varians		97.52	0.35	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		2.48	-.07		99			
300200	**Low discrimination**							**300200**
					OXI			
Dermacoccus nishinomiyaensis		48.81	0.49	VPT 5	93			
Micrococcus sp.		23.83	0.33	SUC 6	89			
Kocuria varians		18.60	0.48		13			
300201	**Low discrimination**							**300201**
					NOV	OXI	PYR	
Kocuria kristinae		66.57	0.34	URE 10	1	99	93	
Staphylococcus hominis subsp. novobiosepticus		12.02	0.17	TRE 95	99	1	1	
Kocuria varians		11.22	0.26	MLT 90	1	13	10	
300202	**Low discrimination**							**300202**
					OXI	NOV		
Kocuria varians		51.12	0.37	VPT 14	13	1		
Staphylococcus hominis subsp. novobiosepticus		36.36	0.24	SUC 12	1	99		
Dermacoccus nishinomiyaensis		4.81	0.08		93	80		
300211	**Acceptable identification**							**300211**
					OXI	PYR		
Staphylococcus capitis subsp. capitis		92.77	0.32	URE 1	1	1		
Kocuria kristinae		5.71	0.06	NIT 89	99	93		
Staphylococcus capitis subsp. ureolyticus		1.52	-.14		1	1		
300220	**Low discrimination**							**300220**
					OXI			
Kocuria varians		87.22	0.39	VPT 14	13			
Dermacoccus nishinomiyaensis		6.58	0.08	SUC 12	93			
Micrococcus sp.		3.21	-.08		89			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
300222	**Good identification**							**300222**
Kocuria varians		100.0	0.28	VPT 14				
300240	**Good identification**							**300240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		93.28	0.71	NIT 90	99			
Staphylococcus warneri		3.06	0.57		1			
Staphylococcus hominis subsp. hominis		1.79	0.47		1			
300241	**Low discrimination**							**300241**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		65.66	0.58	NIT 90	1	99	1	
Kocuria kristinae		33.06	0.54		99	1	93	
Staphylococcus epidermidis		0.77	0.26		1	1	1	
300242	**Good identification**							**300242**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		92.52	0.65	NIT 90	99			
Staphylococcus hominis subsp. hominis		3.87	0.48		1			
Staphylococcus warneri		1.88	0.47		1			
300243	**Low discrimination**							**300243**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		75.94	0.53	NIT 90	1	99	1	
Kocuria kristinae		17.75	0.42		99	1	93	
Staphylococcus epidermidis		5.20	0.37		1	1	1	
300250	**Low discrimination**							**300250**
					NOV			
Staphylococcus warneri		38.38	0.54	TRE 95	1			
Staphylococcus saprophyticus subsp. saprophyticus		21.17	0.28		99			
Staphylococcus hominis subsp. novobiosepticus		16.31	0.30		99			
300252	**Low discrimination**							**300252**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		81.25	0.43	TRE 99	99			
Staphylococcus warneri		8.66	0.44		1			
Staphylococcus hominis subsp. novobiosepticus		5.95	0.24		99			
300253	**Low discrimination**							**300253**
					OXI	NOV	PYR	
Staphylococcus capitis subsp. ureolyticus		88.39	0.33	ARG 99	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		4.41	0.12	NIT 95	1	99	1	
Kocuria kristinae		4.25	0.13		99	1	93	
300260	**Low discrimination**							**300260**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		79.42	0.30	NIT 90	99			
Kocuria varians		16.44	0.25	XYL 1	1			
Staphylococcus warneri		2.61	0.16		1			
300300	**Low discrimination**							**300300**
					OXI	NOV		
Kocuria varians		47.49	0.52	VPT 14	13	1		
Staphylococcus hominis subsp. novobiosepticus		35.38	0.39	SUC 12	1	99		
Dermacoccus nishinomiyaensis		15.83	0.34		93	80		
300301	**Low discrimination**							**300301**
					NOV	OXI	PYR	
Staphylococcus hominis subsp. novobiosepticus		47.52	0.27	NIT 90	99	1	1	
Kocuria kristinae		29.25	0.25	MLT 99	1	99	93	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		22.19	0.30		1	13	10	
300302	**Low discrimination**							**300302**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		57.25	0.34	NIT 90	99			
Kocuria varians		40.25	0.41	MLT 99	1			
Staphylococcus epidermidis		1.00	0.05		1			
300320	**Good identification**							**300320**
					NOV			
Kocuria varians		97.90	0.42	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		2.10	-.02	SUC 12	99			
300322	**Good identification**							**300322**
					NOV			
Kocuria varians		96.07	0.31	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		3.93	-.07	SUC 12	99			
300340	**Very good identification**							**300340**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		98.38	0.80	NIT 90	99			
Staphylococcus hominis subsp. hominis		0.68	0.47		1			
Staphylococcus warneri		0.34	0.47		1			
300341	**Good identification**							**300341**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		92.59	0.68	NIT 90	1	99	1	
Kocuria kristinae		5.18	0.44		99	1	93	
Staphylococcus epidermidis		2.04	0.42		1	1	1	
300342	**Very good identification**							**300342**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		96.49	0.75	NIT 90	99			
Staphylococcus epidermidis		1.68	0.47		1			
Staphylococcus hominis subsp. hominis		1.46	0.49		1			
300343	**Low discrimination**							**300343**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		86.42	0.62	NIT 90	1	99	1	
Staphylococcus epidermidis		11.19	0.52		1	1	1	
Kocuria kristinae		2.25	0.32		99	1	93	
300350	**Low discrimination**							**300350**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		64.09	0.39	NIT 90	99			
Staphylococcus warneri		15.87	0.44	MAN 1	1			
Staphylococcus hominis subsp. hominis		6.57	0.30		1			
300351	**Low discrimination**							**300351**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		78.55	0.27	NIT 90	1	99	1	
Kocuria kristinae		18.13	0.16	MAN 1	99	1	93	
Staphylococcus epidermidis		1.73	0.01		1	1	1	
300352	**Low discrimination**							**300352**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		55.86	0.34	NIT 90	99			
Staphylococcus saprophyticus subsp. saprophyticus		22.12	0.22	MAN 1	99			
Staphylococcus hominis subsp. hominis		12.49	0.32		1			
300360	**Acceptable identification**							**300360**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		90.05	0.39	NIT 90	99			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		9.32	0.29	XYL 1	1			
Staphylococcus hominis subsp. hominis		0.62	0.06		1			
300361	**Low discrimination**							**300361**
					NOV	OXI	PYR	
Staphylococcus hominis subsp. novobiosepticus		89.77	0.27	NIT 90	99	1	1	
Kocuria kristinae		5.02	0.03	XYL 1	1	99	93	
Kocuria varians		3.23	0.07		1	13	10	
300362	**Acceptable identification**							**300362**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		92.02	0.34	NIT 90	99			
Kocuria varians		4.99	0.18	XYL 1	1			
Staphylococcus epidermidis		1.60	0.05		1			
300400	**Low discrimination**							**300400**
					OXI	NOV		
Kocuria varians		31.15	0.43	VPT 14	13	1		
Staphylococcus cohnii subsp. cohnii		31.02	0.45	TRE 7	1	99		
Dermacoccus nishinomiyaensis		23.44	0.33		93	80		
300401	**Low discrimination**							**300401**
					NOV	OXI	PYR	
Staphylococcus cohnii subsp. cohnii		80.20	0.46	URE 1	99	1	1	
Kocuria kristinae		13.33	0.20		1	99	93	
Kocuria varians		6.47	0.21		1	13	10	
300402	**Low discrimination**							**300402**
					NOV			
Kocuria varians		80.35	0.32	VPT 14	1			
Staphylococcus warneri		10.96	0.21	TRE 7	1			
Staphylococcus saprophyticus subsp. saprophyticus		5.83	-.06		99			
300410	**Low discrimination**							**300410**
					OXI	YEP		
Macrococcus equipersicus		36.46	0.43	VPT 1	99	99		
Staphylococcus cohnii subsp. cohnii		32.45	0.47		1	12		
Macrococcus bovicus		26.48	0.39		99	99		
300411	**Good identification**							**300411**
					OXI	YEP		
Staphylococcus cohnii subsp. cohnii		96.86	0.48	URE 1	1	12		
Staphylococcus cohnii subsp. urealyticum		1.39	0.11		1	64		
Macrococcus equipersicus		1.02	0.02		99	99		
300413	**Acceptable identification**							**300413**
Staphylococcus cohnii subsp. urealyticum		93.50	0.31	GLR 99				
Staphylococcus cohnii subsp. cohnii		6.50	0.07					
300420	**Good identification**							**300420**
					NOV			
Kocuria varians		97.22	0.34	VPT 14	1			
Staphylococcus cohnii subsp. cohnii		2.78	0.04	TRE 7	99			
300440	**Low discrimination**							**300440**
					NOV	PYR		
Staphylococcus cohnii subsp. cohnii		46.69	0.55	URE 1	99	1		
Staphylococcus warneri		32.47	0.57		1	8		
Staphylococcus petrasii subsp. pragensis		7.00	0.18		1	99		
300441	**Low discrimination**							**300441**
					OXI	NOV	PYR	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. cohnii		66.71	0.56	URE 1	1	99	1	
Kocuria kristinae		31.52	0.39		99	1	93	
Staphylococcus warneri		0.87	0.22		1	1	8	
300442	**Low discrimination**							**300442**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		69.88	0.35	SUC 91	99			
Staphylococcus warneri		25.22	0.47	MAN 96	1			
Kocuria varians		2.14	0.18		1			
300443	**Low discrimination**							**300443**
					OXI	NOV		
Kocuria kristinae		68.96	0.27	URE 10	99	1		
Staphylococcus cohnii subsp. urealyticum		19.10	0.26	SUC 99	1	99		
Staphylococcus cohnii subsp. cohnii		5.90	0.15		1	99		
300450	**Low discrimination**							**300450**
					OXI	YEP		
Staphylococcus saprophyticus subsp. saprophyticus		41.68	0.48	SUC 91	1	16		
Staphylococcus cohnii subsp. cohnii		19.50	0.58		1	12		
Macrococcus equipersicus		18.70	0.52		99	99		
300451	**Good identification**							**300451**
Staphylococcus cohnii subsp. cohnii		91.92	0.58	URE 1				
Staphylococcus cohnii subsp. urealyticum		2.36	0.26					
Staphylococcus saprophyticus subsp. saprophyticus		1.83	0.07					
300452	**Very good identification**							**300452**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		98.56	0.64	SUC 91	99			
Staphylococcus warneri		1.07	0.44		1			
Staphylococcus cohnii subsp. cohnii		0.08	0.16		99			
300453	**Low discrimination**							**300453**
Staphylococcus cohnii subsp. urealyticum		66.77	0.47	GLR 99				
Staphylococcus saprophyticus subsp. saprophyticus		29.01	0.23					
Staphylococcus cohnii subsp. cohnii		2.59	0.17					
300500	**Low discrimination**							**300500**
					OXI	NOV		
Kocuria varians		82.30	0.47	VPT 14	13	1		
Dermacoccus nishinomiyaensis		7.86	0.18	TRE 7	93	80		
Staphylococcus cohnii subsp. cohnii		6.07	0.25		1	99		
300502	**Good identification**							**300502**
Kocuria varians		97.21	0.35	VPT 14				
Staphylococcus warneri		2.79	0.11	TRE 7				
300510	**Low discrimination**							**300510**
					OXI	YEP		
Macrococcus equipersicus		49.38	0.36	VPT 1	99	99		
Macrococcus bovicus		35.87	0.31		99	99		
Staphylococcus cohnii subsp. cohnii		11.40	0.28		1	12		
300511	**Good identification**							**300511**
Staphylococcus cohnii subsp. cohnii		100.0	0.28	URE 1				
300520	**Good identification**							**300520**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.37	VPT 14				
300540	**Low discrimination**							**300540**
					NOV			
Staphylococcus warneri		40.46	0.47	SUC 95	1			
Staphylococcus cohnii subsp. cohnii		20.47	0.36		99			
Kocuria varians		19.24	0.33		1			
300541	**Low discrimination**							**300541**
					NOV	OXI	PYR	
Kocuria kristinae		55.41	0.30	URE 10	1	99	93	
Staphylococcus cohnii subsp. cohnii		39.09	0.36	SUC 99	99	1	1	
Kocuria varians		2.95	0.11		1	13	10	
300542	**Low discrimination**							**300542**
					OXI	NOV		
Staphylococcus warneri		29.52	0.37	SUC 95	1	1		
Macrococcus caseolyticus		28.01	0.18		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		22.52	0.14		1	99		
300550	**Low discrimination**							**300550**
					OXI			
Macrococcus equipersicus		34.56	0.45	VPT 1	99			
Macrococcus bovicus		21.66	0.39		99			
Staphylococcus saprophyticus subsp. saprophyticus		15.63	0.27		1			
300551	**Low discrimination**							**300551**
					PYR			
Staphylococcus cohnii subsp. cohnii		71.61	0.39	URE 1	1			
Staphylococcus saprophyticus subsp. bovis		12.43	0.18	GAL 10	99			
Staphylococcus cohnii subsp. urealyticum		7.10	0.19		53			
300552	**Acceptable identification**							**300552**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		92.46	0.42	GAL 8	99	7		
Staphylococcus warneri		3.66	0.34	SUC 91	1	8		
Staphylococcus saprophyticus subsp. bovis		1.83	0.18		99	99		
300553	**Low discrimination**							**300553**
					NOV			
Staphylococcus cohnii subsp. urealyticum		83.79	0.39	GLR 99	99			
Staphylococcus saprophyticus subsp. saprophyticus		7.39	0.01		99			
Staphylococcus caprae		5.20	-.02		1			
300600	**Low discrimination**							**300600**
					OXI	PYR		
Staphylococcus warneri		67.98	0.57	MLT 95	1	8		
Staphylococcus petrasii subsp. pragensis		14.66	0.18		1	99		
Kocuria kristinae		5.56	0.20		99	93		
300601	**Very good identification**							**300601**
					OXI	PYR		
Kocuria kristinae		99.53	0.61	URE 10	99	93		
Staphylococcus warneri		0.25	0.22	MLT 90	1	8		
Staphylococcus petrasii subsp. petrasii		0.09	-.02		1	99		
300602	**Low discrimination**							**300602**
					NOV			
Staphylococcus warneri		66.75	0.47	MLT 95	1			
Staphylococcus saprophyticus subsp. saprophyticus		18.89	0.15		99			
Staphylococcus hominis subsp. hominis		4.40	0.17		1			
300603	**Good identification**							**300603**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	PYR		
Kocuria kristinae		99.67	0.48	URE 10	99	93		
Staphylococcus warneri		0.34	0.12	MLT 90	1	8		
300610	**Low discrimination**							**300610**
					NOV			
Staphylococcus warneri		48.65	0.54	MLT 95	1			
Staphylococcus saprophyticus subsp. saprophyticus		26.84	0.28		99			
Staphylococcus pasteuri		11.88	0.27		1			
300611	**Acceptable identification**							**300611**
					OXI	NOV	PYR	
Kocuria kristinae		92.59	0.32	URE 10	99	1	93	
Staphylococcus warneri		4.06	0.19	MAN 4	1	1	8	
Staphylococcus cohnii subsp. cohnii		1.74	0.07	MLT 90	1	99	1	
300612	**Low discrimination**							**300612**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		87.83	0.43	MLT 99	99			
Staphylococcus warneri		9.37	0.44		1			
Staphylococcus pasteuri		2.80	0.19		1			
300640	**Low discrimination**							**300640**
					PYR			
Staphylococcus petrasii subsp. pragensis		47.09	0.59	URE 1	99			
Staphylococcus warneri		41.91	0.83		8			
Staphylococcus hominis subsp. hominis		4.09	0.57		1			
300641	**Very good identification**							**300641**
					OXI	PYR		
Kocuria kristinae		98.01	0.80	URE 10	99	93		
Staphylococcus petrasii subsp. petrasii		0.97	0.39		1	99		
Staphylococcus warneri		0.52	0.48		1	8		
300642	**Low discrimination**							**300642**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		50.37	0.56	MAN 96	99			
Staphylococcus warneri		34.16	0.73		1			
Staphylococcus hominis subsp. hominis		11.73	0.58		1			
300643	**Very good identification**							**300643**
					OXI	PYR		
Kocuria kristinae		96.01	0.68	URE 10	99	93		
Staphylococcus petrasii subsp. petrasii		2.05	0.34		1	99		
Staphylococcus warneri		0.68	0.39		1	8		
300650	**Low discrimination**							**300650**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		71.20	0.69		99			
Staphylococcus warneri		24.77	0.80		1			
Staphylococcus pasteuri		1.96	0.43		1			
300651	**Low discrimination**							**300651**
					OXI	NOV	PYR	
Kocuria kristinae		78.42	0.52	URE 10	99	1	93	
Staphylococcus saprophyticus subsp. saprophyticus		10.32	0.28	MAN 4	1	99	7	
Staphylococcus warneri		7.25	0.46		1	1	8	
300652	**Very good identification**							**300652**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		97.65	0.85		99			
Staphylococcus warneri		2.00	0.71		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus pasteuri		0.19	0.35		1			
300653	**Low discrimination**							**300653**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		70.64	0.43	MNS 1	1	99	7	
Kocuria kristinae		23.69	0.40		99	1	93	
Staphylococcus warneri		2.92	0.36		1	1	8	
300661	**Good identification**							**300661**
					OXI			
Kocuria kristinae		99.02	0.39	URE 10	99			
Staphylococcus petrasii subsp. petrasii		0.98	-.02	XYL 1	1			
300663	**Good identification**							**300663**
Kocuria kristinae		100.0	0.27	URE 10				
300670	**Low discrimination**							**300670**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		72.71	0.28	XYL 1	99			
Staphylococcus warneri		25.29	0.39		1			
Staphylococcus pasteuri		2.00	0.02		1			
300672	**Good identification**							**300672**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		97.81	0.43	XYL 1	99			
Staphylococcus warneri		2.00	0.29		1			
Staphylococcus pasteuri		0.19	-.06		1			
300700	**Low discrimination**							**300700**
					OXI	PYR		
Staphylococcus warneri		67.78	0.47	MLT 95	1	8		
Kocuria varians		20.02	0.29		13	10		
Kocuria kristinae		5.85	0.10		99	93		
300701	**Good identification**							**300701**
					OXI	PYR		
Kocuria kristinae		99.11	0.51	URE 10	99	93		
Kocuria varians		0.30	0.07	MLT 90	13	10		
Staphylococcus warneri		0.24	0.12		1	8		
300702	**Low discrimination**							**300702**
Staphylococcus warneri		60.01	0.37	MLT 95				
Kocuria varians		15.00	0.17					
Staphylococcus hominis subsp. hominis		13.57	0.18					
300703	**Good identification**							**300703**
Kocuria kristinae		100.0	0.39	URE 10				
300710	**Low discrimination**							**300710**
					OXI	NOV		
Staphylococcus warneri		65.78	0.44	MLT 95	1	1		
Macrococcus bovicus		14.54	0.19		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		9.99	0.06		1	99		
300740	**Low discrimination**							**300740**
					NOV			
Staphylococcus warneri		62.12	0.73		1			
Staphylococcus hominis subsp. hominis		20.78	0.58		1			
Staphylococcus hominis subsp. novobiosepticus		9.56	0.39		99			
300741	**Very good identification**							**300741**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	NOV	PYR	
Kocuria kristinae		96.43	0.71	URE 10	99	1	93	
Staphylococcus petrasii subsp. petrasii		1.91	0.36		1	1	99	
Staphylococcus hominis subsp. novobiosepticus		0.92	0.27		1	99	1	
300742	**Low discrimination**							**300742**
					NOV			
Staphylococcus hominis subsp. hominis		40.37	0.59	NIT 99	1			
Staphylococcus warneri		34.27	0.63		1			
Staphylococcus saprophyticus subsp. saprophyticus		13.91	0.34		99			
300743	**Good identification**							**300743**
					OXI	NOV	PYR	
Kocuria kristinae		91.63	0.58	URE 10	99	1	93	
Staphylococcus petrasii subsp. petrasii		3.91	0.30		1	1	99	
Staphylococcus hominis subsp. novobiosepticus		1.88	0.21		1	99	1	
300750	**Low discrimination**							**300750**
					NOV	PYR		
Staphylococcus warneri		48.68	0.70		1	8		
Staphylococcus saprophyticus subsp. saprophyticus		38.54	0.47		99	7		
Staphylococcus saprophyticus subsp. bovis		6.86	0.43		99	99		
300751	**Low discrimination**							**300751**
					OXI	NOV	PYR	
Kocuria kristinae		74.83	0.42	URE 10	99	1	93	
Staphylococcus saprophyticus subsp. bovis		11.31	0.31	MAN 4	1	99	99	
Staphylococcus warneri		6.56	0.35		1	1	8	
300752	**Good identification**							**300752**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		90.85	0.63	GAL 8	99			
Staphylococcus warneri		6.75	0.60		1			
Staphylococcus hominis subsp. hominis		1.64	0.42		1			
300753	**Low discrimination**							**300753**
					OXI	PYR	NOV	
Kocuria kristinae		43.77	0.30	URE 10	99	93	1	
Staphylococcus saprophyticus subsp. saprophyticus		34.05	0.22	MAN 4	1	7	99	
Staphylococcus saprophyticus subsp. bovis		6.61	0.18		1	99	99	
300760	**Low discrimination**							**300760**
					NOV			
Staphylococcus warneri		62.22	0.32	XYL 1	1			
Staphylococcus hominis subsp. hominis		20.81	0.16		1			
Staphylococcus hominis subsp. novobiosepticus		9.58	-.02		99			
300761	**Good identification**							**300761**
					OXI			
Kocuria kristinae		98.06	0.30	URE 10	99			
Staphylococcus petrasii subsp. petrasii		1.94	-.05	XYL 1	1			
300770	**Low discrimination**							**300770**
					NOV	PYR		
Staphylococcus warneri		51.75	0.29	XYL 1	1	8		
Staphylococcus saprophyticus subsp. saprophyticus		40.97	0.06		99	7		
Staphylococcus saprophyticus subsp. bovis		7.29	0.02		99	99		
301000	**Low discrimination**							**301000**
					NOV	YEP		
Dermacoccus nishinomiyaensis		58.65	0.33	VPT 5	80	1		
Micrococcus sp.		18.48	0.13	ESL 1	1	41		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Micrococcus equipersicus		12.41	0.24		99	99		
301010	**Good identification**							**301010**
Micrococcus equipersicus		97.53	0.50	VPT 1				
Micrococcus bovicus		2.47	0.16					
301040	**Low discrimination**							**301040**
					NOV	OXI	YEP	
Micrococcus equipersicus		80.25	0.33	VPT 1	99	99	99	
Staphylococcus hominis subsp. novobiosepticus		10.94	0.08	MAN 95	99	1	1	
Kocuria varians		5.49	0.11		1	13	80	
301050	**Very good identification**							**301050**
Micrococcus equipersicus		97.86	0.59	VPT 1				
Micrococcus bovicus		2.14	0.23					
301100	**Low discrimination**							**301100**
					OXI	YEP	NOV	
Kocuria varians		48.76	0.29	VPT 14	13	80	1	
Dermacoccus nishinomiyaensis		34.72	0.18	ESL 1	93	1	80	
Micrococcus equipersicus		16.53	0.16		99	99	99	
301110	**Good identification**							**301110**
Micrococcus equipersicus		97.53	0.43	VPT 1				
Micrococcus bovicus		2.47	0.08					
301150	**Good identification**							**301150**
Micrococcus equipersicus		97.86	0.51	VPT 1				
Micrococcus bovicus		2.14	0.16					
301210	**Acceptable identification**							**301210**
Micrococcus equipersicus		93.99	0.29	VPT 1				
Micrococcus bovicus		6.02	0.03	SUC 9				
301240	**Good identification**							**301240**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. novobiosepticus		98.33	0.49	ESL 8	1	99	1	
Micrococcus equipersicus		0.72	0.12	NIT 90	99	99	99	
Staphylococcus warneri		0.38	0.16		1	1	43	
301241	**Low discrimination**							**301241**
					OXI	NOV		
Kocuria kristinae		59.15	0.42	URE 10	99	1		
Staphylococcus hominis subsp. novobiosepticus		40.85	0.36	TRE 95	1	99		
301242	**Good identification**							**301242**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		99.52	0.43	ESL 8	99			
Staphylococcus hominis subsp. hominis		0.48	0.07	NIT 90	1			
301243	**Low discrimination**							**301243**
					OXI	NOV		
Staphylococcus hominis subsp. novobiosepticus		59.80	0.31	ESL 8	1	99		
Kocuria kristinae		40.20	0.29	NIT 90	99	1		
301250	**Low discrimination**							**301250**
					OXI	YEP		
Micrococcus equipersicus		87.14	0.38	VPT 1	99	99		
Staphylococcus hominis subsp. novobiosepticus		6.32	0.08	SUC 9	1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		4.81	0.11		99	99		
301340	**Very good identification**							**301340**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. novobiosepticus		99.82	0.59	ESL 8	1	99	1	
Macrococcus equipersicus		0.11	0.04	NIT 90	99	99	99	
Staphylococcus hominis subsp. hominis		0.08	0.06		1	1	49	
301341	**Low discrimination**							**301341**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		85.95	0.46	ESL 8	1	99	1	
Kocuria kristinae		13.83	0.32	NIT 90	99	1	93	
Staphylococcus epidermidis		0.22	0.01		1	1	1	
301342	**Very good identification**							**301342**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		99.62	0.53	ESL 8	99			
Staphylococcus epidermidis		0.20	0.05	NIT 90	1			
Staphylococcus hominis subsp. hominis		0.18	0.08		1			
301343	**Acceptable identification**							**301343**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		91.77	0.41	ESL 8	1	99	1	
Kocuria kristinae		6.85	0.19	NIT 90	99	1	93	
Staphylococcus epidermidis		1.38	0.11		1	1	1	
301350	**Low discrimination**							**301350**
					OXI	YEP		
Macrococcus equipersicus		63.98	0.31	VPT 1	99	99		
Staphylococcus hominis subsp. novobiosepticus		32.49	0.17	SUC 9	1	1		
Macrococcus bovicus		3.53	0.03		99	99		
301410	**Acceptable identification**							**301410**
Macrococcus equipersicus		94.53	0.50	VPT 1				
Macrococcus bovicus		3.59	0.19					
Macrococcus carouselicus		1.48	0.08					
301440	**Low discrimination**							**301440**
					OXI	NOV	YEP	
Macrococcus equipersicus		88.86	0.33	VPT 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		6.57	0.14	MAN 95	1	99	12	
Staphylococcus warneri		4.57	0.16		1	1	43	
301441	**Acceptable identification**							**301441**
					OXI	NOV		
Kocuria kristinae		92.12	0.27	URE 10	99	1		
Staphylococcus cohnii subsp. cohnii		7.88	0.15	SUC 99	1	99		
301450	**Good identification**							**301450**
					OXI			
Macrococcus equipersicus		94.86	0.59	VPT 1	99			
Macrococcus bovicus		3.11	0.27		99			
Staphylococcus saprophyticus subsp. saprophyticus		1.01	0.07		1			
301453	**Good identification**							**301453**
					NOV			
Staphylococcus cohnii subsp. urealyticum		97.22	0.39	GLR 99	99			
Aerococcus viridans		2.78	0.18		1			
301510	**Acceptable identification**							**301510**
Macrococcus equipersicus		94.91	0.43	VPT 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Macrococcus bovicus		3.60	0.12					
Macrococcus carouselicus		1.49	0.00					
301550	**Good identification**							**301550**
					OXI			
Macrococcus equipersicus		95.21	0.51	VPT 1	99			
Macrococcus bovicus		3.12	0.19		99			
Staphylococcus kloosii		1.67	0.33		1			
301552	**Low discrimination**							**301552**
					OXI			
Staphylococcus kloosii		45.96	0.33	VPT 10	1			
Staphylococcus saprophyticus subsp. saprophyticus		29.54	0.01	PHS 99	1			
Macrococcus equipersicus		24.50	0.10		99			
301553	**Acceptable identification**							**301553**
					NOV			
Staphylococcus cohnii subsp. urealyticum		93.74	0.31	GLR 99	99			
Aerococcus viridans		6.26	0.18		1			
301601	**Good identification**							**301601**
Kocuria kristinae		100.0	0.48	URE 10				
301603	**Good identification**							**301603**
Kocuria kristinae		100.0	0.36	URE 10				
301610	**Low discrimination**							**301610**
Macrococcus equipersicus		67.98	0.29	VPT 1				
Macrococcus carouselicus		21.86	0.14	SUC 9				
Macrococcus bovicus		6.53	0.07					
301641	**Very good identification**							**301641**
					OXI	NOV	PYR	
Kocuria kristinae		99.92	0.68	URE 10	99	1	93	
Staphylococcus petrasii subsp. petrasii		0.04	-.02		1	1	99	
Staphylococcus hominis subsp. novobiosepticus		0.04	-.05		1	99	1	
301643	**Very good identification**							**301643**
Kocuria kristinae		100.0	0.56	URE 10				
301650	**Low discrimination**							**301650**
					OXI	NOV		
Staphylococcus saprophyticus subsp. saprophyticus		41.76	0.28	ESL 1	1	99		
Macrococcus equipersicus		38.57	0.38		99	99		
Staphylococcus warneri		14.53	0.39		1	1		
301651	**Good identification**							**301651**
Kocuria kristinae		100.0	0.40	URE 10				
301652	**Good identification**							**301652**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		97.81	0.43	ESL 1	99			
Staphylococcus warneri		2.00	0.29		1			
Staphylococcus pasteuri		0.19	-.06		1			
301653	**Low discrimination**							**301653**
					OXI	NOV		
Kocuria kristinae		84.50	0.27	URE 10	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		10.18	0.02	MAN 4	1	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		2.76	0.07		1	99	
301661	**Good identification**						**301661**
Kocuria kristinae		100.0	0.27	URE 10			
301701	**Acceptable identification**						**301701**
					OXI		
Kocuria kristinae		93.88	0.39	URE 10	99		
Rothia mucilaginosa		6.12	0.10	MLT 90	1		
301741	**Very good identification**						**301741**
					OXI	NOV	
Kocuria kristinae		97.94	0.58	URE 10	99	1	
Rothia mucilaginosa		1.66	0.18		1	1	
Staphylococcus hominis subsp. novobiosepticus		0.32	0.05		1	99	
301743	**Good identification**						**301743**
					OXI	NOV	
Kocuria kristinae		99.29	0.46	URE 10	99	1	
Staphylococcus hominis subsp. novobiosepticus		0.71	-.01		1	99	
301750	**Low discrimination**						**301750**
					OXI	NOV	
Macrococcus equipersicus		61.76	0.31	VPT 1	99	99	
Staphylococcus warneri		17.14	0.29	SUC 9	1	1	
Staphylococcus saprophyticus subsp. saprophyticus		13.57	0.06		1	99	
301751	**Good identification**						**301751**
Kocuria kristinae		100.0	0.30	URE 10			
302000	**Low discrimination**						**302000**
					OXI		
Kocuria varians		57.81	0.80	VPT 14	13		
Dermacoccus nishinomiyaensis		37.34	0.69		93		
Kocuria rosea		3.67	0.47		35		
302001	**Low discrimination**						**302001**
					OXI		
Kocuria varians		86.40	0.58	VPT 14	13		
Dermacoccus nishinomiyaensis		6.48	0.28	MNS 8	93		
Kocuria rosea		3.32	0.20		35		
302002	**Very good identification**						**302002**
					OXI		
Kocuria varians		97.21	0.69	VPT 14	13		
Dermacoccus nishinomiyaensis		2.25	0.28		93		
Kocuria rosea		0.22	0.05		35		
302003	**Good identification**						**302003**
Kocuria varians		100.0	0.47	VPT 14			
302010	**Low discrimination**						**302010**
					OXI		
Kocuria varians		39.31	0.39	VPT 14	13		
Kocuria rosea		27.42	0.27	MAN 1	35		
Dermacoccus nishinomiyaensis		25.39	0.28		93		
302011	**Good identification**						**302011**
Staphylococcus capitis subsp. capitis		97.14	0.45	URE 1			
Kocuria varians		2.01	0.17				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		0.85	0.01				
302012	**Good identification**						**302012**
Kocuria varians		100.0	0.28	VPT 14			
302020	**Good identification**						**302020**
					OXI		
Kocuria varians		91.20	0.71	VPT 14	13		
Kocuria rosea		7.06	0.39		35		
Dermacoccus nishinomiyaensis		1.69	0.28		93		
302021	**Good identification**						**302021**
Kocuria varians		95.53	0.49	VPT 14			
Kocuria rosea		4.47	0.13	MNS 8			
302022	**Very good identification**						**302022**
Kocuria varians		99.72	0.60	VPT 14			
Kocuria rosea		0.28	-.02				
302023	**Good identification**						**302023**
Kocuria varians		100.0	0.38	VPT 14			
302030	**Low discrimination**						**302030**
Kocuria varians		54.03	0.30	VPT 14			
Kocuria rosea		45.98	0.19	MAN 1			
302040	**Low discrimination**						**302040**
					NOV		
Kocuria varians		63.91	0.67	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		27.26	0.49		99		
Staphylococcus hominis subsp. hominis		5.76	0.47		1		
302041	**Low discrimination**						**302041**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		53.35	0.37	SUC 99	99		
Kocuria varians		43.50	0.45		1		
Staphylococcus epidermidis		2.23	0.16		1		
302042	**Low discrimination**						**302042**
					NOV		
Kocuria varians		45.04	0.55	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		36.68	0.44		99		
Staphylococcus hominis subsp. hominis		16.89	0.48		1		
302043	**Low discrimination**						**302043**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		56.04	0.31	SUC 99	99		
Kocuria varians		23.94	0.33		1		
Staphylococcus epidermidis		13.80	0.27		1		
302050	**Low discrimination**						**302050**
					NOV	OXI	
Staphylococcus hominis subsp. hominis		26.11	0.30	SUC 99	1	1	
Macrococcus bovicus		24.35	0.23	MAN 13	99	99	
Kocuria varians		19.59	0.26		1	13	
302052	**Low discrimination**						**302052**
					NOV		
Staphylococcus hominis subsp. hominis		65.02	0.31	SUC 99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		11.72	0.14	MAN 13	1			
Staphylococcus hominis subsp. novobiosepticus		9.55	0.03		99			
302053	**Good identification**							**302053**
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	ARG 99				
302060	**Very good identification**							**302060**
					NOV			
Kocuria varians		98.19	0.57	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		1.20	0.08		99			
Kocuria rosea		0.35	-.02		1			
302061	**Good identification**							**302061**
					NOV			
Kocuria varians		96.59	0.35	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		3.41	-.04	MNS 8	99			
302062	**Good identification**							**302062**
					NOV			
Kocuria varians		96.69	0.46	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		2.26	0.03		99			
Staphylococcus hominis subsp. hominis		1.04	0.07		1			
302100	**Good identification**							**302100**
					OXI	NOV		
Kocuria varians		92.20	0.84	VPT 14	13	1		
Dermacoccus nishinomiyaensis		7.56	0.54		93	80		
Staphylococcus hominis subsp. novobiosepticus		0.17	0.18		1	99		
302101	**Very good identification**							**302101**
					OXI	NOV		
Kocuria varians		98.53	0.62	VPT 14	13	1		
Dermacoccus nishinomiyaensis		0.94	0.13	MNS 8	93	80		
Staphylococcus hominis subsp. novobiosepticus		0.54	0.05		1	99		
302102	**Very good identification**							**302102**
					OXI	NOV		
Kocuria varians		99.29	0.73	VPT 14	13	1		
Staphylococcus hominis subsp. novobiosepticus		0.36	0.12		1	99		
Dermacoccus nishinomiyaensis		0.29	0.13		93	80		
302103	**Good identification**							**302103**
					COA	NOV		
Kocuria varians		96.98	0.51	VPT 14	1	1		
Staphylococcus schleiferi subsp. coagulans		1.55	0.12	MNS 8	99	1		
Staphylococcus hominis subsp. novobiosepticus		1.01	0.00		1	99		
302110	**Low discrimination**							**302110**
					OXI	YEP	NOV	
Kocuria varians		89.65	0.43	VPT 14	13	80	1	
Dermacoccus nishinomiyaensis		7.35	0.13	MAN 1	93	1	80	
Micrococcus bovicus		3.00	0.08		99	99	99	
302112	**Good identification**							**302112**
Kocuria varians		100.0	0.31	VPT 14				
302120	**Very good identification**							**302120**
					OXI			
Kocuria varians		99.71	0.74	VPT 14	13			
Dermacoccus nishinomiyaensis		0.24	0.13		93			
Kocuria rosea		0.05	-.02		35			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
302121	**Good identification**						**302121**
Kocuria varians		100.0	0.53	VPT 14			
302122	**Very good identification**						**302122**
Kocuria varians		100.0	0.63	VPT 14			
302123	**Good identification**						**302123**
Kocuria varians		100.0	0.41	VPT 14			
302130	**Good identification**						**302130**
Kocuria varians		100.0	0.33	VPT 14			
302140	**Low discrimination**						**302140**
					NOV		
Kocuria varians		51.58	0.70	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		44.01	0.59		99		
Staphylococcus hominis subsp. hominis		3.36	0.47		1		
302141	**Low discrimination**						**302141**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		66.98	0.47	SUC 99	99		
Kocuria varians		27.31	0.48		1		
Staphylococcus epidermidis		5.30	0.32		1		
302142	**Low discrimination**						**302142**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		53.61	0.53	SUC 99	99		
Kocuria varians		32.92	0.59		1		
Staphylococcus hominis subsp. hominis		8.92	0.49		1		
302143	**Low discrimination**						**302143**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		59.10	0.41	SUC 99	99		
Staphylococcus epidermidis		27.48	0.42		1		
Kocuria varians		12.62	0.37		1		
302150	**Low discrimination**						**302150**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		32.02	0.31	TRE 99	99	99	
Kocuria varians		20.60	0.29		1	10	
Staphylococcus hominis subsp. hominis		19.83	0.30		1	1	
302152	**Low discrimination**						**302152**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		50.00	0.32	SUC 99	1	1	
Staphylococcus hominis subsp. novobiosepticus		20.33	0.12	MAN 13	1	99	
Staphylococcus saprophyticus subsp. bovis		17.20	0.18		99	99	
302160	**Very good identification**						**302160**
					NOV		
Kocuria varians		97.43	0.61	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		2.39	0.18		99		
Staphylococcus hominis subsp. hominis		0.18	0.06		1		
302161	**Acceptable identification**						**302161**
					NOV		
Kocuria varians		93.41	0.39	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		6.59	0.05	MNS 8	99		
302162	**Acceptable identification**						**302162**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Kocuria varians		94.82	0.50	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		4.44	0.12		99		
Staphylococcus hominis subsp. hominis		0.74	0.08		1		
302163	**Low discrimination**						**302163**
					NOV		
Kocuria varians		83.53	0.28	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		11.24	0.00	MNS 8	99		
Staphylococcus epidermidis		5.23	0.01		1		
302200	**Low discrimination**						**302200**
					OXI	NOV	
Kocuria varians		45.23	0.62	VPT 14	13	1	
Staphylococcus hominis subsp. novobiosepticus		31.06	0.49	SUC 12	1	99	
Dermacoccus nishinomiyaensis		13.68	0.44		93	80	
302201	**Low discrimination**						**302201**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		52.65	0.37	MLT 99	99		
Kocuria varians		26.67	0.41		1		
Staphylococcus capitis subsp. capitis		13.36	0.28		1		
302202	**Low discrimination**						**302202**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		44.06	0.44	MLT 99	99		
Kocuria varians		33.61	0.51		1		
Staphylococcus hominis subsp. hominis		20.29	0.48		1		
302203	**Low discrimination**						**302203**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		60.82	0.31	MLT 99	99		
Kocuria varians		16.14	0.29		1		
Staphylococcus epidermidis		14.97	0.27		1		
302210	**Low discrimination**						**302210**
					NOV		
Staphylococcus hominis subsp. hominis		34.65	0.30	MAN 13	1		
Kocuria varians		16.15	0.21	MLT 99	1		
Staphylococcus hominis subsp. novobiosepticus		11.09	0.08		99		
302211	**Good identification**						**302211**
					NOV		
Staphylococcus capitis subsp. capitis		96.02	0.51	URE 1	1		
Staphylococcus capitis subsp. ureolyticus		3.69	0.13		1		
Staphylococcus hominis subsp. novobiosepticus		0.29	-.04		99		
302212	**Low discrimination**						**302212**
					NOV		
Staphylococcus hominis subsp. hominis		72.64	0.31	MAN 13	1		
Staphylococcus hominis subsp. novobiosepticus		10.66	0.03	MLT 99	99		
Kocuria varians		8.14	0.10		1		
302213	**Acceptable identification**						**302213**
Staphylococcus capitis subsp. ureolyticus		92.57	0.31	ARG 99			
Staphylococcus capitis subsp. capitis		6.70	0.16	MLT 96			
Staphylococcus simulans		0.72	0.00				
302220	**Good identification**						**302220**
					NOV		
Kocuria varians		91.27	0.53	VPT 14	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Kocuria rosea		5.75	0.19	SUC 12	1	
Staphylococcus hominis subsp. novobiosepticus		1.80	0.08		99	
302221	**Acceptable identification**					**302221**
					NOV	
Kocuria varians		91.33	0.31	VPT 14	1	
Staphylococcus hominis subsp. novobiosepticus		5.18	-.04	SUC 12	99	
Kocuria rosea		3.49	-.07	MNS 8	1	
302222	**Acceptable identification**					**302222**
					NOV	
Kocuria varians		94.78	0.42	VPT 14	1	
Staphylococcus hominis subsp. novobiosepticus		3.57	0.03	SUC 12	99	
Staphylococcus hominis subsp. hominis		1.65	0.07		1	
302240	**Low discrimination**					**302240**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		81.84	0.90		99	
Staphylococcus hominis subsp. hominis		17.29	0.88		1	
Staphylococcus epidermidis		0.46	0.52		1	
302241	**Good identification**					**302241**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		93.73	0.78		99	
Staphylococcus epidermidis		3.92	0.57		1	
Staphylococcus hominis subsp. hominis		1.62	0.53		1	
302242	**Low discrimination**					**302242**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		66.83	0.85		99	
Staphylococcus hominis subsp. hominis		30.77	0.89		1	
Staphylococcus epidermidis		2.21	0.62		1	
302243	**Low discrimination**					**302243**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		76.53	0.72		99	
Staphylococcus epidermidis		18.84	0.68		1	
Staphylococcus hominis subsp. hominis		2.88	0.54		1	
302250	**Low discrimination**					**302250**
					NOV	
Staphylococcus hominis subsp. hominis		72.18	0.71	MAN 13	1	
Staphylococcus hominis subsp. novobiosepticus		23.10	0.49		99	
Staphylococcus warneri		2.59	0.46		1	
302251	**Low discrimination**					**302251**
					NOV	
Staphylococcus capitis subsp. ureolyticus		68.60	0.41	ARG 99	1	
Staphylococcus hominis subsp. novobiosepticus		22.06	0.37	LAC 88	99	
Staphylococcus hominis subsp. hominis		5.63	0.36		1	
302252	**Low discrimination**					**302252**
					NOV	
Staphylococcus hominis subsp. hominis		84.38	0.72	MAN 13	1	
Staphylococcus hominis subsp. novobiosepticus		12.39	0.44		99	
Staphylococcus capitis subsp. ureolyticus		1.33	0.18		1	
302253	**Very good identification**					**302253**
					NOV	
Staphylococcus capitis subsp. ureolyticus		95.86	0.59	ARG 99	1	
Staphylococcus hominis subsp. novobiosepticus		2.26	0.31		99	
Staphylococcus hominis subsp. hominis		1.26	0.37		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
302260	**Low discrimination**					**302260**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		75.24	0.49	XYL 1	99	
Staphylococcus hominis subsp. hominis		15.89	0.47		1	
Kocuria varians		8.45	0.39		1	
302261	**Acceptable identification**					**302261**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		91.06	0.37	XYL 1	99	
Staphylococcus epidermidis		3.81	0.16		1	
Kocuria varians		3.56	0.18		1	
302262	**Low discrimination**					**302262**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		64.41	0.44	XYL 1	99	
Staphylococcus hominis subsp. hominis		29.66	0.48		1	
Kocuria varians		3.79	0.28		1	
302263	**Low discrimination**					**302263**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		75.64	0.31	XYL 1	99	
Staphylococcus epidermidis		18.62	0.27		1	
Staphylococcus hominis subsp. hominis		2.84	0.13		1	
302270	**Low discrimination**					**302270**
					NOV	
Staphylococcus hominis subsp. hominis		75.76	0.30	MAN 13	1	
Staphylococcus hominis subsp. novobiosepticus		24.25	0.08	XYL 1	99	
302272	**Low discrimination**					**302272**
					NOV	
Staphylococcus hominis subsp. hominis		87.20	0.31	MAN 13	1	
Staphylococcus hominis subsp. novobiosepticus		12.80	0.03	XYL 1	99	
302300	**Low discrimination**					**302300**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		54.23	0.59	MLT 99	99	
Kocuria varians		39.49	0.66		1	
Staphylococcus hominis subsp. hominis		4.14	0.47		1	
302301	**Low discrimination**					**302301**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		74.29	0.47	MLT 99	99	
Kocuria varians		18.82	0.44		1	
Staphylococcus epidermidis		5.87	0.32		1	
302302	**Low discrimination**					**302302**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		62.10	0.53	MLT 99	99	
Kocuria varians		23.69	0.55		1	
Staphylococcus hominis subsp. hominis		10.33	0.49		1	
302303	**Low discrimination**					**302303**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		61.88	0.41	MLT 99	99	
Staphylococcus epidermidis		28.77	0.42		1	
Kocuria varians		8.21	0.33		1	
302310	**Low discrimination**					**302310**
					NOV	
Staphylococcus hominis subsp. hominis		37.72	0.30	MAN 13	1	
Staphylococcus hominis subsp. novobiosepticus		33.43	0.18	MLT 99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		24.34	0.25		1		
302312	**Low discrimination**						**302312**
					NOV		
Staphylococcus hominis subsp. hominis		64.04	0.32	MAN 13	1		
Staphylococcus hominis subsp. novobiosepticus		26.03	0.12	MLT 99	99		
Kocuria varians		9.93	0.14		1		
302320	**Very good identification**						**302320**
					NOV		
Kocuria varians		95.92	0.57	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		3.79	0.18	SUC 12	99		
Staphylococcus hominis subsp. hominis		0.29	0.06		1		
302321	**Low discrimination**						**302321**
					NOV		
Kocuria varians		89.81	0.35	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		10.19	0.05	SUC 12	99		
302322	**Acceptable identification**						**302322**
					NOV		
Kocuria varians		91.92	0.45	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		6.93	0.12	SUC 12	99		
Staphylococcus hominis subsp. hominis		1.15	0.08		1		
302340	**Good identification**						**302340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		91.85	1.00		99		
Staphylococcus hominis subsp. hominis		7.01	0.88		1		
Staphylococcus epidermidis		0.98	0.67		1		
302341	**Good identification**						**302341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		92.03	0.88		99		
Staphylococcus epidermidis		7.28	0.73		1		
Staphylococcus hominis subsp. hominis		0.57	0.54		1		
302342	**Low discrimination**						**302342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		81.30	0.95		99		
Staphylococcus hominis subsp. hominis		13.52	0.90		1		
Staphylococcus epidermidis		5.09	0.78		1		
302343	**Low discrimination**						**302343**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		67.57	0.82		99		
Staphylococcus epidermidis		31.42	0.83		1		
Staphylococcus hominis subsp. hominis		0.92	0.55		1		
302350	**Low discrimination**						**302350**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		50.64	0.71	MAN 13	1	1	
Staphylococcus hominis subsp. novobiosepticus		44.88	0.59		1	99	
Staphylococcus saprophyticus subsp. bovis		3.30	0.43		99	99	
302351	**Low discrimination**						**302351**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		79.52	0.47	MAN 1	99		
Staphylococcus hominis subsp. hominis		7.32	0.37		1		
Staphylococcus epidermidis		6.29	0.32		1		
302352	**Low discrimination**						**302352**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		69.00	0.73	MAN 13	1		
Staphylococcus hominis subsp. novobiosepticus		28.05	0.53		99		
Staphylococcus epidermidis		1.76	0.36		1		
302353	**Low discrimination**						**302353**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		54.14	0.41	MAN 1	99		
Staphylococcus epidermidis		25.17	0.42		1		
Staphylococcus hominis subsp. hominis		10.87	0.38		1		
302360	**Low discrimination**						**302360**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		87.48	0.59	XYL 1	99		
Staphylococcus hominis subsp. hominis		6.67	0.47		1		
Kocuria varians		4.91	0.43		1		
302361	**Acceptable identification**						**302361**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		90.51	0.47	XYL 1	99		
Staphylococcus epidermidis		7.16	0.32		1		
Kocuria varians		1.77	0.21		1		
302362	**Low discrimination**						**302362**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		79.47	0.53	XYL 1	99		
Staphylococcus hominis subsp. hominis		13.22	0.49		1		
Staphylococcus epidermidis		4.97	0.36		1		
302363	**Low discrimination**						**302363**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		67.17	0.41	XYL 1	99		
Staphylococcus epidermidis		31.23	0.42		1		
Staphylococcus hominis subsp. hominis		0.91	0.14		1		
302370	**Low discrimination**						**302370**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		49.97	0.30	MAN 13	1	1	
Staphylococcus hominis subsp. novobiosepticus		44.29	0.18	XYL 1	1	99	
Staphylococcus saprophyticus subsp. bovis		3.26	0.02		99	99	
302372	**Low discrimination**						**302372**
					NOV		
Staphylococcus hominis subsp. hominis		71.10	0.32	MAN 13	1		
Staphylococcus hominis subsp. novobiosepticus		28.90	0.12	XYL 1	99		
302400	**Low discrimination**						**302400**
					OXI		
Kocuria varians		85.02	0.57	VPT 14	13		
Dermacoccus nishinomiyaensis		7.37	0.28	TRE 7	93		
Staphylococcus auricularis		3.79	0.28		1		
302401	**Low discrimination**						**302401**
					NOV		
Kocuria varians		76.46	0.35	VPT 14	1		
Staphylococcus cohnii subsp. cohnii		14.62	0.23	TRE 7	99		
Staphylococcus auricularis		4.35	0.08	MNS 8	1		
302402	**Acceptable identification**						**302402**
Kocuria varians		93.94	0.46	VPT 14			
Staphylococcus hominis subsp. hominis		3.29	0.17	TRE 7			
Staphylococcus auricularis		1.65	0.08				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
302420	**Good identification**							**302420**
Kocuria varians		98.97	0.48	VPT 14				
Kocuria rosea		1.03	-.02	TRE 7				
302422	**Good identification**							**302422**
Kocuria varians		100.0	0.36	VPT 14				
302440	**Low discrimination**							**302440**
Staphylococcus hominis subsp. hominis		56.59	0.57	SUC 99				
Kocuria varians		14.93	0.44					
Staphylococcus warneri		13.09	0.49					
302441	**Low discrimination**							**302441**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. cohnii		31.18	0.33	URE 1	1	99	1	
Staphylococcus petrasii subsp. petrasii		17.46	0.18	NIT 7	1	1	99	
Kocuria kristinae		14.73	0.16		99	1	93	
302442	**Low discrimination**							**302442**
Staphylococcus hominis subsp. hominis		83.17	0.58	SUC 99				
Staphylococcus warneri		5.46	0.40					
Kocuria varians		5.27	0.32					
302450	**Low discrimination**							**302450**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		80.17	0.59		99	99		
Staphylococcus warneri		6.57	0.46		1	8		
Staphylococcus hominis subsp. hominis		5.86	0.40		1	1		
302451	**Low discrimination**							**302451**
					PYR			
Staphylococcus saprophyticus subsp. bovis		84.21	0.47		99			
Staphylococcus cohnii subsp. cohnii		13.28	0.35		1			
Staphylococcus cohnii subsp. urealyticum		0.80	0.11		53			
302452	**Low discrimination**							**302452**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		58.79	0.47		99	99		
Staphylococcus hominis subsp. hominis		20.19	0.41		1	1		
Staphylococcus saprophyticus subsp. saprophyticus		13.91	0.23		99	7		
302453	**Low discrimination**							**302453**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		69.42	0.34		99	99		
Staphylococcus cohnii subsp. urealyticum		26.69	0.31		99	53		
Staphylococcus hominis subsp. hominis		1.95	0.06		1	1		
302460	**Acceptable identification**							**302460**
Kocuria varians		90.17	0.34	VPT 14				
Staphylococcus hominis subsp. hominis		9.83	0.16	TRE 7				
302500	**Very good identification**							**302500**
					OXI	YEP	NOV	
Kocuria varians		95.93	0.61	VPT 14	13	80	1	
Macrococcus caseolyticus		1.91	0.14	TRE 7	99	99	99	
Dermacoccus nishinomiyaensis		1.06	0.13		93	1	80	
302501	**Good identification**							**302501**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		96.58	0.39	VPT 14				
Staphylococcus schleiferi subsp. schleiferi		1.80	-.02	TRE 7				
Rothia mucilaginosa		1.63	-.09	MNS 8				
302502	**Low discrimination**							**302502**
					NOV	OXI		
Kocuria varians		88.40	0.49	VPT 14	1	13		
Macrococcus caseolyticus		9.37	0.18	TRE 7	99	99		
Staphylococcus hominis subsp. hominis		2.23	0.18		1	1		
302503	**Good identification**							**302503**
Kocuria varians		100.0	0.28	VPT 14				
302510	**Low discrimination**							**302510**
					NOV	PYR	OXI	
Staphylococcus saprophyticus subsp. bovis		69.80	0.31	MLT 99	99	99	1	
Kocuria varians		15.40	0.20		1	10	13	
Macrococcus bovicus		10.25	0.12		99	1	99	
302520	**Good identification**							**302520**
Kocuria varians		100.0	0.51	VPT 14				
302521	**Good identification**							**302521**
Kocuria varians		100.0	0.30	VPT 14				
302522	**Good identification**							**302522**
Kocuria varians		100.0	0.40	VPT 14				
302540	**Low discrimination**							**302540**
					NOV	OXI		
Macrococcus caseolyticus		67.63	0.55	URE 1	99	99		
Staphylococcus hominis subsp. hominis		20.59	0.58		1	1		
Kocuria varians		7.52	0.47		1	13		
302542	**Low discrimination**							**302542**
					NOV	OXI		
Macrococcus caseolyticus		78.60	0.59	URE 1	99	99		
Staphylococcus hominis subsp. hominis		18.73	0.59		1	1		
Kocuria varians		1.64	0.36		1	13		
302550	**Very good identification**							**302550**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		96.51	0.72		99	99		
Staphylococcus hominis subsp. hominis		1.91	0.41		1	1		
Staphylococcus warneri		0.63	0.36		1	8		
302551	**Very good identification**							**302551**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		99.11	0.59		99	99		
Staphylococcus cohnii subsp. cohnii		0.43	0.15		99	1		
Staphylococcus caprae		0.17	-.09		1	75		
302552	**Low discrimination**							**302552**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		87.90	0.59		1	99	99	
Staphylococcus hominis subsp. hominis		8.18	0.42		1	1	1	
Macrococcus caseolyticus		2.32	0.18		99	99	80	
302553	**Low discrimination**							**302553**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus saprophyticus subsp. bovis		84.41	0.47		99		
Staphylococcus caprae		11.00	0.18		1		
Staphylococcus cohnii subsp. urealyticum		3.48	0.24		99		
302560	**Low discrimination**						**302560**
					NOV	OXI	
Kocuria varians		74.78	0.38	VPT 14	1	13	
Macrococcus caseolyticus		19.33	0.14	TRE 7	99	99	
Staphylococcus hominis subsp. hominis		5.89	0.16		1	1	
302570	**Good identification**						**302570**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	XYL 1			
302600	**Low discrimination**						**302600**
Staphylococcus hominis subsp. hominis		65.59	0.57	MLT 99			
Staphylococcus warneri		15.17	0.49				
Kocuria varians		10.75	0.39				
302601	**Low discrimination**						**302601**
					OXI	PYR	
Kocuria kristinae		78.48	0.38	URE 10	99	93	
Staphylococcus petrasii subsp. petrasii		8.46	0.18	NIT 7	1	99	
Staphylococcus hominis subsp. hominis		4.87	0.22	MLT 90	1	1	
302602	**Low discrimination**						**302602**
Staphylococcus hominis subsp. hominis		88.24	0.58	MLT 99			
Staphylococcus warneri		5.80	0.40				
Kocuria varians		3.48	0.28				
302610	**Low discrimination**						**302610**
Staphylococcus pasteuri		36.29	0.35	GLR 99			
Staphylococcus warneri		26.02	0.46	MLT 86			
Staphylococcus hominis subsp. hominis		23.21	0.40				
302612	**Low discrimination**						**302612**
Staphylococcus hominis subsp. hominis		46.72	0.41	MAN 13			
Staphylococcus pasteuri		25.42	0.27	MLT 99			
Staphylococcus warneri		14.87	0.37				
302613	**Low discrimination**						**302613**
					NOV	PYR	
Staphylococcus simulans		77.02	0.27	VPT 4	1	70	
Staphylococcus saprophyticus subsp. bovis		7.53	0.06	ARG 92	99	99	
Staphylococcus capitis subsp. ureolyticus		5.23	-.11	BGA 90	1	1	
302620	**Low discrimination**						**302620**
Kocuria varians		85.08	0.30	VPT 14			
Staphylococcus hominis subsp. hominis		14.92	0.16	SUC 12			
302640	**Good identification**						**302640**
					NOV		
Staphylococcus hominis subsp. hominis		92.16	0.98		1		
Staphylococcus warneri		4.09	0.76		1		
Staphylococcus hominis subsp. novobiosepticus		1.39	0.49		99		
302641	**Low discrimination**						**302641**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		37.44	0.59	ARG 99	1	99	
Kocuria kristinae		31.58	0.57		99	93	
Staphylococcus hominis subsp. hominis		21.55	0.63		1	1	
302642	**Very good identification**						**302642**
					NOV		
Staphylococcus hominis subsp. hominis		97.59	0.99		1		
Staphylococcus warneri		1.23	0.66		1		
Staphylococcus hominis subsp. novobiosepticus		0.68	0.44		99		
302643	**Low discrimination**						**302643**
					PYR		
Staphylococcus lugdunensis		36.94	0.59	ORN 99	96		
Staphylococcus hominis subsp. hominis		28.23	0.65		1		
Staphylococcus petrasii subsp. petrasii		22.50	0.53		99		
302650	**Low discrimination**						**302650**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		52.92	0.81	MAN 13	1	1	
Staphylococcus saprophyticus subsp. bovis		29.24	0.72		99	99	
Staphylococcus warneri		11.39	0.73		1	8	
302651	**Low discrimination**						**302651**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		69.10	0.59		99	99	
Staphylococcus petrasii subsp. petrasii		13.19	0.39		99	1	
Staphylococcus hominis subsp. hominis		10.21	0.46		1	1	
302652	**Low discrimination**						**302652**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		77.68	0.82	MAN 13	1	1	
Staphylococcus saprophyticus subsp. bovis		9.14	0.59		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		5.47	0.43		99	7	
302653	**Low discrimination**						**302653**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		39.45	0.47		99	99	
Staphylococcus hominis subsp. hominis		27.37	0.48		1	1	
Staphylococcus petrasii subsp. petrasii		16.22	0.34		99	1	
302660	**Good identification**						**302660**
					NOV		
Staphylococcus hominis subsp. hominis		91.83	0.57	XYL 1	1		
Staphylococcus warneri		4.08	0.35		1		
Staphylococcus hominis subsp. novobiosepticus		1.39	0.08		99		
302662	**Very good identification**						**302662**
					NOV		
Staphylococcus hominis subsp. hominis		97.79	0.58	XYL 1	1		
Staphylococcus warneri		1.23	0.25		1		
Staphylococcus hominis subsp. novobiosepticus		0.68	0.03		99		
302670	**Low discrimination**						**302670**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		53.63	0.40	MAN 13	1	1	
Staphylococcus saprophyticus subsp. bovis		29.63	0.31	XYL 1	99	99	
Staphylococcus warneri		11.54	0.32		1	8	
302672	**Low discrimination**						**302672**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		77.95	0.41	MAN 13	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. bovis		9.17	0.18	XYL 1	99	99		
Staphylococcus saprophyticus subsp. saprophyticus		5.49	0.02		99	7		
302700	**Low discrimination**							**302700**
Staphylococcus hominis subsp. hominis		69.00	0.58	MLT 99				
Kocuria varians		15.66	0.43					
Rothia mucilaginosa		5.55	0.22					
302701	**Low discrimination**							**302701**
					OXI			
Rothia mucilaginosa		57.69	0.32	URE 1	1			
Kocuria kristinae		24.22	0.28	ESL 90	99			
Staphylococcus petrasii subsp. petrasii		5.22	0.14		1			
302702	**Low discrimination**							**302702**
					NOV	OXI		
Staphylococcus hominis subsp. hominis		89.20	0.59	MLT 99	1	1		
Kocuria varians		4.86	0.32		1	13		
Macrococcus caseolyticus		2.52	0.14		99	99		
302710	**Low discrimination**							**302710**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		59.99	0.43	MLT 99	99	99		
Staphylococcus hominis subsp. hominis		29.41	0.41		1	1		
Staphylococcus warneri		9.61	0.36		1	8		
302711	**Low discrimination**							**302711**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		86.27	0.31	MLT 99	1	99	99	
Kocuria kristinae		4.78	-.01		99	1	93	
Staphylococcus hominis subsp. hominis		3.45	0.06		1	1	1	
302712	**Low discrimination**							**302712**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		65.48	0.42	MAN 13	1	1		
Staphylococcus saprophyticus subsp. bovis		28.45	0.31	MLT 99	99	99		
Staphylococcus warneri		6.08	0.26		1	8		
302720	**Low discrimination**							**302720**
Kocuria varians		88.76	0.34	VPT 14				
Staphylococcus hominis subsp. hominis		11.25	0.16	SUC 12				
302740	**Good identification**							**302740**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. hominis		92.01	0.99		1	1	49	
Staphylococcus hominis subsp. novobiosepticus		3.85	0.59		1	99	1	
Macrococcus caseolyticus		2.04	0.52		99	99	99	
302741	**Low discrimination**							**302741**
					PYR	NOV		
Staphylococcus petrasii subsp. petrasii		29.91	0.55	ARG 99	99	1		
Staphylococcus hominis subsp. hominis		27.98	0.64		1	1		
Staphylococcus hominis subsp. novobiosepticus		14.33	0.47		1	99		
302742	**Good identification**							**302742**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. hominis		94.78	1.00		1	1	49	
Macrococcus caseolyticus		2.68	0.55		99	99	99	
Staphylococcus hominis subsp. novobiosepticus		1.82	0.53		1	99	1	
302743	**Low discrimination**							**302743**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR			
Staphylococcus lugdunensis		38.11	0.59	ORN 99	96			
Staphylococcus hominis subsp. hominis		31.55	0.65		1			
Staphylococcus petrasii subsp. petrasii		15.47	0.50		99			
302750	**Low discrimination**							**302750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		65.47	0.84		99	99		
Staphylococcus hominis subsp. hominis		32.09	0.82		1	1		
Staphylococcus warneri		2.01	0.63		1	8		
302751	**Good identification**							**302751**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		92.20	0.72		99	99		
Staphylococcus hominis subsp. hominis		3.69	0.47		1	1		
Staphylococcus petrasii subsp. petrasii		2.93	0.36		99	1		
302752	**Low discrimination**							**302752**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		68.27	0.83	MAN 13	1	1		
Staphylococcus saprophyticus subsp. bovis		29.66	0.72		99	99		
Staphylococcus warneri		1.22	0.53		1	8		
302753	**Low discrimination**							**302753**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		72.46	0.59		99	99		
Staphylococcus hominis subsp. hominis		13.62	0.48		1	1		
Staphylococcus petrasii subsp. petrasii		4.97	0.30		99	1		
302760	**Good identification**							**302760**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. hominis		90.92	0.58	XYL 1	1	1	49	
Staphylococcus hominis subsp. novobiosepticus		3.80	0.18		1	99	1	
Micrococcus caseolyticus		2.01	0.11		99	99	99	
302762	**Good identification**							**302762**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. hominis		94.76	0.59	XYL 1	1	1	49	
Micrococcus caseolyticus		2.68	0.14		99	99	99	
Staphylococcus hominis subsp. novobiosepticus		1.82	0.12		1	99	1	
302770	**Low discrimination**							**302770**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		65.75	0.43	XYL 1	99	99		
Staphylococcus hominis subsp. hominis		32.23	0.41		1	1		
Staphylococcus warneri		2.02	0.21		1	8		
302771	**Low discrimination**							**302771**
					NOV	PYR		
Staphylococcus xylosus		71.43	0.36	VPT 14	99	65		
Staphylococcus saprophyticus subsp. bovis		26.66	0.31	GLR 95	99	99		
Staphylococcus hominis subsp. hominis		1.07	0.06		1	1		
302772	**Low discrimination**							**302772**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		67.31	0.42	MAN 13	1	1		
Staphylococcus saprophyticus subsp. bovis		29.24	0.31	XYL 1	99	99		
Staphylococcus xylosus		2.25	0.04		99	65		
302773	**Good identification**							**302773**
					NOV	PYR		
Staphylococcus xylosus		96.25	0.45	VPT 14	99	65		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		3.16	0.18	GLR 95	99	99	
Staphylococcus hominis subsp. hominis		0.59	0.07		1	1	
303000	**Low discrimination**						**303000**
					OXI		
Kocuria varians		57.85	0.39	VPT 14	13		
Dermacoccus nishinomiyaensis		37.37	0.28	ESL 1	93		
Kocuria rosea		3.67	0.05		35		
303002	**Good identification**						**303002**
Kocuria varians		100.0	0.28	VPT 14			
303020	**Acceptable identification**						**303020**
Kocuria varians		92.82	0.30	VPT 14			
Kocuria rosea		7.18	-.02	ESL 1			
303040	**Low discrimination**						**303040**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		77.11	0.27	ESL 8	99		
Kocuria varians		21.00	0.26	SUC 99	1		
Staphylococcus hominis subsp. hominis		1.89	0.05		1		
303100	**Acceptable identification**						**303100**
					OXI	NOV	
Kocuria varians		91.05	0.43	VPT 14	13	1	
Dermacoccus nishinomiyaensis		7.47	0.13	ESL 1	93	80	
Staphylococcus hominis subsp. novobiosepticus		1.48	-.04		1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
303102	**Good identification**							**303102**
Kocuria varians		100.0	0.31	VPT 14				
303120	**Good identification**							**303120**
Kocuria varians		100.0	0.33	VPT 14				
303140	**Low discrimination**							**303140**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		87.34	0.37	ESL 8	99			
Kocuria varians		11.89	0.29	SUC 99	1			
Staphylococcus hominis subsp. hominis		0.77	0.06		1			
303142	**Acceptable identification**							**303142**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		91.69	0.32	ESL 8	99			
Kocuria varians		6.54	0.18	SUC 99	1			
Staphylococcus hominis subsp. hominis		1.77	0.08		1			
303200	**Low discrimination**							**303200**
					OXI	NOV		
Staphylococcus hominis subsp. novobiosepticus		80.33	0.27	ESL 8	1	99		
Kocuria varians		13.59	0.21	MLT 99	13	1		
Dermacoccus nishinomiyaensis		4.11	0.03		93	80		
303240	**Very good identification**							**303240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		97.51	0.68	ESL 8	99			
Staphylococcus hominis subsp. hominis		2.39	0.47		1			
Staphylococcus epidermidis		0.06	0.11		1			
303241	**Very good identification**							**303241**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		98.14	0.56	ESL 8	1	99	1	
Kocuria kristinae		1.19	0.19		99	1	93	
Staphylococcus epidermidis		0.48	0.16		1	1	1	
303242	**Good identification**							**303242**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		94.58	0.63	ESL 8	99			
Staphylococcus hominis subsp. hominis		5.06	0.48		1			
Staphylococcus epidermidis		0.36	0.21		1			
303243	**Good identification**							**303243**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		96.09	0.50	ESL 8	1	99	1	
Staphylococcus epidermidis		2.75	0.27		1	1	1	
Kocuria kristinae		0.54	0.06		99	1	93	
303250	**Low discrimination**							**303250**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		73.37	0.27	ESL 8	99			
Staphylococcus hominis subsp. hominis		26.63	0.30	MAN 1	1			
303260	**Good identification**							**303260**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		97.61	0.27	ESL 8	99			
Staphylococcus hominis subsp. hominis		2.40	0.05	XYL 1	1			
303300	**Low discrimination**							**303300**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		89.79	0.37	ESL 8	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		7.60	0.25	MLT 99	1		
Rothia mucilaginosa		1.82	0.00		1		
303302	**Acceptable identification**						**303302**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		94.02	0.32	ESL 8	99		
Kocuria varians		4.17	0.14	MLT 99	1		
Staphylococcus hominis subsp. hominis		1.82	0.08		1		
303340	**Very good identification**						**303340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.94	0.78	ESL 8	99		
Staphylococcus hominis subsp. hominis		0.88	0.47		1		
Staphylococcus epidermidis		0.12	0.26		1		
303341	**Very good identification**						**303341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.33	0.66	ESL 8	99		
Staphylococcus epidermidis		0.90	0.32		1		
Rothia mucilaginosa		0.57	0.18		1		
303342	**Very good identification**						**303342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		97.41	0.73	ESL 8	99		
Staphylococcus hominis subsp. hominis		1.88	0.49		1		
Staphylococcus epidermidis		0.71	0.36		1		
303343	**Good identification**						**303343**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		94.68	0.60	ESL 8	99		
Staphylococcus epidermidis		5.11	0.42		1		
Staphylococcus hominis subsp. hominis		0.15	0.14		1		
303350	**Low discrimination**						**303350**
					PYR	NOV	
Staphylococcus hominis subsp. novobiosepticus		87.75	0.37	ESL 8	1	99	
Staphylococcus hominis subsp. hominis		11.50	0.30	MAN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		0.75	0.02		99	99	
303352	**Low discrimination**						**303352**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		77.78	0.32	ESL 8	99		
Staphylococcus hominis subsp. hominis		22.22	0.32	MAN 1	1		
303360	**Good identification**						**303360**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.49	0.37	ESL 8	99		
Staphylococcus hominis subsp. hominis		0.87	0.06	XYL 1	1		
Kocuria varians		0.64	0.02		1		
303362	**Good identification**						**303362**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.11	0.32	ESL 8	99		
Staphylococcus hominis subsp. hominis		1.90	0.08	XYL 1	1		
303550	**Low discrimination**						**303550**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		87.44	0.31	ESL 1	1	99	
Macrococcus equipersicus		9.23	0.10		99	1	
Macrococcus bovicus		3.33	0.00		99	1	
303640	**Low discrimination**						**303640**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		81.98	0.57	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		10.66	0.27		99		
Staphylococcus warneri		3.64	0.35		1		
303641	**Low discrimination**						**303641**
					OXI		
Kocuria kristinae		83.41	0.45	URE 10	99		
Rothia mucilaginosa		6.25	0.18	NIT 7	1		
Staphylococcus petrasii subsp. petrasii		4.00	0.18		1		
303642	**Good identification**						**303642**
					NOV		
Staphylococcus hominis subsp. hominis		92.98	0.58	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		5.54	0.22		99		
Staphylococcus warneri		1.17	0.25		1		
303643	**Low discrimination**						**303643**
					OXI	PYR	
Kocuria kristinae		66.82	0.32	URE 10	99	93	
Staphylococcus lugdunensis		11.32	0.18	NIT 7	1	96	
Staphylococcus hominis subsp. hominis		8.65	0.23		1	1	
303650	**Low discrimination**						**303650**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		53.63	0.40	ESL 1	1	1	
Staphylococcus saprophyticus subsp. bovis		29.63	0.31	MAN 13	99	99	
Staphylococcus warneri		11.54	0.32		1	8	
303652	**Low discrimination**						**303652**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		77.95	0.41	ESL 1	1	1	
Staphylococcus saprophyticus subsp. bovis		9.17	0.18	MAN 13	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		5.49	0.02		99	7	
303700	**Good identification**						**303700**
					NOV		
Rothia mucilaginosa		97.36	0.42	URE 1	1		
Staphylococcus hominis subsp. hominis		1.36	0.16		1		
Staphylococcus hominis subsp. novobiosepticus		0.49	-.04		99		
303701	**Good identification**						**303701**
					OXI		
Rothia mucilaginosa		98.85	0.51	URE 1	1		
Kocuria kristinae		1.15	0.15		99		
303713	**Good identification**						**303713**
Staphylococcus lentus		100.0	0.30	VPT 7			
303733	**Low discrimination**						**303733**
					OXI		
Staphylococcus lentus		64.62	0.30	VPT 7	99		
Staphylococcus equorum		21.62	0.15	URE 1	1		
Staphylococcus xylosus		13.76	0.10		1		
303740	**Low discrimination**						**303740**
					NOV		
Rothia mucilaginosa		54.55	0.49	URE 1	1		
Staphylococcus hominis subsp. hominis		32.31	0.58		1		
Staphylococcus hominis subsp. novobiosepticus		11.63	0.37		99		
303741	**Good identification**						**303741**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Rothia mucilaginosa		93.31	0.59	URE 1	1	1		
Kocuria kristinae		4.20	0.35		99	1		
Staphylococcus hominis subsp. novobiosepticus		1.66	0.25		1	99		
303742	**Low discrimination**							**303742**
					OXI	NOV	YEP	
Staphylococcus hominis subsp. hominis		82.22	0.59	ESL 1	1	1	49	
Staphylococcus hominis subsp. novobiosepticus		13.58	0.32		1	99	1	
Macrococcus caseolyticus		2.32	0.14		99	99	99	
303750	**Low discrimination**							**303750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		62.94	0.43	ESL 1	99	99		
Staphylococcus hominis subsp. hominis		30.85	0.41		1	1		
Rothia mucilaginosa		3.52	0.08		1	75		
303751	**Low discrimination**							**303751**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		40.94	0.31	ESL 1	1	99	99	
Rothia mucilaginosa		27.48	0.18		1	1	75	
Staphylococcus lentus		15.79	0.25		99	99	6	
303752	**Low discrimination**							**303752**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		68.24	0.42	ESL 1	1	1		
Staphylococcus saprophyticus subsp. bovis		29.65	0.31	MAN 13	99	99		
Staphylococcus warneri		1.22	0.12		1	8		
303753	**Low discrimination**							**303753**
					OXI	PYR		
Staphylococcus lentus		73.47	0.36	VPT 7	99	6		
Staphylococcus saprophyticus subsp. bovis		13.43	0.18	URE 1	1	99		
Staphylococcus xylosus		8.91	0.11		1	65		
303771	**Low discrimination**							**303771**
					OXI			
Staphylococcus equorum		41.68	0.27	VPT 1	1			
Staphylococcus xylosus		31.50	0.23	GLR 90	1			
Staphylococcus lentus		18.15	0.25		99			
303773	**Low discrimination**							**303773**
					OXI			
Staphylococcus equorum		39.55	0.36	VPT 1	1			
Staphylococcus xylosus		33.09	0.33	GLR 90	1			
Staphylococcus lentus		23.74	0.36		99			
304000	**Acceptable identification**							**304000**
					OXI			
Dermacoccus nishinomiyaensis		91.35	0.47	VPT 5	93			
Micrococcus sp.		5.52	0.13	PHS 5	89			
Kocuria varians		3.13	0.25		13			
304052	**Good identification**							**304052**
Staphylococcus kloosii		100.0	0.28	VPT 10				
304100	**Low discrimination**							**304100**
					OXI			
Dermacoccus nishinomiyaensis		78.77	0.33	VPT 5	93			
Kocuria varians		21.23	0.29	PHS 5	13			
304150	**Low discrimination**							**304150**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		78.63	0.33	VPT 10	1			
Macrococcus equipersicus		21.37	0.04	TRE 99	99			
304152	**Low discrimination**							**304152**
					NOV	YEP		
Staphylococcus kloosii		59.58	0.33	VPT 10	99	99		
Staphylococcus caprae		40.42	0.02	TRE 99	1	1		
304240	**Low discrimination**							**304240**
					COA	NOV		
Staphylococcus hominis subsp. novobiosepticus		42.14	0.30	NIT 90	1	99		
Staphylococcus epidermidis		37.08	0.36	PHS 1	1	1		
Staphylococcus aureus subsp. anaerobius		16.92	0.16		99	1		
304241	**Low discrimination**							**304241**
					OXI	NOV	PYR	
Staphylococcus epidermidis		81.30	0.42	NIT 97	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		12.44	0.17		1	99	1	
Kocuria kristinae		6.26	0.13		99	1	93	
304242	**Low discrimination**							**304242**
					NOV			
Staphylococcus epidermidis		82.10	0.47	NIT 97	1			
Staphylococcus hominis subsp. novobiosepticus		15.87	0.24		99			
Staphylococcus hominis subsp. hominis		2.03	0.17		1			
304243	**Good identification**							**304243**
					OXI	NOV	PYR	
Staphylococcus epidermidis		96.89	0.52	NIT 97	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		2.52	0.12		1	99	1	
Kocuria kristinae		0.59	0.01		99	1	93	
304340	**Low discrimination**							**304340**
					NOV			
Staphylococcus epidermidis		61.94	0.52	NIT 97	1			
Staphylococcus hominis subsp. novobiosepticus		37.27	0.39		99			
Staphylococcus hominis subsp. hominis		0.79	0.16		1			
304341	**Good identification**							**304341**
					OXI	NOV	PYR	
Staphylococcus epidermidis		92.12	0.57	NIT 97	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		7.46	0.27		1	99	1	
Kocuria kristinae		0.42	0.03		99	1	93	
304342	**Good identification**							**304342**
					NOV			
Staphylococcus epidermidis		90.33	0.62	NIT 97	1			
Staphylococcus hominis subsp. novobiosepticus		9.24	0.34		99			
Staphylococcus hominis subsp. hominis		0.43	0.18		1			
304343	**Very good identification**							**304343**
					NOV			
Staphylococcus epidermidis		98.64	0.68	NIT 97	1			
Staphylococcus hominis subsp. novobiosepticus		1.36	0.21		99			
304410	**Low discrimination**							**304410**
					OXI	YEP		
Staphylococcus kloosii		52.29	0.42	VPT 10	1	99		
Staphylococcus cohnii subsp. cohnii		39.82	0.27	MLT 95	1	12		
Macrococcus equipersicus		4.57	0.02		99	99		
304411	**Low discrimination**							**304411**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. cohnii		82.11	0.27	URE 1			
Staphylococcus cohnii subsp. urealyticum		17.89	0.14	PHS 9			
304412	**Good identification**						**304412**
Staphylococcus kloosii		100.0	0.42	VPT 10			
304413	**Good identification**						**304413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	GLR 99			
304430	**Good identification**						**304430**
Staphylococcus kloosii		100.0	0.29	VPT 10			
304432	**Good identification**						**304432**
Staphylococcus kloosii		100.0	0.30	VPT 10			
304440	**Low discrimination**						**304440**
					NOV	YEP	
Staphylococcus cohnii subsp. cohnii		85.30	0.35	URE 1	99	12	
Staphylococcus kloosii		8.64	0.27	PHS 9	99	99	
Staphylococcus warneri		6.06	0.16		1	43	
304441	**Acceptable identification**						**304441**
					OXI	NOV	
Staphylococcus cohnii subsp. cohnii		91.13	0.35	URE 1	1	99	
Staphylococcus cohnii subsp. urealyticum		4.47	0.09	PHS 9	1	99	
Kocuria kristinae		4.40	-.02		99	1	
304442	**Low discrimination**						**304442**
Staphylococcus kloosii		62.58	0.28	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		37.42	-.06	MAN 99			
304443	**Good identification**						**304443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	GLR 99			
304450	**Low discrimination**						**304450**
					YEP		
Staphylococcus kloosii		84.27	0.68	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		10.70	0.37		12		
Staphylococcus saprophyticus subsp. saprophyticus		2.34	0.07		16		
304451	**Low discrimination**						**304451**
					YEP		
Staphylococcus cohnii subsp. cohnii		68.34	0.38	URE 1	12		
Staphylococcus cohnii subsp. urealyticum		26.65	0.30	PHS 9	64		
Staphylococcus kloosii		5.02	0.27		99		
304452	**Low discrimination**						**304452**
Staphylococcus kloosii		86.96	0.69	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		12.60	0.23				
Staphylococcus cohnii subsp. urealyticum		0.44	0.10				
304453	**Good identification**						**304453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		96.35	0.51	GLR 99	99	64	
Staphylococcus kloosii		1.95	0.28		99	99	
Staphylococcus caprae		1.71	-.02		1	1	
304470	**Very good identification**						**304470**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.55	VPT 10			
304472	**Very good identification**						**304472**
Staphylococcus kloosii		100.0	0.56	VPT 10			
304510	**Good identification**						**304510**
					YEP		
Staphylococcus kloosii		95.82	0.47	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		4.18	0.07	MLT 95	12		
304512	**Good identification**						**304512**
Staphylococcus kloosii		100.0	0.48	VPT 10			
304513	**Low discrimination**						**304513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		81.02	0.28	GLR 99	99		
Staphylococcus caprae		18.98	-.02		1		
304530	**Good identification**						**304530**
Staphylococcus kloosii		100.0	0.35	VPT 10			
304532	**Good identification**						**304532**
Staphylococcus kloosii		100.0	0.36	VPT 10			
304540	**Low discrimination**						**304540**
					YEP		
Staphylococcus kloosii		63.89	0.33	VPT 10	99		
Staphylococcus cohnii subsp. cohnii		36.11	0.15	MAN 99	12		
304542	**Low discrimination**						**304542**
					NOV	YEP	
Staphylococcus kloosii		75.68	0.33	VPT 10	99	99	
Staphylococcus caprae		24.32	-.05	MAN 99	1	1	
304550	**Very good identification**						**304550**
					PYR	YEP	
Staphylococcus kloosii		97.18	0.74	VPT 10	52	99	
Staphylococcus saprophyticus subsp. bovis		1.34	0.18		99	80	
Staphylococcus cohnii subsp. cohnii		0.71	0.17		1	12	
304552	**Good identification**						**304552**
					NOV	YEP	
Staphylococcus kloosii		93.34	0.75	VPT 10	99	99	
Staphylococcus caprae		5.76	0.22		1	1	
Staphylococcus saprophyticus subsp. saprophyticus		0.61	0.01		99	16	
304553	**Low discrimination**						**304553**
					NOV	YEP	
Staphylococcus caprae		78.95	0.39	ARG 99	1	1	
Staphylococcus cohnii subsp. urealyticum		19.29	0.43	NIT 90	99	64	
Staphylococcus kloosii		1.76	0.33		99	99	
304570	**Very good identification**						**304570**
Staphylococcus kloosii		100.0	0.61	VPT 10			
304572	**Very good identification**						**304572**
Staphylococcus kloosii		100.0	0.62	VPT 10			
304641	**Good identification**						**304641**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	PYR	
Kocuria kristinae		98.35	0.39	URE 10	99	93	
Staphylococcus petrasii subsp. petrasii		0.98	-.02	PHS 1	1	99	
Staphylococcus epidermidis		0.68	0.01		1	1	
304643	**Acceptable identification**						**304643**
					OXI	PYR	
Kocuria kristinae		91.96	0.27	URE 10	99	93	
Staphylococcus epidermidis		8.04	0.11	PHS 1	1	1	
304650	**Low discrimination**						**304650**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		54.19	0.28	PHS 1	99		
Staphylococcus kloosii		19.13	0.47		99		
Staphylococcus warneri		18.85	0.39		1		
304652	**Acceptable identification**						**304652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		91.03	0.43	PHS 1	99		
Staphylococcus kloosii		6.14	0.48		99		
Staphylococcus warneri		1.86	0.29		1		
304670	**Good identification**						**304670**
Staphylococcus kloosii		100.0	0.35	VPT 10			
304672	**Low discrimination**						**304672**
Staphylococcus kloosii		62.55	0.36	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		37.45	0.02	SUC 9			
304710	**Good identification**						**304710**
Staphylococcus kloosii		100.0	0.27	VPT 10			
304712	**Good identification**						**304712**
Staphylococcus kloosii		100.0	0.28	VPT 10			
304741	**Low discrimination**						**304741**
					OXI	PYR	
Kocuria kristinae		87.94	0.30	URE 10	99	93	
Staphylococcus epidermidis		10.32	0.16	PHS 1	1	1	
Staphylococcus petrasii subsp. petrasii		1.74	-.05		1	99	
304742	**Low discrimination**						**304742**
Staphylococcus hominis subsp. hominis		46.90	0.28	NIT 99			
Staphylococcus epidermidis		31.59	0.21	PHS 3			
Staphylococcus warneri		13.00	0.22				
304750	**Low discrimination**						**304750**
					NOV	PYR	
Staphylococcus kloosii		53.20	0.53	VPT 10	99	52	
Staphylococcus saprophyticus subsp. bovis		29.72	0.31	SUC 9	99	99	
Staphylococcus warneri		8.53	0.29		1	8	
304752	**Low discrimination**						**304752**
					NOV	YEP	
Staphylococcus kloosii		47.41	0.54	VPT 10	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		31.47	0.22	SUC 9	99	16	
Staphylococcus caprae		10.93	0.13		1	1	
304753	**Low discrimination**						**304753**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COA	NOV	
Staphylococcus caprae		68.11	0.30	ARG 99	1	1	
Staphylococcus aureus subsp. aureus		29.20	0.18	NIT 90	99	4	
Staphylococcus cohnii subsp. urealyticum		1.39	0.12		1	99	
304770	**Good identification**						**304770**
Staphylococcus kloosii		100.0	0.41	VPT 10			
304772	**Good identification**						**304772**
Staphylococcus kloosii		100.0	0.41	VPT 10			
305052	**Good identification**						**305052**
Staphylococcus kloosii		100.0	0.28	VPT 10			
305150	**Low discrimination**						**305150**
					OXI		
Staphylococcus kloosii		63.39	0.33	VPT 10	1		
Macrococcus equipersicus		36.61	0.10	TRE 99	99		
305152	**Good identification**						**305152**
Staphylococcus kloosii		100.0	0.33	VPT 10			
305410	**Low discrimination**						**305410**
					OXI		
Staphylococcus kloosii		84.34	0.42	VPT 10	1		
Macrococcus equipersicus		15.66	0.09	MLT 95	99		
305412	**Good identification**						**305412**
Staphylococcus kloosii		100.0	0.42	VPT 10			
305413	**Good identification**						**305413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	GLR 99			
305430	**Good identification**						**305430**
Staphylococcus kloosii		100.0	0.29	VPT 10			
305432	**Good identification**						**305432**
Staphylococcus kloosii		100.0	0.30	VPT 10			
305440	**Good identification**						**305440**
Staphylococcus kloosii		100.0	0.27	VPT 10			
305442	**Good identification**						**305442**
Staphylococcus kloosii		100.0	0.28	VPT 10			
305450	**Very good identification**						**305450**
					OXI		
Staphylococcus kloosii		97.43	0.68	VPT 10	1		
Macrococcus equipersicus		2.57	0.18		99		
305452	**Very good identification**						**305452**
Staphylococcus kloosii		100.0	0.69	VPT 10			
305453	**Good identification**						**305453**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		95.07	0.42	GLR 99			
Staphylococcus kloosii		4.93	0.28				
305470	**Very good identification**						**305470**
Staphylococcus kloosii		100.0	0.55	VPT 10			
305472	**Very good identification**						**305472**
Staphylococcus kloosii		100.0	0.56	VPT 10			
305510	**Good identification**						**305510**
					OXI		
Staphylococcus kloosii		96.06	0.47	VPT 10	1		
Macrococcus equipersicus		3.94	0.01	MLT 95	99		
305512	**Good identification**						**305512**
Staphylococcus kloosii		100.0	0.48	VPT 10			
305530	**Good identification**						**305530**
Staphylococcus kloosii		100.0	0.35	VPT 10			
305532	**Good identification**						**305532**
Staphylococcus kloosii		100.0	0.36	VPT 10			
305540	**Good identification**						**305540**
Staphylococcus kloosii		100.0	0.33	VPT 10			
305542	**Good identification**						**305542**
Staphylococcus kloosii		100.0	0.33	VPT 10			
305550	**Very good identification**						**305550**
					OXI		
Staphylococcus kloosii		99.42	0.74	VPT 10	1		
Macrococcus equipersicus		0.58	0.10		99		
305551	**Low discrimination**						**305551**
Staphylococcus kloosii		68.69	0.33	VPT 10			
Staphylococcus cohnii subsp. urealyticum		31.31	0.14	MNS 1			
305552	**Very good identification**						**305552**
Staphylococcus kloosii		100.0	0.75	VPT 10			
305553	**Low discrimination**						**305553**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		74.55	0.35	GLR 99	99	64	
Staphylococcus kloosii		17.53	0.33		99	99	
Staphylococcus caprae		7.93	-.02		1	1	
305570	**Very good identification**						**305570**
Staphylococcus kloosii		100.0	0.61	VPT 10			
305572	**Very good identification**						**305572**
Staphylococcus kloosii		100.0	0.62	VPT 10			
305641	**Good identification**						**305641**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Kocuria kristinae		100.0	0.27	URE 10		
305650	**Good identification**					**305650**
Staphylococcus kloosii		100.0	0.47	VPT 10		
305652	**Low discrimination**					**305652**
Staphylococcus kloosii		86.98	0.48	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		13.02	0.02	SUC 9		
305670	**Good identification**					**305670**
Staphylococcus kloosii		100.0	0.35	VPT 10		
305672	**Good identification**					**305672**
Staphylococcus kloosii		100.0	0.36	VPT 10		
305710	**Good identification**					**305710**
Staphylococcus kloosii		100.0	0.27	VPT 10		
305712	**Good identification**					**305712**
Staphylococcus kloosii		100.0	0.28	VPT 10		
305750	**Very good identification**					**305750**
Staphylococcus kloosii		100.0	0.53	VPT 10		
305752	**Very good identification**					**305752**
Staphylococcus kloosii		100.0	0.54	VPT 10		
305770	**Good identification**					**305770**
Staphylococcus kloosii		100.0	0.41	VPT 10		
305772	**Good identification**					**305772**
Staphylococcus kloosii		100.0	0.41	VPT 10		
306000	**Low discrimination**					**306000**
					OXI	
Dermacoccus nishinomiyaensis		75.67	0.42	VPT 5	93	
Kocuria varians		22.48	0.39	PHS 5	13	
Kocuria rosea		1.43	0.05		35	
306001	**Low discrimination**					**306001**
Staphylococcus schleiferi subsp. schleiferi		78.85	0.29	URE 1		
Kocuria varians		11.49	0.17	ARG 86		
Staphylococcus capitis subsp. capitis		5.17	0.04			
306002	**Low discrimination**					**306002**
					OXI	
Kocuria varians		89.24	0.28	VPT 14	13	
Dermacoccus nishinomiyaensis		10.76	0.01	PHS 1	93	
306011	**Good identification**					**306011**
Staphylococcus capitis subsp. capitis		100.0	0.27	URE 1		
306020	**Low discrimination**					**306020**
					OXI	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Kocuria varians		85.17	0.30	VPT 14	13	
Dermacoccus nishinomiyaensis		8.24	0.01	PHS 1	93	
Kocuria rosea		6.59	-.02		35	
306041	**Acceptable identification**					**306041**
					NOV	
Staphylococcus epidermidis		92.82	0.32	SUC 99	1	
Staphylococcus hominis subsp. novobiosepticus		3.95	-.04		99	
Kocuria varians		3.22	0.04		1	
306042	**Low discrimination**					**306042**
Staphylococcus epidermidis		83.63	0.36	SUC 99		
Staphylococcus hominis subsp. hominis		6.34	0.17			
Kocuria varians		5.52	0.14			
306043	**Good identification**					**306043**
Staphylococcus epidermidis		100.0	0.42	SUC 99		
306100	**Low discrimination**					**306100**
					OXI	
Kocuria varians		69.51	0.43	VPT 14	13	
Dermacoccus nishinomiyaensis		29.71	0.27	PHS 1	93	
Staphylococcus epidermidis		0.79	0.01		1	
306101	**Low discrimination**					**306101**
					COA	
Staphylococcus schleiferi subsp. coagulans		55.22	0.37	ARG 99	99	
Staphylococcus schleiferi subsp. schleiferi		35.23	0.29	LAC 86	1	
Kocuria varians		7.70	0.21		1	
306102	**Low discrimination**					**306102**
					COA	
Kocuria varians		79.11	0.31	VPT 14	1	
Staphylococcus schleiferi subsp. coagulans		10.86	0.12	PHS 1	99	
Staphylococcus epidermidis		10.03	0.11		1	
306103	**Very good identification**					**306103**
					COA	
Staphylococcus schleiferi subsp. coagulans		97.68	0.53	ARG 99	99	
Staphylococcus epidermidis		1.69	0.16		1	
Kocuria varians		0.63	0.10		1	
306111	**Good identification**					**306111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	ARG 99		
306113	**Good identification**					**306113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.14	0.53	ARG 99	99	
Staphylococcus caprae		0.86	-.02		1	
306120	**Good identification**					**306120**
Kocuria varians		100.0	0.33	VPT 14		
306140	**Low discrimination**					**306140**
					NOV	
Staphylococcus epidermidis		71.28	0.42	SUC 99	1	
Kocuria varians		13.99	0.29		1	
Staphylococcus hominis subsp. novobiosepticus		11.94	0.18		99	
306141	**Good identification**					**306141**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus epidermidis		96.92	0.47	SUC 99	1		
Staphylococcus hominis subsp. novobiosepticus		2.19	0.05		99		
Kocuria varians		0.89	0.07		1		
306142	**Acceptable identification**						**306142**
					NOV		
Staphylococcus epidermidis		93.76	0.52	SUC 99	1		
Staphylococcus hominis subsp. novobiosepticus		2.67	0.12		99		
Kocuria varians		1.64	0.18		1		
306143	**Very good identification**						**306143**
					COA		
Staphylococcus epidermidis		96.72	0.58	SUC 99	1		
Staphylococcus caprae		2.34	0.13		1		
Staphylococcus schleiferi subsp. coagulans		0.57	0.12		99		
306153	**Good identification**						**306153**
					COA		
Staphylococcus caprae		96.74	0.39	ARG 99	1		
Staphylococcus epidermidis		2.12	0.16	TRE 90	1		
Staphylococcus schleiferi subsp. coagulans		1.14	0.12		99		
306201	**Low discrimination**						**306201**
					NOV		
Staphylococcus epidermidis		85.12	0.32	MLT 99	1		
Staphylococcus capitis subsp. capitis		11.25	0.09		1		
Staphylococcus hominis subsp. novobiosepticus		3.63	-.04		99		
306202	**Low discrimination**						**306202**
					NOV		
Staphylococcus epidermidis		85.42	0.36	MLT 99	1		
Staphylococcus hominis subsp. hominis		6.48	0.17		1		
Staphylococcus hominis subsp. novobiosepticus		4.60	0.03		99		
306203	**Good identification**						**306203**
					COA	YEP	
Staphylococcus epidermidis		97.90	0.42	MLT 99	1	1	
Staphylococcus chromogenes		1.07	-.02		1	95	
Staphylococcus schleiferi subsp. coagulans		1.03	0.02		99	1	
306211	**Good identification**						**306211**
Staphylococcus capitis subsp. capitis		100.0	0.32	URE 1			
306240	**Low discrimination**						**306240**
					NOV		
Staphylococcus epidermidis		65.71	0.67		1		
Staphylococcus hominis subsp. novobiosepticus		20.79	0.49		99		
Staphylococcus hominis subsp. hominis		13.44	0.57		1		
306241	**Very good identification**						**306241**
					NOV		
Staphylococcus epidermidis		95.71	0.73		1		
Staphylococcus hominis subsp. novobiosepticus		4.08	0.37		99		
Staphylococcus hominis subsp. hominis		0.22	0.22		1		
306242	**Low discrimination**						**306242**
					NOV		
Staphylococcus epidermidis		88.52	0.78		1		
Staphylococcus hominis subsp. hominis		6.71	0.58		1		
Staphylococcus hominis subsp. novobiosepticus		4.76	0.44		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
306243	**Very good identification**						**306243**
					NOV		
Staphylococcus epidermidis		99.18	0.83		1		
Staphylococcus hominis subsp. novobiosepticus		0.72	0.31		99		
Staphylococcus hominis subsp. hominis		0.08	0.23		1		
306250	**Low discrimination**						**306250**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		63.38	0.40	PHS 3	1	1	
Staphylococcus epidermidis		20.94	0.26	MAN 13	1	1	
Staphylococcus saprophyticus subsp. bovis		9.05	0.18		99	99	
306251	**Low discrimination**						**306251**
					PYR	NOV	
Staphylococcus epidermidis		78.96	0.32	MAN 1	1	1	
Staphylococcus capitis subsp. ureolyticus		10.46	0.00		1	1	
Staphylococcus saprophyticus subsp. bovis		4.60	0.06		99	99	
306252	**Low discrimination**						**306252**
					NOV		
Staphylococcus hominis subsp. hominis		50.77	0.41	PHS 3	1		
Staphylococcus epidermidis		45.25	0.36	MAN 13	1		
Staphylococcus hominis subsp. novobiosepticus		2.43	0.03		99		
306253	**Low discrimination**						**306253**
					COA		
Staphylococcus epidermidis		62.45	0.42	MAN 1	1		
Staphylococcus capitis subsp. ureolyticus		19.15	0.18		1		
Staphylococcus delphini		16.49	0.18		99		
306261	**Good identification**						**306261**
					NOV		
Staphylococcus epidermidis		95.92	0.32	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		4.09	-.04		99		
306262	**Low discrimination**						**306262**
					NOV		
Staphylococcus epidermidis		88.52	0.36	XYL 1	1		
Staphylococcus hominis subsp. hominis		6.71	0.17		1		
Staphylococcus hominis subsp. novobiosepticus		4.76	0.03		99		
306263	**Good identification**						**306263**
Staphylococcus epidermidis		100.0	0.42	XYL 1			
306300	**Low discrimination**						**306300**
					NOV		
Staphylococcus epidermidis		73.91	0.42	MLT 99	1		
Staphylococcus hominis subsp. novobiosepticus		12.38	0.18		99		
Kocuria varians		9.01	0.25		1		
306301	**Low discrimination**						**306301**
					COA	NOV	
Staphylococcus epidermidis		88.56	0.47	MLT 99	1	1	
Staphylococcus schleiferi subsp. coagulans		8.40	0.27		99	1	
Staphylococcus hominis subsp. novobiosepticus		2.00	0.05		1	99	
306302	**Acceptable identification**						**306302**
					NOV		
Staphylococcus epidermidis		94.58	0.52	MLT 99	1		
Staphylococcus hominis subsp. novobiosepticus		2.69	0.12		99		
Staphylococcus hominis subsp. hominis		1.37	0.18		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
306303	**Low discrimination**						**306303**
					COA	YEP	
Staphylococcus epidermidis		82.83	0.58	MLT 99	1	1	
Staphylococcus schleiferi subsp. coagulans		15.24	0.43		99	1	
Staphylococcus chromogenes		1.44	0.17		1	95	
306313	**Acceptable identification**						**306313**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		91.04	0.42	ARG 99	99	1	
Staphylococcus epidermidis		5.42	0.16		1	1	
Staphylococcus chromogenes		2.62	0.06		1	95	
306340	**Low discrimination**						**306340**
					NOV		
Staphylococcus epidermidis		82.86	0.83		1		
Staphylococcus hominis subsp. novobiosepticus		13.88	0.59		99		
Staphylococcus hominis subsp. hominis		3.24	0.57		1		
306341	**Very good identification**						**306341**
					NOV		
Staphylococcus epidermidis		97.75	0.88		1		
Staphylococcus hominis subsp. novobiosepticus		2.20	0.47		99		
Staphylococcus hominis subsp. hominis		0.04	0.22		1		
306342	**Very good identification**						**306342**
					NOV		
Staphylococcus epidermidis		95.88	0.93		1		
Staphylococcus hominis subsp. novobiosepticus		2.73	0.53		99		
Staphylococcus hominis subsp. hominis		1.39	0.59		1		
306343	**Excellent identification**						**306343**
					NOV		
Staphylococcus epidermidis		99.57	0.99		1		
Staphylococcus hominis subsp. novobiosepticus		0.38	0.41		99		
Staphylococcus hominis subsp. hominis		0.02	0.24		1		
306350	**Low discrimination**						**306350**
					PYR	NOV	
Staphylococcus epidermidis		48.75	0.42	MAN 1	1	1	
Staphylococcus hominis subsp. hominis		28.21	0.40		1	1	
Staphylococcus saprophyticus subsp. bovis		14.88	0.31		99	99	
306351	**Acceptable identification**						**306351**
					NOV	PYR	
Staphylococcus epidermidis		91.05	0.47	MAN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		3.74	0.18		99	99	
Staphylococcus caprae		2.58	0.04		1	75	
306352	**Low discrimination**						**306352**
Staphylococcus epidermidis		76.33	0.52	MAN 1			
Staphylococcus hominis subsp. hominis		16.37	0.42				
Staphylococcus caprae		3.29	0.13				
306353	**Low discrimination**						**306353**
					COA		
Staphylococcus epidermidis		76.58	0.58	MAN 1	1		
Staphylococcus caprae		13.03	0.30		1		
Staphylococcus aureus subsp. aureus		5.59	0.18		99		
306360	**Low discrimination**						**306360**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		82.24	0.42	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		13.77	0.18		99		
Staphylococcus hominis subsp. hominis		3.22	0.16		1		
306361	**Good identification**						**306361**
					NOV		
Staphylococcus epidermidis		97.80	0.47	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		2.21	0.05		99		
306362	**Good identification**						**306362**
					NOV		
Staphylococcus epidermidis		95.88	0.52	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		2.73	0.12		99		
Staphylococcus hominis subsp. hominis		1.39	0.18		1		
306363	**Very good identification**						**306363**
					NOV		
Staphylococcus epidermidis		99.62	0.58	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		0.38	0.00		99		
306401	**Good identification**						**306401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	URE 1			
306442	**Good identification**						**306442**
Staphylococcus hominis subsp. hominis		100.0	0.27	PHS 3			
306450	**Good identification**						**306450**
					PYR	YEP	
Staphylococcus saprophyticus subsp. bovis		96.28	0.47		99	80	
Staphylococcus kloosii		1.47	0.27		52	99	
Staphylococcus cohnii subsp. cohnii		1.39	0.14		1	12	
306451	**Low discrimination**						**306451**
					PYR		
Staphylococcus saprophyticus subsp. bovis		87.20	0.34		99		
Staphylococcus cohnii subsp. cohnii		5.44	0.14		1		
Staphylococcus cohnii subsp. urealyticum		4.97	0.14		53		
306452	**Low discrimination**						**306452**
					NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		84.78	0.34		99	80	
Staphylococcus caprae		6.03	0.00		1	1	
Staphylococcus kloosii		5.59	0.28		99	99	
306453	**Low discrimination**						**306453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		42.20	0.35	GLR 99	99		
Staphylococcus caprae		38.14	0.18		1		
Staphylococcus saprophyticus subsp. bovis		18.30	0.22		99		
306501	**Good identification**						**306501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	URE 1			
306543	**Acceptable identification**						**306543**
					YEP		
Staphylococcus caprae		93.93	0.33	ARG 99	1		
Staphylococcus epidermidis		4.35	0.16	MAN 95	1		
Staphylococcus chromogenes		0.87	-.02		95		
306550	**Very good identification**						**306550**
					NOV	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		96.81	0.59		99	80	
Staphylococcus caprae		2.24	0.15		1	1	
Staphylococcus kloosii		0.72	0.33		99	99	
306551	**Low discrimination**						**306551**
					NOV		
Staphylococcus saprophyticus subsp. bovis		59.35	0.47		99		
Staphylococcus caprae		40.29	0.33		1		
Staphylococcus cohnii subsp. urealyticum		0.36	0.07		99		
306552	**Low discrimination**						**306552**
					NOV	YEP	
Staphylococcus caprae		62.76	0.41	ARG 99	1	1	
Staphylococcus saprophyticus subsp. bovis		35.68	0.47	MNS 88	99	80	
Staphylococcus kloosii		1.14	0.33		99	99	
306553	**Very good identification**						**306553**
					COA	NOV	
Staphylococcus caprae		95.99	0.59	ARG 99	1	1	
Staphylococcus saprophyticus subsp. bovis		1.86	0.34		1	99	
Staphylococcus aureus subsp. aureus		1.69	0.18		99	4	
306602	**Good identification**						**306602**
Staphylococcus hominis subsp. hominis		100.0	0.27	PHS 3			
306640	**Very good identification**						**306640**
Staphylococcus hominis subsp. hominis		95.49	0.67	PHS 3			
Staphylococcus epidermidis		1.49	0.26				
Staphylococcus warneri		1.39	0.35				
306641	**Low discrimination**						**306641**
					PYR		
Staphylococcus epidermidis		38.08	0.32	TRE 1	1		
Staphylococcus hominis subsp. hominis		26.84	0.32		1		
Staphylococcus petrasii subsp. petrasii		15.23	0.18		99		
306642	**Very good identification**						**306642**
Staphylococcus hominis subsp. hominis		95.25	0.68	PHS 3			
Staphylococcus epidermidis		4.01	0.36				
Staphylococcus warneri		0.39	0.25				
306643	**Low discrimination**						**306643**
					YEP		
Staphylococcus epidermidis		62.44	0.42	TRE 1	1		
Staphylococcus hominis subsp. hominis		16.32	0.33		49		
Staphylococcus chromogenes		7.84	0.20		95		
306650	**Low discrimination**						**306650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		80.21	0.59		99	99	
Staphylococcus hominis subsp. hominis		17.96	0.50		1	1	
Staphylococcus warneri		1.26	0.32		1	8	
306651	**Good identification**						**306651**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		97.03	0.47		99	99	
Staphylococcus hominis subsp. hominis		1.77	0.15		1	1	
Staphylococcus petrasii subsp. petrasii		0.75	-.02		99	1	
306652	**Low discrimination**						**306652**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Staphylococcus hominis subsp. hominis		49.61	0.51	PHS 3	1	1	
Staphylococcus saprophyticus subsp. bovis		47.18	0.47	MAN 13	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		1.14	0.02		99	7	
306662	**Good identification**						**306662**
Staphylococcus hominis subsp. hominis		100.0	0.27	PHS 3			
306702	**Low discrimination**						**306702**
					YEP		
Staphylococcus hominis subsp. hominis		76.84	0.28	PHS 3	49		
Staphylococcus epidermidis		16.90	0.11	MLT 99	1		
Staphylococcus chromogenes		6.26	-.02		95		
306703	**Low discrimination**						**306703**
					YEP		
Staphylococcus chromogenes		80.35	0.39	VPT 1	95		
Staphylococcus hyicus		10.19	0.17	ARG 90	1		
Staphylococcus epidermidis		4.07	0.16		1		
306710	**Acceptable identification**						**306710**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		94.28	0.31	MLT 99	99	99	
Staphylococcus hominis subsp. hominis		5.72	0.09		1	1	
306740	**Low discrimination**						**306740**
					PYR	NOV	
Staphylococcus hominis subsp. hominis		88.31	0.68	PHS 3	1	1	
Staphylococcus epidermidis		7.20	0.42		1	1	
Staphylococcus saprophyticus subsp. bovis		2.20	0.31		99	99	
306741	**Low discrimination**						**306741**
					YEP		
Staphylococcus epidermidis		73.10	0.47	TRE 1	1		
Staphylococcus hominis subsp. hominis		9.85	0.33		49		
Staphylococcus chromogenes		5.13	0.20		95		
306742	**Low discrimination**						**306742**
					OXI	NOV	
Staphylococcus hominis subsp. hominis		80.17	0.69	PHS 3	1	1	
Staphylococcus epidermidis		17.64	0.52		1	1	
Micrococcus caseolyticus		0.74	0.14		99	99	
306743	**Low discrimination**						**306743**
					YEP		
Staphylococcus epidermidis		69.21	0.58	TRE 1	1		
Staphylococcus chromogenes		13.80	0.39		95		
Staphylococcus caprae		5.58	0.24		1		
306750	**Good identification**						**306750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		93.95	0.72		99	99	
Staphylococcus hominis subsp. hominis		5.70	0.51		1	1	
Staphylococcus caprae		0.20	0.06		1	75	
306751	**Low discrimination**						**306751**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		87.89	0.59		1	99	
Staphylococcus caprae		5.52	0.24		1	1	
Staphylococcus aureus subsp. aureus		4.99	0.18		99	4	
306752	**Low discrimination**						**306752**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		66.22	0.59		99	99	
Staphylococcus hominis subsp. hominis		18.86	0.52		1	1	
Staphylococcus caprae		10.77	0.32		1	75	
306753	**Low discrimination**						**306753**
					COA	NOV	
Staphylococcus aureus subsp. aureus		78.73	0.59	ARG 99	99	4	
Staphylococcus caprae		16.70	0.50		1	1	
Staphylococcus saprophyticus subsp. bovis		3.50	0.47		1	99	
306762	**Low discrimination**						**306762**
Staphylococcus hominis subsp. hominis		81.97	0.28	PHS 3			
Staphylococcus epidermidis		18.03	0.11	XYL 1			
306770	**Acceptable identification**						**306770**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		94.28	0.31	XYL 1	99	99	
Staphylococcus hominis subsp. hominis		5.72	0.09		1	1	
306773	**Low discrimination**						**306773**
					COA	NOV	
Staphylococcus xylosus		45.19	0.34	VPT 14	1	99	
Staphylococcus aureus subsp. aureus		29.54	0.18	GLR 95	99	4	
Staphylococcus lutrae		12.58	0.18		99	1	
307101	**Good identification**						**307101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	ARG 99			
307103	**Very good identification**						**307103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	ARG 99			
307111	**Good identification**						**307111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	ARG 99			
307113	**Good identification**						**307113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	ARG 99			
307240	**Low discrimination**						**307240**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		69.33	0.27	ESL 8	99		
Staphylococcus epidermidis		25.46	0.26	PHS 1	1		
Staphylococcus hominis subsp. hominis		5.21	0.15		1		
307241	**Low discrimination**						**307241**
					NOV		
Staphylococcus epidermidis		73.17	0.32	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		26.83	0.15		99		
307242	**Low discrimination**						**307242**
					NOV		
Staphylococcus epidermidis		64.98	0.36	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		30.09	0.22		99		
Staphylococcus hominis subsp. hominis		4.93	0.17		1		
307243	**Acceptable identification**						**307243**
					NOV		
Staphylococcus epidermidis		94.13	0.42	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		5.87	0.09		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
307301	**Acceptable identification**						**307301**
					COA		
Staphylococcus schleiferi subsp. coagulans		90.02	0.27	ARG 99	99		
Staphylococcus epidermidis		9.98	0.06	LAC 86	1		
307303	**Acceptable identification**						**307303**
					COA		
Staphylococcus schleiferi subsp. coagulans		94.60	0.43	ARG 99	99		
Staphylococcus epidermidis		5.41	0.16		1		
307313	**Good identification**						**307313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ARG 99			
307340	**Low discrimination**						**307340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		58.12	0.37	ESL 8	99		
Staphylococcus epidermidis		40.31	0.42	PHS 1	1		
Staphylococcus hominis subsp. hominis		1.58	0.16		1		
307341	**Low discrimination**						**307341**
					NOV		
Staphylococcus epidermidis		83.75	0.47	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		16.25	0.25		99		
307342	**Low discrimination**						**307342**
					NOV		
Staphylococcus epidermidis		79.39	0.52	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		19.46	0.32		99		
Staphylococcus hominis subsp. hominis		1.15	0.18		1		
307343	**Very good identification**						**307343**
					COA	NOV	
Staphylococcus epidermidis		96.64	0.58	ESL 1	1	1	
Staphylococcus hominis subsp. novobiosepticus		3.19	0.19		1	99	
Staphylococcus schleiferi subsp. coagulans		0.17	0.02		99	1	
307450	**Low discrimination**						**307450**
Staphylococcus kloosii		60.11	0.27	VPT 10			
Staphylococcus saprophyticus subsp. bovis		39.89	0.06	NIT 1			
307452	**Good identification**						**307452**
Staphylococcus kloosii		100.0	0.28	VPT 10			
307453	**Good identification**						**307453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	GLR 99			
307552	**Low discrimination**						**307552**
					NOV	YEP	
Staphylococcus kloosii		53.46	0.33	VPT 10	99	99	
Staphylococcus caprae		29.67	0.00	NIT 1	1	1	
Staphylococcus saprophyticus subsp. bovis		16.87	0.06		99	80	
307642	**Acceptable identification**						**307642**
Staphylococcus hominis subsp. hominis		93.32	0.27	ESL 1			
Staphylococcus piscifermentans		6.68	0.11	PHS 3			
307742	**Low discrimination**						**307742**
Staphylococcus hominis subsp. hominis		75.81	0.28	ESL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		16.68	0.11	PHS 3				
Staphylococcus piscifermentans		7.51	0.14					
307750	**Acceptable identification**							**307750**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		94.28	0.31	ESL 1	99	99		
Staphylococcus hominis subsp. hominis		5.72	0.09		1	1		
307763	**Good identification**							**307763**
Staphylococcus gallinarum		96.75	0.32	VPT 1				
Staphylococcus equorum		1.63	-.02	MAN 91				
Staphylococcus xylosus		1.62	-.03					
307771	**Low discrimination**							**307771**
Staphylococcus gallinarum		82.58	0.45	VPT 1				
Staphylococcus equorum		12.99	0.31					
Staphylococcus xylosus		1.85	0.12					
307773	**Low discrimination**							**307773**
Staphylococcus gallinarum		82.80	0.53	VPT 1				
Staphylococcus equorum		13.68	0.39					
Staphylococcus xylosus		2.15	0.21					
310000	**Low discrimination**							**310000**
					OXI			
Dermacoccus nishinomiyaensis		66.96	0.33	VPT 5	93			
Micrococcus sp.		21.10	0.13	ORN 1	89			
Kocuria varians		11.94	0.25		13			
310100	**Low discrimination**							**310100**
					OXI			
Kocuria varians		58.41	0.29	VPT 14	13			
Dermacoccus nishinomiyaensis		41.59	0.18	ORN 1	93			
310240	**Acceptable identification**							**310240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		93.18	0.30	ORN 1	99			
Staphylococcus warneri		3.06	0.16	NIT 90	1			
Staphylococcus epidermidis		1.97	0.03		1			
310340	**Good identification**							**310340**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		95.52	0.39	ORN 1	99			
Staphylococcus epidermidis		3.82	0.18	NIT 90	1			
Staphylococcus hominis subsp. hominis		0.66	0.06		1			
310341	**Low discrimination**							**310341**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		72.41	0.27	ORN 1	1	99	1	
Staphylococcus epidermidis		21.51	0.24	NIT 90	1	1	1	
Kocuria kristinae		4.05	0.03		99	1	93	
310342	**Low discrimination**							**310342**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		79.98	0.34	ORN 1	99			
Staphylococcus epidermidis		18.81	0.29	NIT 90	1			
Staphylococcus hominis subsp. hominis		1.21	0.08		1			
310343	**Low discrimination**							**310343**
					NOV	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		54.35	0.34	ORN 12	1	1	
Staphylococcus hominis subsp. novobiosepticus		31.11	0.21	NIT 97	99	1	
Staphylococcus lugdunensis		14.54	0.18		1	96	
310641	**Low discrimination**						**310641**
					OXI		
Kocuria kristinae		53.26	0.39	URE 10	99		
Staphylococcus lugdunensis		46.22	0.39	ORN 1	1		
Staphylococcus petrasii subsp. petrasii		0.53	-.02		1		
310643	**Very good identification**						**310643**
					OXI		
Staphylococcus lugdunensis		96.90	0.59	NIT 99	1		
Kocuria kristinae		3.10	0.27		99		
310650	**Low discrimination**						**310650**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		72.71	0.28	ORN 1	99		
Staphylococcus warneri		25.29	0.39		1		
Staphylococcus pasteuri		2.00	0.02		1		
310652	**Good identification**						**310652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		97.81	0.43	ORN 1	99		
Staphylococcus warneri		2.00	0.29		1		
Staphylococcus pasteuri		0.19	-.06		1		
310740	**Low discrimination**						**310740**
					NOV		
Staphylococcus warneri		62.70	0.32	ORN 1	1		
Staphylococcus hominis subsp. hominis		20.97	0.16		1		
Staphylococcus hominis subsp. novobiosepticus		9.65	-.02		99		
310741	**Low discrimination**						**310741**
					OXI		
Staphylococcus lugdunensis		71.85	0.39	NIT 99	1		
Kocuria kristinae		27.60	0.30	LAC 90	99		
Staphylococcus petrasii subsp. petrasii		0.55	-.05		1		
310743	**Very good identification**						**310743**
					OXI		
Staphylococcus lugdunensis		98.94	0.59	NIT 99	1		
Kocuria kristinae		1.06	0.17		99		
310750	**Low discrimination**						**310750**
					NOV	PYR	
Staphylococcus warneri		51.75	0.29	ORN 1	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		40.97	0.06		99	7	
Staphylococcus saprophyticus subsp. bovis		7.29	0.02		99	99	
311641	**Good identification**						**311641**
					OXI		
Kocuria kristinae		96.61	0.27	URE 10	99		
Staphylococcus lugdunensis		3.39	-.02	ORN 1	1		
312000	**Low discrimination**						**312000**
					OXI		
Kocuria varians		57.85	0.39	VPT 14	13		
Dermacoccus nishinomiyaensis		37.37	0.28	ORN 1	93		
Kocuria rosea		3.67	0.05		35		
312002	**Good identification**						**312002**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Kocuria varians		100.0	0.28	VPT 14			
312020	**Acceptable identification**						**312020**
Kocuria varians		92.82	0.30	VPT 14			
Kocuria rosea		7.18	-.02	ORN 1			
312100	**Acceptable identification**						**312100**
					OXI		
Kocuria varians		92.42	0.43	VPT 14	13		
Dermacoccus nishinomiyaensis		7.58	0.13	ORN 1	93		
312102	**Good identification**						**312102**
Kocuria varians		100.0	0.31	VPT 14			
312120	**Good identification**						**312120**
Kocuria varians		100.0	0.33	VPT 14			
312140	**Low discrimination**						**312140**
					NOV		
Kocuria varians		49.00	0.29	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		41.80	0.18	ORN 1	99		
Staphylococcus epidermidis		6.01	0.08		1		
312240	**Low discrimination**						**312240**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		77.35	0.49	ORN 1	99		
Staphylococcus hominis subsp. hominis		16.34	0.47		1		
Staphylococcus epidermidis		5.88	0.34		1		
312241	**Low discrimination**						**312241**
					NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		39.92	0.37	ORN 1	99	1	
Staphylococcus lugdunensis		36.83	0.39		1	96	
Staphylococcus epidermidis		22.56	0.39		1	1	
312242	**Low discrimination**						**312242**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		51.37	0.44	ORN 1	99		
Staphylococcus hominis subsp. hominis		23.65	0.48		1		
Staphylococcus epidermidis		22.98	0.44		1		
312243	**Low discrimination**						**312243**
					NOV	PYR	
Staphylococcus lugdunensis		77.89	0.59	TRE 99	1	96	
Staphylococcus epidermidis		16.79	0.50		1	1	
Staphylococcus hominis subsp. novobiosepticus		5.05	0.31		99	1	
312250	**Low discrimination**						**312250**
					NOV		
Staphylococcus hominis subsp. hominis		75.76	0.30	ORN 1	1		
Staphylococcus hominis subsp. novobiosepticus		24.25	0.08	MAN 13	99		
312252	**Low discrimination**						**312252**
					NOV		
Staphylococcus hominis subsp. hominis		82.48	0.31	ORN 1	1		
Staphylococcus hominis subsp. novobiosepticus		12.11	0.03	MAN 13	99		
Staphylococcus epidermidis		5.42	0.03		1		
312340	**Low discrimination**						**312340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		81.81	0.59	ORN 1	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		11.76	0.49		1		
Staphylococcus hominis subsp. hominis		6.24	0.47		1		
312341	**Low discrimination**						**312341**
					NOV	PYR	
Staphylococcus epidermidis		44.83	0.55	ORN 12	1	1	
Staphylococcus hominis subsp. novobiosepticus		42.00	0.47		99	1	
Staphylococcus lugdunensis		12.92	0.39		1	96	
312342	**Low discrimination**						**312342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		49.40	0.53	ORN 1	99		
Staphylococcus epidermidis		41.74	0.60		1		
Staphylococcus hominis subsp. hominis		8.22	0.49		1		
312343	**Low discrimination**						**312343**
					NOV	PYR	
Staphylococcus epidermidis		50.50	0.65	ORN 12	1	1	
Staphylococcus lugdunensis		41.35	0.59		1	96	
Staphylococcus hominis subsp. novobiosepticus		8.05	0.41		99	1	
312350	**Low discrimination**						**312350**
					NOV		
Staphylococcus hominis subsp. hominis		48.10	0.30	ORN 1	1		
Staphylococcus hominis subsp. novobiosepticus		42.63	0.18	MAN 13	99		
Staphylococcus epidermidis		6.13	0.08		1		
312352	**Low discrimination**						**312352**
					NOV		
Staphylococcus hominis subsp. hominis		57.14	0.32	ORN 1	1		
Staphylococcus hominis subsp. novobiosepticus		23.23	0.12	MAN 13	99		
Staphylococcus epidermidis		19.63	0.19		1		
312441	**Good identification**						**312441**
Staphylococcus lugdunensis		100.0	0.39	SUC 99			
312443	**Very good identification**						**312443**
Staphylococcus lugdunensis		100.0	0.59	SUC 99			
312541	**Good identification**						**312541**
Staphylococcus lugdunensis		100.0	0.39	SUC 99			
312543	**Very good identification**						**312543**
Staphylococcus lugdunensis		100.0	0.59	SUC 99			
312550	**Good identification**						**312550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.31	ORN 1			
312601	**Good identification**						**312601**
					OXI		
Staphylococcus lugdunensis		99.05	0.39	MLT 99	1		
Kocuria kristinae		0.96	-.04	LAC 90	99		
312603	**Very good identification**						**312603**
Staphylococcus lugdunensis		100.0	0.59	MLT 99			
312640	**Low discrimination**						**312640**
					PYR		
Staphylococcus hominis subsp. hominis		70.36	0.57	ORN 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lugdunensis		24.26	0.39		96		
Staphylococcus warneri		3.12	0.35		8		
312641	**Very good identification**						**312641**
					OXI		
Staphylococcus lugdunensis		99.74	0.80	LAC 90	1		
Staphylococcus petrasii subsp. petrasii		0.10	0.18		1		
Kocuria kristinae		0.09	0.16		99		
312642	**Low discrimination**						**312642**
					PYR		
Staphylococcus lugdunensis		72.11	0.59	MNS 99	96		
Staphylococcus hominis subsp. hominis		27.28	0.58		1		
Staphylococcus warneri		0.34	0.25		8		
312643	**Excellent identification**						**312643**
					PYR		
Staphylococcus lugdunensis		99.98	1.00		96		
Staphylococcus hominis subsp. hominis		0.01	0.23		1		
Staphylococcus petrasii subsp. petrasii		0.01	0.12		99		
312650	**Low discrimination**						**312650**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		52.97	0.40	ORN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		29.26	0.31	MAN 13	99	99	
Staphylococcus warneri		11.40	0.32		1	8	
312651	**Acceptable identification**						**312651**
					NOV		
Staphylococcus lugdunensis		92.58	0.39	MAN 1	1		
Staphylococcus saprophyticus subsp. bovis		5.54	0.18	LAC 90	99		
Staphylococcus petrasii subsp. petrasii		1.06	-.02		1		
312652	**Low discrimination**						**312652**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		68.42	0.41	ORN 1	1	1	
Staphylococcus lugdunensis		12.23	0.18	MAN 13	1	96	
Staphylococcus saprophyticus subsp. bovis		8.05	0.18		99	99	
312653	**Very good identification**						**312653**
					NOV	PYR	
Staphylococcus lugdunensis		99.72	0.59	MAN 1	1	96	
Staphylococcus saprophyticus subsp. bovis		0.17	0.06		99	99	
Staphylococcus hominis subsp. hominis		0.12	0.06		1	1	
312661	**Good identification**						**312661**
Staphylococcus lugdunensis		100.0	0.39	XYL 1			
312663	**Very good identification**						**312663**
Staphylococcus lugdunensis		100.0	0.59	XYL 1			
312701	**Good identification**						**312701**
Staphylococcus lugdunensis		99.24	0.39	MLT 99			
Rothia mucilaginosa		0.76	-.09	LAC 90			
312703	**Very good identification**						**312703**
Staphylococcus lugdunensis		100.0	0.59	MLT 99			
312740	**Low discrimination**						**312740**
					NOV	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		71.09	0.58	ORN 1	1	1		
Staphylococcus lugdunensis		22.63	0.39		1	96		
Staphylococcus hominis subsp. novobiosepticus		2.97	0.18		99	1		
312741	**Very good identification**							**312741**
					PYR			
Staphylococcus lugdunensis		99.75	0.80	LAC 90	96			
Staphylococcus petrasii subsp. petrasii		0.07	0.14		99			
Staphylococcus hominis subsp. hominis		0.07	0.23		1			
312742	**Low discrimination**							**312742**
					OXI	PYR	NOV	
Staphylococcus lugdunensis		69.57	0.59	MNS 99	1	96	1	
Staphylococcus hominis subsp. hominis		28.51	0.59		1	1	1	
Macrococcus caseolyticus		0.81	0.14		99	80	99	
312743	**Excellent identification**							**312743**
					PYR			
Staphylococcus lugdunensis		99.97	1.00		96			
Staphylococcus epidermidis		0.01	0.24		1			
Staphylococcus hominis subsp. hominis		0.01	0.24		1			
312750	**Low discrimination**							**312750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		65.30	0.43	ORN 1	99	99		
Staphylococcus hominis subsp. hominis		32.01	0.41		1	1		
Staphylococcus warneri		2.01	0.21		1	8		
312751	**Low discrimination**							**312751**
					NOV	PYR		
Staphylococcus lugdunensis		79.58	0.39	MAN 1	1	96		
Staphylococcus saprophyticus subsp. bovis		19.05	0.31	LAC 90	99	99		
Staphylococcus hominis subsp. hominis		0.76	0.06		1	1		
312752	**Low discrimination**							**312752**
					NOV	PYR		
Staphylococcus hominis subsp. hominis		61.84	0.42	ORN 1	1	1		
Staphylococcus saprophyticus subsp. bovis		26.86	0.31	MAN 13	99	99		
Staphylococcus lugdunensis		10.20	0.18		1	96		
312753	**Very good identification**							**312753**
					NOV	PYR		
Staphylococcus lugdunensis		99.22	0.59	MAN 1	1	96		
Staphylococcus saprophyticus subsp. bovis		0.66	0.18		99	99		
Staphylococcus hominis subsp. hominis		0.12	0.07		1	1		
312761	**Good identification**							**312761**
Staphylococcus lugdunensis		100.0	0.39	XYL 1				
312763	**Very good identification**							**312763**
Staphylococcus lugdunensis		100.0	0.59	XYL 1				
313240	**Good identification**							**313240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		97.61	0.27	ORN 1	99			
Staphylococcus hominis subsp. hominis		2.40	0.05	ESL 8	1			
313340	**Good identification**							**313340**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		97.51	0.37	ORN 1	99			
Staphylococcus epidermidis		1.63	0.08	ESL 8	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus hominis subsp. hominis		0.86	0.06		1	
313342	**Low discrimination**					**313342**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		89.49	0.32	ORN 1	99	
Staphylococcus epidermidis		8.78	0.19	ESL 8	1	
Staphylococcus hominis subsp. hominis		1.73	0.08		1	
313641	**Good identification**					**313641**
					OXI	
Staphylococcus lugdunensis		97.88	0.39	ESL 1	1	
Kocuria kristinae		2.12	0.04	LAC 90	99	
313643	**Very good identification**					**313643**
Staphylococcus lugdunensis		100.0	0.59	ESL 1		
313741	**Low discrimination**					**313741**
					OXI	
Staphylococcus lugdunensis		85.62	0.39	ESL 1	1	
Rothia mucilaginosa		13.76	0.18	LAC 90	1	
Kocuria kristinae		0.62	-.06		99	
313743	**Very good identification**					**313743**
Staphylococcus lugdunensis		100.0	0.59	ESL 1		
314242	**Good identification**					**314242**
Staphylococcus epidermidis		100.0	0.29	ORN 12		
314243	**Good identification**					**314243**
Staphylococcus epidermidis		100.0	0.34	ORN 12		
314340	**Good identification**					**314340**
					NOV	
Staphylococcus epidermidis		95.73	0.34	ORN 12	1	
Staphylococcus hominis subsp. novobiosepticus		4.27	-.02	NIT 97	99	
314341	**Good identification**					**314341**
Staphylococcus epidermidis		100.0	0.39	ORN 12		
314342	**Good identification**					**314342**
					NOV	
Staphylococcus epidermidis		99.25	0.44	ORN 12	1	
Staphylococcus hominis subsp. novobiosepticus		0.75	-.07	NIT 97	99	
314343	**Good identification**					**314343**
Staphylococcus epidermidis		100.0	0.50	ORN 12		
314450	**Good identification**					**314450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
314452	**Good identification**					**314452**
Staphylococcus kloosii		100.0	0.28	VPT 10		
314550	**Good identification**					**314550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
314552	**Good identification**					**314552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.33	VPT 10			
315450	**Good identification**						**315450**
Staphylococcus kloosii		100.0	0.27	VPT 10			
315452	**Good identification**						**315452**
Staphylococcus kloosii		100.0	0.28	VPT 10			
315550	**Good identification**						**315550**
Staphylococcus kloosii		100.0	0.33	VPT 10			
315552	**Good identification**						**315552**
Staphylococcus kloosii		100.0	0.33	VPT 10			
316141	**Good identification**						**316141**
Staphylococcus epidermidis		100.0	0.29	ORN 12			
316142	**Good identification**						**316142**
Staphylococcus epidermidis		100.0	0.34	ORN 12			
316143	**Good identification**						**316143**
Staphylococcus epidermidis		100.0	0.40	ORN 12			
316240	**Good identification**						**316240**
					NOV		
Staphylococcus epidermidis		96.29	0.49	ORN 12	1		
Staphylococcus hominis subsp. novobiosepticus		2.26	0.08		99		
Staphylococcus hominis subsp. hominis		1.46	0.15		1		
316241	**Very good identification**						**316241**
					NOV	PYR	
Staphylococcus epidermidis		99.40	0.55	ORN 12	1	1	
Staphylococcus hominis subsp. novobiosepticus		0.31	-.04		99	1	
Staphylococcus lugdunensis		0.29	-.02		1	96	
316242	**Very good identification**						**316242**
					NOV		
Staphylococcus epidermidis		99.05	0.60	ORN 12	1		
Staphylococcus hominis subsp. hominis		0.56	0.17		1		
Staphylococcus hominis subsp. novobiosepticus		0.40	0.03		99		
316243	**Very good identification**						**316243**
					PYR		
Staphylococcus epidermidis		99.18	0.65	ORN 12	1		
Staphylococcus lugdunensis		0.82	0.18		96		
316301	**Good identification**						**316301**
Staphylococcus epidermidis		100.0	0.29	ORN 12			
316302	**Good identification**						**316302**
Staphylococcus epidermidis		100.0	0.34	ORN 12			
316303	**Good identification**						**316303**
					COA		
Staphylococcus epidermidis		98.66	0.40	ORN 12	1		
Staphylococcus schleiferi subsp. coagulans		1.35	0.02	MLT 99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
316340	**Very good identification**						**316340**
					NOV		
Staphylococcus epidermidis		98.49	0.65	ORN 12	1		
Staphylococcus hominis subsp. novobiosepticus		1.22	0.18		99		
Staphylococcus hominis subsp. hominis		0.29	0.16		1		
316341	**Very good identification**						**316341**
					NOV	PYR	
Staphylococcus epidermidis		99.78	0.71	ORN 12	1	1	
Staphylococcus hominis subsp. novobiosepticus		0.17	0.05		99	1	
Staphylococcus lugdunensis		0.05	-.02		1	96	
316342	**Very good identification**						**316342**
					NOV		
Staphylococcus epidermidis		99.68	0.75	ORN 12	1		
Staphylococcus hominis subsp. novobiosepticus		0.21	0.12		99		
Staphylococcus hominis subsp. hominis		0.11	0.18		1		
316343	**Very good identification**						**316343**
					NOV	PYR	
Staphylococcus epidermidis		99.83	0.81	ORN 12	1	1	
Staphylococcus lugdunensis		0.15	0.18		1	96	
Staphylococcus hominis subsp. novobiosepticus		0.03	0.00		99	1	
316351	**Good identification**						**316351**
Staphylococcus epidermidis		100.0	0.29	ORN 12			
316352	**Good identification**						**316352**
Staphylococcus epidermidis		100.0	0.34	ORN 12			
316353	**Good identification**						**316353**
Staphylococcus epidermidis		98.76	0.40	ORN 12			
Staphylococcus caprae		1.25	-.11	MAN 1			
316361	**Good identification**						**316361**
Staphylococcus epidermidis		100.0	0.29	ORN 12			
316362	**Good identification**						**316362**
Staphylococcus epidermidis		100.0	0.34	ORN 12			
316363	**Good identification**						**316363**
Staphylococcus epidermidis		100.0	0.40	ORN 12			
316641	**Good identification**						**316641**
					PYR		
Staphylococcus lugdunensis		96.61	0.39	PHS 1	96		
Staphylococcus epidermidis		3.39	0.14	LAC 90	1		
316642	**Low discrimination**						**316642**
					PYR		
Staphylococcus hominis subsp. hominis		41.13	0.27	ORN 1	1		
Staphylococcus lugdunensis		35.52	0.18	PHS 3	96		
Staphylococcus epidermidis		23.35	0.19		1		
316643	**Very good identification**						**316643**
					PYR		
Staphylococcus lugdunensis		98.78	0.59	PHS 1	96		
Staphylococcus epidermidis		1.22	0.24		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
316741	**Low discrimination**						**316741**
					PYR		
Staphylococcus lugdunensis		83.43	0.39	PHS 1	96		
Staphylococcus epidermidis		16.57	0.29	LAC 90	1		
316742	**Low discrimination**						**316742**
					PYR		
Staphylococcus epidermidis		62.30	0.34	ORN 12	1		
Staphylococcus hominis subsp. hominis		20.98	0.28	TRE 1	1		
Staphylococcus lugdunensis		16.72	0.18		96		
316743	**Good identification**						**316743**
					PYR	YEP	
Staphylococcus lugdunensis		93.38	0.59	PHS 1	96	50	
Staphylococcus epidermidis		6.53	0.40		1	1	
Staphylococcus chromogenes		0.10	-.02		45	95	
316750	**Acceptable identification**						**316750**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		94.28	0.31	ORN 1	99	99	
Staphylococcus hominis subsp. hominis		5.72	0.09		1	1	
317341	**Good identification**						**317341**
Staphylococcus epidermidis		100.0	0.29	ORN 12			
317342	**Good identification**						**317342**
Staphylococcus epidermidis		100.0	0.34	ORN 12			
317343	**Good identification**						**317343**
Staphylococcus epidermidis		100.0	0.40	ORN 12			
320000	**Low discrimination**						**320000**
					OXI		
Kocuria varians		81.74	0.56	VPT 14	13		
Dermacoccus nishinomiyaensis		13.89	0.33		93		
Micrococcus sp.		4.38	0.13		89		
320001	**Good identification**						**320001**
Kocuria varians		100.0	0.34	VPT 14			
320002	**Good identification**						**320002**
Kocuria varians		100.0	0.45	VPT 14			
320020	**Good identification**						**320020**
Kocuria varians		100.0	0.47	VPT 14			
320022	**Good identification**						**320022**
Kocuria varians		100.0	0.36	VPT 14			
320040	**Acceptable identification**						**320040**
					NOV		
Kocuria varians		93.54	0.43	VPT 14	1		
Staphylococcus warneri		2.78	0.18		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.97	-.06		99		
320042	**Low discrimination**						**320042**
					NOV		
Kocuria varians		68.59	0.31	VPT 14	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus saprophyticus subsp. saprophyticus		29.00	0.10		99	
Staphylococcus warneri		2.41	0.08		1	
320052	**Good identification**					**320052**
Staphylococcus saprophyticus subsp. saprophyticus		100.0	0.38	SUC 91		
320060	**Good identification**					**320060**
Kocuria varians		100.0	0.33	VPT 14		
320100	**Very good identification**					**320100**
					OXI	
Kocuria varians		97.89	0.60	VPT 14	13	
Dermacoccus nishinomiyaensis		2.11	0.18		93	
320101	**Good identification**					**320101**
Kocuria varians		100.0	0.38	VPT 14		
320102	**Good identification**					**320102**
Kocuria varians		100.0	0.49	VPT 14		
320103	**Good identification**					**320103**
Kocuria varians		100.0	0.27	VPT 14		
320120	**Good identification**					**320120**
Kocuria varians		100.0	0.50	VPT 14		
320121	**Good identification**					**320121**
Kocuria varians		100.0	0.29	VPT 14		
320122	**Good identification**					**320122**
Kocuria varians		100.0	0.39	VPT 14		
320140	**Good identification**					**320140**
					NOV	
Kocuria varians		98.62	0.46	VPT 14	1	
Staphylococcus hominis subsp. novobiosepticus		1.38	-.02		99	
320142	**Good identification**					**320142**
					NOV	
Kocuria varians		95.12	0.35	VPT 14	1	
Staphylococcus hominis subsp. novobiosepticus		2.55	-.07		99	
Staphylococcus saprophyticus subsp. saprophyticus		2.33	-.12		99	
320160	**Good identification**					**320160**
Kocuria varians		100.0	0.37	VPT 14		
320200	**Low discrimination**					**320200**
					OXI	
Kocuria varians		85.75	0.38	VPT 14	13	
Dermacoccus nishinomiyaensis		6.82	0.08	SUC 12	93	
Staphylococcus warneri		4.10	0.18		1	
320202	**Low discrimination**					**320202**
					NOV	
Kocuria varians		88.81	0.27	VPT 14	1	
Staphylococcus saprophyticus subsp. saprophyticus		6.17	-.11	SUC 12	99	
Staphylococcus warneri		5.02	0.08		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
320220	**Good identification**							**320220**
	Kocuria varians	100.0	0.29	VPT 14				
320240	**Low discrimination**							**320240**
					NOV			
	Staphylococcus hominis subsp. novobiosepticus	41.40	0.30	BGA 1	99			
	Staphylococcus warneri	33.63	0.45	NIT 90	1			
	Staphylococcus saprophyticus subsp. saprophyticus	12.69	0.15		99			
320241	**Acceptable identification**							**320241**
					OXI	NOV	PYR	
	Kocuria kristinae	92.11	0.44	URE 10	99	1	93	
	Staphylococcus hominis subsp. novobiosepticus	5.54	0.17	TRE 95	1	99	1	
	Staphylococcus epidermidis	1.60	0.14		1	1	1	
320242	**Low discrimination**							**320242**
					NOV			
	Staphylococcus saprophyticus subsp. saprophyticus	63.23	0.31	TRE 99	99			
	Staphylococcus hominis subsp. novobiosepticus	19.60	0.24	MAN 96	99			
	Staphylococcus warneri	9.86	0.35		1			
320243	**Low discrimination**							**320243**
					OXI	NOV	PYR	
	Kocuria kristinae	72.41	0.32	URE 10	99	1	93	
	Staphylococcus epidermidis	15.92	0.24	TRE 95	1	1	1	
	Staphylococcus hominis subsp. novobiosepticus	9.39	0.12		1	99	1	
320250	**Acceptable identification**							**320250**
					NOV			
	Staphylococcus saprophyticus subsp. saprophyticus	92.60	0.43	TRE 99	99			
	Staphylococcus warneri	7.40	0.42		1			
320252	**Very good identification**							**320252**
					NOV			
	Staphylococcus saprophyticus subsp. saprophyticus	99.53	0.59	TRE 99	99			
	Staphylococcus warneri	0.47	0.32		1			
320300	**Good identification**							**320300**
					NOV			
	Kocuria varians	97.79	0.42	VPT 14	1			
	Staphylococcus hominis subsp. novobiosepticus	2.21	-.02	SUC 12	99			
320302	**Good identification**							**320302**
					NOV			
	Kocuria varians	95.87	0.31	VPT 14	1			
	Staphylococcus hominis subsp. novobiosepticus	4.13	-.07	SUC 12	99			
320320	**Good identification**							**320320**
	Kocuria varians	100.0	0.33	VPT 14				
320340	**Low discrimination**							**320340**
					NOV			
	Staphylococcus hominis subsp. novobiosepticus	78.56	0.39	BGA 1	99			
	Kocuria varians	7.72	0.28	NIT 90	1			
	Staphylococcus warneri	6.72	0.34		1			
320341	**Low discrimination**							**320341**
					OXI	NOV	PYR	
	Kocuria kristinae	53.91	0.34	URE 10	99	1	93	
	Staphylococcus hominis subsp. novobiosepticus	29.20	0.27	TRE 95	1	99	1	
	Staphylococcus epidermidis	15.90	0.29		1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
320342	**Low discrimination**							**320342**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		60.82	0.34	BGA 1	99			
Staphylococcus epidermidis		26.23	0.34	NIT 90	1			
Staphylococcus saprophyticus subsp. saprophyticus		5.69	0.09		99			
320343	**Low discrimination**							**320343**
					OXI	NOV	PYR	
Staphylococcus epidermidis		63.30	0.40	NIT 97	1	1	1	
Staphylococcus hominis subsp. novobiosepticus		19.76	0.21		1	99	1	
Kocuria kristinae		16.94	0.22		99	1	93	
320352	**Good identification**							**320352**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		97.89	0.37	GAL 8	99			
Staphylococcus warneri		1.67	0.22	TRE 99	1			
Staphylococcus hominis subsp. novobiosepticus		0.44	-.07		99			
320400	**Low discrimination**							**320400**
					NOV			
Kocuria varians		84.88	0.33	VPT 14	1			
Staphylococcus warneri		7.35	0.18	TRE 7	1			
Staphylococcus saprophyticus subsp. saprophyticus		5.21	-.06		99			
320412	**Good identification**							**320412**
Staphylococcus saprophyticus subsp. saprophyticus		100.0	0.38	SUC 91				
320413	**Acceptable identification**							**320413**
Staphylococcus cohnii subsp. urealyticum		91.41	0.35	GLR 99				
Staphylococcus saprophyticus subsp. saprophyticus		8.59	-.03					
320440	**Low discrimination**							**320440**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		75.61	0.35	SUC 91	99			
Staphylococcus warneri		20.47	0.45	MAN 96	1			
Kocuria varians		2.73	0.20		1			
320441	**Low discrimination**							**320441**
					OXI	NOV		
Kocuria kristinae		86.67	0.30	URE 10	99	1		
Staphylococcus cohnii subsp. cohnii		5.56	0.15	SUC 99	1	99		
Staphylococcus saprophyticus subsp. saprophyticus		3.30	-.06		1	99		
320442	**Good identification**							**320442**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		98.26	0.51	SUC 91	99			
Staphylococcus warneri		1.56	0.35	MAN 96	1			
Kocuria varians		0.18	0.08		1			
320443	**Low discrimination**							**320443**
					OXI	NOV		
Staphylococcus cohnii subsp. urealyticum		40.10	0.30	GLR 99	1	99		
Kocuria kristinae		32.17	0.17	MAN 91	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		27.73	0.10		1	99		
320450	**Very good identification**							**320450**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		98.66	0.64	SUC 91	99	7		
Staphylococcus warneri		0.81	0.42		1	8		
Staphylococcus saprophyticus subsp. bovis		0.26	0.22		99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
320452	**Very good identification**							**320452**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		99.92	0.79	SUC 91	99			
Staphylococcus warneri		0.05	0.32		1			
Staphylococcus kloosii		0.01	0.28		99			
320453	**Low discrimination**							**320453**
Staphylococcus saprophyticus subsp. saprophyticus		62.14	0.38	SUC 91				
Staphylococcus cohnii subsp. urealyticum		37.86	0.51	MNS 1				
320472	**Good identification**							**320472**
Staphylococcus saprophyticus subsp. saprophyticus		100.0	0.38	SUC 91				
320500	**Good identification**							**320500**
Kocuria varians		100.0	0.37	VPT 14				
320513	**Good identification**							**320513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	GLR 99				
320520	**Good identification**							**320520**
Kocuria varians		100.0	0.27	VPT 14				
320542	**Acceptable identification**							**320542**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		91.84	0.29	GAL 8	99			
Staphylococcus warneri		5.31	0.24	SUC 91	1			
Kocuria varians		2.85	0.12	MAN 96	1			
320550	**Low discrimination**							**320550**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		85.50	0.42	GAL 8	99	7		
Staphylococcus saprophyticus subsp. bovis		10.18	0.34	SUC 91	99	99		
Staphylococcus warneri		2.54	0.31		1	8		
320552	**Very good identification**							**320552**
					PYR			
Staphylococcus saprophyticus subsp. saprophyticus		99.03	0.58	GAL 8	7			
Staphylococcus saprophyticus subsp. bovis		0.52	0.22	SUC 91	99			
Staphylococcus kloosii		0.28	0.33		52			
320553	**Low discrimination**							**320553**
					PYR			
Staphylococcus cohnii subsp. urealyticum		72.66	0.43	GLR 99	53			
Staphylococcus saprophyticus subsp. saprophyticus		24.20	0.16		7			
Staphylococcus saprophyticus subsp. bovis		3.14	0.09		99			
320600	**Low discrimination**							**320600**
					OXI	NOV	PYR	
Staphylococcus warneri		63.72	0.45	MLT 95	1	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		24.05	0.15		1	99	7	
Kocuria kristinae		6.95	0.10		99	1	93	
320601	**Good identification**							**320601**
					OXI	PYR		
Kocuria kristinae		99.81	0.51	URE 10	99	93		
Staphylococcus warneri		0.19	0.10	MLT 90	1	8		
320602	**Low discrimination**							**320602**
					OXI	NOV	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. saprophyticus		84.78	0.31	MAN 96	1	99	7	
Staphylococcus warneri		13.22	0.35	MLT 99	1	1	8	
Kocuria kristinae		1.08	-.03		99	1	93	
320603	**Good identification**							**320603**
					OXI	NOV		
Kocuria kristinae		97.89	0.39	URE 10	99	1		
Staphylococcus simulans		1.33	0.05	MLT 90	1	1		
Staphylococcus saprophyticus subsp. saprophyticus		0.78	-.11		1	99		
320610	**Acceptable identification**							**320610**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		92.60	0.43	MLT 99	99			
Staphylococcus warneri		7.40	0.42		1			
320612	**Very good identification**							**320612**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		99.45	0.59	MLT 99	99			
Staphylococcus warneri		0.47	0.32		1			
Staphylococcus simulans		0.09	0.07		1			
320640	**Low discrimination**							**320640**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		64.00	0.56	MAN 96	1	99	7	
Staphylococcus warneri		32.55	0.71		1	1	8	
Kocuria kristinae		1.68	0.30		99	1	93	
320641	**Very good identification**							**320641**
					OXI	PYR		
Kocuria kristinae		98.49	0.71	URE 10	99	93		
Staphylococcus petrasii subsp. petrasii		0.73	0.27		1	99		
Staphylococcus warneri		0.39	0.36		1	8		
320642	**Very good identification**							**320642**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		96.93	0.72	MAN 96	1	99	7	
Staphylococcus warneri		2.90	0.61		1	1	8	
Kocuria kristinae		0.11	0.17		99	1	93	
320643	**Low discrimination**							**320643**
					OXI	NOV	PYR	
Kocuria kristinae		89.95	0.58	URE 10	99	1	93	
Staphylococcus saprophyticus subsp. saprophyticus		7.92	0.31		1	99	7	
Staphylococcus petrasii subsp. petrasii		1.44	0.21		1	1	99	
320650	**Very good identification**							**320650**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		98.38	0.85		99	7		
Staphylococcus warneri		1.51	0.68		1	8		
Staphylococcus saprophyticus subsp. bovis		0.10	0.34		99	99		
320651	**Low discrimination**							**320651**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		66.13	0.43	MNS 1	1	99	7	
Kocuria kristinae		29.56	0.42		99	1	93	
Staphylococcus warneri		2.05	0.33		1	1	8	
320652	**Excellent identification**							**320652**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		99.91	1.00		99	7		
Staphylococcus warneri		0.09	0.58		1	8		
Staphylococcus saprophyticus subsp. bovis		0.01	0.22		99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
320653	**Very good identification**							**320653**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		97.37	0.59	MNS 1	1	99	7	
Kocuria kristinae		1.92	0.30		99	1	93	
Staphylococcus warneri		0.18	0.23		1	1	8	
320661	**Good identification**							**320661**
Kocuria kristinae		100.0	0.30	URE 10				
320662	**Good identification**							**320662**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		97.10	0.31	MAN 96	99			
Staphylococcus warneri		2.91	0.20	XYL 1	1			
320670	**Good identification**							**320670**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		98.49	0.43	XYL 1	99			
Staphylococcus warneri		1.51	0.27		1			
320672	**Very good identification**							**320672**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		99.91	0.59	XYL 1	99			
Staphylococcus warneri		0.09	0.17		1			
320700	**Low discrimination**							**320700**
					OXI	PYR		
Staphylococcus warneri		62.00	0.34	MLT 95	1	8		
Kocuria varians		24.42	0.19		13	10		
Kocuria kristinae		7.14	0.00		99	93		
320701	**Good identification**							**320701**
Kocuria kristinae		100.0	0.41	URE 10				
320703	**Good identification**							**320703**
Kocuria kristinae		100.0	0.29	URE 10				
320712	**Good identification**							**320712**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		98.32	0.37	GAL 8	99			
Staphylococcus warneri		1.68	0.22	MLT 99	1			
320740	**Low discrimination**							**320740**
					OXI	NOV	PYR	
Staphylococcus warneri		61.15	0.61		1	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		33.11	0.34		1	99	7	
Kocuria kristinae		3.34	0.20		99	1	93	
320741	**Very good identification**							**320741**
					OXI	PYR		
Kocuria kristinae		98.07	0.61	URE 10	99	93		
Staphylococcus petrasii subsp. petrasii		1.46	0.23		1	99		
Staphylococcus warneri		0.37	0.26		1	8		
320742	**Low discrimination**							**320742**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		89.21	0.50	GAL 8	99			
Staphylococcus warneri		9.69	0.51	MAN 96	1			
Staphylococcus hominis subsp. hominis		0.46	0.18		1			
320743	**Acceptable identification**							**320743**
					OXI	NOV	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Kocuria kristinae		93.70	0.48	URE 10	99	1	93	
Staphylococcus petrasii subsp. petrasii		3.00	0.18		1	1	99	
Staphylococcus saprophyticus subsp. saprophyticus		2.15	0.09		1	99	7	
320750	**Good identification**							**320750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		90.68	0.63	GAL 8	99	7		
Staphylococcus warneri		5.05	0.58		1	8		
Staphylococcus saprophyticus subsp. bovis		4.27	0.47		99	99		
320751	**Low discrimination**							**320751**
					OXI	PYR	NOV	
Kocuria kristinae		41.31	0.32	URE 10	99	93	1	
Staphylococcus saprophyticus subsp. bovis		28.09	0.34	MAN 4	1	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		24.10	0.22		1	7	99	
320752	**Very good identification**							**320752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		99.47	0.78	GAL 8	99	7		
Staphylococcus warneri		0.33	0.48		1	8		
Staphylococcus saprophyticus subsp. bovis		0.21	0.34		99	99		
320753	**Low discrimination**							**320753**
					OXI	PYR	NOV	
Staphylococcus saprophyticus subsp. saprophyticus		83.09	0.37	GAL 8	1	7	99	
Kocuria kristinae		6.28	0.20	MNS 1	99	93	1	
Staphylococcus saprophyticus subsp. bovis		4.27	0.22		1	99	99	
320771	**Good identification**							**320771**
Staphylococcus xylosus		100.0	0.29	VPT 14				
320772	**Good identification**							**320772**
Staphylococcus saprophyticus subsp. saprophyticus		99.48	0.37	GAL 8				
Staphylococcus xylosus		0.52	-.02	XYL 1				
320773	**Good identification**							**320773**
Staphylococcus xylosus		98.08	0.39	VPT 14				
Staphylococcus saprophyticus subsp. saprophyticus		1.92	-.04	GLR 95				
321241	**Good identification**							**321241**
					OXI	NOV		
Kocuria kristinae		97.95	0.32	URE 10	99	1		
Staphylococcus hominis subsp. novobiosepticus		2.05	-.05	TRE 95	1	99		
321413	**Good identification**							**321413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	GLR 99				
321452	**Good identification**							**321452**
Staphylococcus saprophyticus subsp. saprophyticus		98.79	0.38	ESL 1				
Staphylococcus kloosii		1.22	0.28	SUC 91				
321453	**Acceptable identification**							**321453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		94.50	0.42	GLR 99	99	53		
Staphylococcus saprophyticus subsp. saprophyticus		4.03	-.03		99	7		
Aerococcus viridans		1.48	0.16		1	95		
321550	**Low discrimination**							**321550**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		43.98	0.33	VPT 10	1			
Staphylococcus saprophyticus subsp. saprophyticus		30.62	0.01	PHS 99	1			
Macrococcus equipersicus		25.40	0.10		99			
321553	**Good identification**							**321553**
					NOV			
Staphylococcus cohnii subsp. urealyticum		96.49	0.35	GLR 99	99			
Aerococcus viridans		3.51	0.16		1			
321601	**Good identification**							**321601**
Kocuria kristinae		100.0	0.39	URE 10				
321641	**Very good identification**							**321641**
Kocuria kristinae		100.0	0.58	URE 10				
321642	**Acceptable identification**							**321642**
					OXI	NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		94.46	0.31	ESL 1	1	99	7	
Staphylococcus warneri		2.83	0.20	MAN 96	1	1	8	
Kocuria kristinae		2.71	0.05		99	1	93	
321643	**Good identification**							**321643**
					OXI	NOV		
Kocuria kristinae		99.65	0.46	URE 10	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		0.35	-.11		1	99		
321650	**Good identification**							**321650**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		98.49	0.43	ESL 1	99			
Staphylococcus warneri		1.51	0.27		1			
321651	**Acceptable identification**							**321651**
					OXI	NOV		
Kocuria kristinae		91.71	0.30	URE 10	99	1		
Staphylococcus saprophyticus subsp. saprophyticus		8.29	0.02	MAN 4	1	99		
321652	**Very good identification**							**321652**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		99.91	0.59	ESL 1	99			
Staphylococcus warneri		0.09	0.17		1			
321701	**Good identification**							**321701**
					OXI			
Kocuria kristinae		95.34	0.29	URE 10	99			
Rothia mucilaginosa		4.66	-.02	MLT 90	1			
321741	**Good identification**							**321741**
					OXI			
Kocuria kristinae		98.75	0.48	URE 10	99			
Rothia mucilaginosa		1.25	0.05		1			
321743	**Good identification**							**321743**
Kocuria kristinae		100.0	0.36	URE 10				
321752	**Good identification**							**321752**
Staphylococcus saprophyticus subsp. saprophyticus		100.0	0.37	ESL 1				
322000	**Very good identification**							**322000**
					OXI			
Kocuria varians		97.84	0.70	VPT 14	13			
Dermacoccus nishinomiyaensis		1.92	0.28		93			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Kocuria rosea		0.19	0.05		35	
322001	**Good identification**					**322001**
Kocuria varians		100.0	0.49	VPT 14		
322002	**Very good identification**					**322002**
Kocuria varians		100.0	0.59	VPT 14		
322003	**Good identification**					**322003**
Kocuria varians		100.0	0.37	VPT 14		
322010	**Good identification**					**322010**
Kocuria varians		100.0	0.29	VPT 14		
322020	**Very good identification**					**322020**
Kocuria varians		99.77	0.61	VPT 14		
Kocuria rosea		0.23	-.02			
322021	**Good identification**					**322021**
Kocuria varians		100.0	0.39	VPT 14		
322022	**Good identification**					**322022**
Kocuria varians		100.0	0.50	VPT 14		
322023	**Good identification**					**322023**
Kocuria varians		100.0	0.28	VPT 14		
322040	**Very good identification**					**322040**
					NOV	
Kocuria varians		98.21	0.57	VPT 14	1	
Staphylococcus hominis subsp. novobiosepticus		1.27	0.08		99	
Staphylococcus hominis subsp. hominis		0.27	0.05		1	
322041	**Acceptable identification**					**322041**
					NOV	
Kocuria varians		92.97	0.35	VPT 14	1	
Staphylococcus epidermidis		3.58	0.04	MNS 8	1	
Staphylococcus hominis subsp. novobiosepticus		3.46	-.04		99	
322042	**Acceptable identification**					**322042**
					NOV	
Kocuria varians		94.67	0.46	VPT 14	1	
Staphylococcus hominis subsp. novobiosepticus		2.34	0.03		99	
Staphylococcus epidermidis		1.92	0.09		1	
322060	**Good identification**					**322060**
Kocuria varians		100.0	0.47	VPT 14		
322062	**Good identification**					**322062**
Kocuria varians		100.0	0.36	VPT 14		
322100	**Very good identification**					**322100**
					OXI	
Kocuria varians		99.75	0.74	VPT 14	13	
Dermacoccus nishinomiyaensis		0.25	0.13		93	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
322101	**Good identification**						**322101**
					COA		
Kocuria varians		99.59	0.52	VPT 14	1		
Staphylococcus schleiferi subsp. coagulans		0.41	0.02	MNS 8	99		
322102	**Very good identification**						**322102**
Kocuria varians		100.0	0.63	VPT 14			
322103	**Acceptable identification**						**322103**
					COA		
Kocuria varians		91.84	0.41	VPT 14	1		
Staphylococcus schleiferi subsp. coagulans		8.16	0.18	MNS 8	99		
322110	**Good identification**						**322110**
Kocuria varians		100.0	0.33	VPT 14			
322120	**Very good identification**						**322120**
Kocuria varians		100.0	0.65	VPT 14			
322121	**Good identification**						**322121**
Kocuria varians		100.0	0.43	VPT 14			
322122	**Very good identification**						**322122**
Kocuria varians		100.0	0.53	VPT 14			
322123	**Good identification**						**322123**
Kocuria varians		100.0	0.32	VPT 14			
322140	**Very good identification**						**322140**
					NOV		
Kocuria varians		96.65	0.60	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		2.50	0.18		99		
Staphylococcus epidermidis		0.66	0.14		1		
322141	**Low discrimination**						**322141**
					NOV		
Kocuria varians		81.98	0.39	VPT 14	1		
Staphylococcus epidermidis		11.92	0.19	MNS 8	1		
Staphylococcus hominis subsp. novobiosepticus		6.09	0.05		99		
322142	**Low discrimination**						**322142**
					NOV		
Kocuria varians		88.18	0.49	VPT 14	1		
Staphylococcus epidermidis		6.74	0.24		1		
Staphylococcus hominis subsp. novobiosepticus		4.35	0.12		99		
322143	**Low discrimination**						**322143**
					NOV		
Staphylococcus epidermidis		58.85	0.30	SUC 99	1		
Kocuria varians		36.04	0.27		1		
Staphylococcus hominis subsp. novobiosepticus		5.11	0.00		99		
322150	**Low discrimination**						**322150**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		87.49	0.34	TRE 99	99	99	
Kocuria varians		12.51	0.19		1	10	
322160	**Good identification**						**322160**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Kocuria varians		100.0	0.51	VPT 14				
322161	**Good identification**							**322161**
Kocuria varians		100.0	0.29	VPT 14				
322162	**Good identification**							**322162**
Kocuria varians		100.0	0.40	VPT 14				
322200	**Good identification**							**322200**
					OXI	NOV		
Kocuria varians		96.29	0.53	VPT 14	13	1		
Staphylococcus hominis subsp. novobiosepticus		2.00	0.08	SUC 12	1	99		
Dermacoccus nishinomiyaensis		0.88	0.03		93	80		
322201	**Low discrimination**							**322201**
					OXI	PYR		
Kocuria varians		82.42	0.31	VPT 14	13	10		
Kocuria kristinae		7.54	0.01	SUC 12	99	93		
Staphylococcus epidermidis		5.11	0.04	MNS 8	1	1		
322202	**Acceptable identification**							**322202**
					NOV			
Kocuria varians		91.69	0.41	VPT 14	1			
Staphylococcus hominis subsp. novobiosepticus		3.64	0.03	SUC 12	99			
Staphylococcus epidermidis		2.99	0.09		1			
322220	**Good identification**							**322220**
Kocuria varians		100.0	0.43	VPT 14				
322222	**Good identification**							**322222**
Kocuria varians		100.0	0.32	VPT 14				
322240	**Low discrimination**							**322240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		66.85	0.49	BGA 1	99			
Staphylococcus hominis subsp. hominis		14.12	0.47		1			
Staphylococcus epidermidis		9.32	0.39		1			
322241	**Low discrimination**							**322241**
					OXI	NOV	PYR	
Staphylococcus epidermidis		45.63	0.45		1	1	1	
Staphylococcus hominis subsp. novobiosepticus		44.04	0.37		1	99	1	
Kocuria kristinae		6.12	0.21		99	1	93	
322242	**Low discrimination**							**322242**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		42.30	0.44	BGA 1	99			
Staphylococcus epidermidis		34.69	0.50		1			
Staphylococcus hominis subsp. hominis		19.48	0.48		1			
322243	**Low discrimination**							**322243**
					NOV			
Staphylococcus epidermidis		80.90	0.55		1			
Staphylococcus hominis subsp. novobiosepticus		13.28	0.31		99			
Staphylococcus capitis subsp. ureolyticus		3.20	0.13		1			
322250	**Low discrimination**							**322250**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		51.52	0.34	TRE 99	99	99		
Staphylococcus hominis subsp. hominis		19.63	0.30		1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		17.42	0.34		1	8	
322253	**Good identification**						**322253**
Staphylococcus capitis subsp. ureolyticus		97.37	0.41	ARG 99			
Staphylococcus epidermidis		1.04	0.14	BGA 12			
Staphylococcus simulans		0.94	0.13				
322260	**Low discrimination**						**322260**
					NOV		
Kocuria varians		75.37	0.30	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		20.34	0.08	SUC 12	99		
Staphylococcus hominis subsp. hominis		4.30	0.05		1		
322300	**Good identification**						**322300**
					NOV		
Kocuria varians		94.72	0.56	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		3.94	0.18	SUC 12	99		
Staphylococcus epidermidis		1.04	0.14		1		
322301	**Low discrimination**						**322301**
					NOV		
Kocuria varians		73.87	0.34	VPT 14	1		
Staphylococcus epidermidis		17.29	0.19	SUC 12	1		
Staphylococcus hominis subsp. novobiosepticus		8.84	0.05	MNS 8	99		
322302	**Low discrimination**						**322302**
					NOV		
Kocuria varians		82.25	0.45	VPT 14	1		
Staphylococcus epidermidis		10.12	0.24	SUC 12	1		
Staphylococcus hominis subsp. novobiosepticus		6.54	0.12		99		
322303	**Low discrimination**						**322303**
					NOV		
Staphylococcus epidermidis		64.70	0.30	MLT 99	1		
Kocuria varians		24.61	0.23		1		
Staphylococcus hominis subsp. novobiosepticus		5.62	0.00		99		
322320	**Good identification**						**322320**
Kocuria varians		100.0	0.47	VPT 14			
322322	**Good identification**						**322322**
Kocuria varians		100.0	0.36	VPT 14			
322340	**Low discrimination**						**322340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		71.49	0.59	BGA 1	99		
Staphylococcus epidermidis		18.83	0.55		1		
Staphylococcus hominis subsp. hominis		5.45	0.47		1		
322341	**Low discrimination**						**322341**
					NOV		
Staphylococcus epidermidis		65.18	0.60		1		
Staphylococcus hominis subsp. novobiosepticus		33.31	0.47		99		
Kocuria varians		0.62	0.21		1		
322342	**Low discrimination**						**322342**
					NOV		
Staphylococcus epidermidis		56.42	0.65		1		
Staphylococcus hominis subsp. novobiosepticus		36.42	0.53		99		
Staphylococcus hominis subsp. hominis		6.06	0.49		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
322343	**Good identification**						**322343**
					NOV		
Staphylococcus epidermidis		91.65	0.71		1		
Staphylococcus hominis subsp. novobiosepticus		7.96	0.41		99		
Staphylococcus hominis subsp. hominis		0.11	0.14		1		
322350	**Low discrimination**						**322350**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		79.98	0.47	TRE 99	99	99	
Staphylococcus hominis subsp. hominis		8.25	0.30		1	1	
Staphylococcus hominis subsp. novobiosepticus		7.32	0.18		1	99	
322351	**Low discrimination**						**322351**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		77.14	0.34	TRE 99	99	99	
Staphylococcus epidermidis		13.81	0.19		1	1	
Staphylococcus hominis subsp. novobiosepticus		7.06	0.05		1	99	
322352	**Low discrimination**						**322352**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		48.27	0.34	TRE 99	99	99	
Staphylococcus hominis subsp. hominis		23.39	0.32		1	1	
Staphylococcus epidermidis		14.73	0.24		1	1	
322353	**Low discrimination**						**322353**
					PYR	NOV	
Staphylococcus epidermidis		52.49	0.30	MAN 1	1	1	
Staphylococcus saprophyticus subsp. bovis		23.15	0.22		99	99	
Staphylococcus capitis subsp. ureolyticus		8.78	0.00		1	1	
322360	**Low discrimination**						**322360**
					NOV		
Kocuria varians		58.04	0.33	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		31.32	0.18	SUC 12	99		
Staphylococcus epidermidis		8.25	0.14		1		
322363	**Low discrimination**						**322363**
					NOV		
Staphylococcus epidermidis		84.58	0.30	XYL 1	1		
Staphylococcus xylosus		8.07	0.04		99		
Staphylococcus hominis subsp. novobiosepticus		7.35	0.00		99		
322373	**Good identification**						**322373**
Staphylococcus xylosus		100.0	0.29	VPT 14			
322400	**Good identification**						**322400**
Kocuria varians		97.15	0.47	VPT 14			
Staphylococcus auricularis		2.11	0.12	TRE 7			
Staphylococcus warneri		0.74	0.11				
322402	**Good identification**						**322402**
Kocuria varians		100.0	0.36	VPT 14			
322420	**Good identification**						**322420**
Kocuria varians		100.0	0.38	VPT 14			
322422	**Good identification**						**322422**
Kocuria varians		100.0	0.27	VPT 14			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
322440	**Low discrimination**							**322440**
					NOV	PYR		
Kocuria varians		41.44	0.34	VPT 14	1	10		
Staphylococcus warneri		27.25	0.37	TRE 7	1	8		
Staphylococcus saprophyticus subsp. bovis		14.59	0.22		99	99		
322450	**Good identification**							**322450**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		94.96	0.63		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		3.75	0.23		99	7		
Staphylococcus warneri		1.30	0.34		1	8		
322451	**Good identification**							**322451**
Staphylococcus saprophyticus subsp. bovis		99.06	0.50					
Staphylococcus cohnii subsp. urealyticum		0.94	0.14					
322452	**Low discrimination**							**322452**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		52.21	0.50		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		46.66	0.38		99	7		
Staphylococcus warneri		0.95	0.24		1	8		
322453	**Low discrimination**							**322453**
					NOV			
Staphylococcus saprophyticus subsp. bovis		68.17	0.38		99			
Staphylococcus cohnii subsp. urealyticum		26.21	0.35		99			
Staphylococcus simulans		3.17	0.11		1			
322500	**Good identification**							**322500**
Kocuria varians		100.0	0.51	VPT 14				
322501	**Acceptable identification**							**322501**
Kocuria varians		92.27	0.29	VPT 14				
Staphylococcus schleiferi subsp. schleiferi		7.73	0.02	TRE 7				
322502	**Good identification**							**322502**
Kocuria varians		100.0	0.40	VPT 14				
322510	**Good identification**							**322510**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		95.33	0.34	MLT 99	99	99		
Kocuria varians		4.67	0.10		1	10		
322520	**Good identification**							**322520**
Kocuria varians		100.0	0.42	VPT 14				
322522	**Good identification**							**322522**
Kocuria varians		100.0	0.30	VPT 14				
322540	**Low discrimination**							**322540**
					NOV	OXI	PYR	
Kocuria varians		41.11	0.37	VPT 14	1	13	10	
Staphylococcus saprophyticus subsp. bovis		38.59	0.34	TRE 7	99	1	99	
Macrococcus caseolyticus		11.20	0.14		99	99	80	
322550	**Very good identification**							**322550**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		99.81	0.75		99	99		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		0.11	0.24		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		0.09	0.01		99	7	
322551	**Very good identification**						**322551**
Staphylococcus saprophyticus subsp. bovis		99.77	0.63				
Staphylococcus xylosus		0.13	0.00				
Staphylococcus cohnii subsp. urealyticum		0.10	0.07				
322552	**Very good identification**						**322552**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		97.96	0.63		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		1.90	0.16		99	7	
Staphylococcus warneri		0.14	0.14		1	8	
322553	**Acceptable identification**						**322553**
Staphylococcus saprophyticus subsp. bovis		94.68	0.50				
Staphylococcus cohnii subsp. urealyticum		3.90	0.28				
Staphylococcus xylosus		1.42	0.09				
322560	**Good identification**						**322560**
Kocuria varians		100.0	0.28	VPT 14			
322570	**Good identification**						**322570**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	XYL 1			
322573	**Low discrimination**						**322573**
					COA	NOV	
Staphylococcus xylosus		69.87	0.31	VPT 14	1	99	
Staphylococcus lutrae		26.05	0.18	GLR 95	99	1	
Staphylococcus saprophyticus subsp. bovis		4.08	0.09	SUC 95	1	99	
322600	**Low discrimination**						**322600**
Staphylococcus warneri		44.28	0.37	MLT 95			
Kocuria varians		41.83	0.29				
Staphylococcus hominis subsp. hominis		7.73	0.16				
322601	**Low discrimination**						**322601**
					OXI	PYR	
Kocuria kristinae		88.96	0.28	URE 10	99	93	
Staphylococcus petrasii subsp. petrasii		7.19	0.05	NIT 7	1	99	
Kocuria varians		3.85	0.08	MLT 90	13	10	
322603	**Low discrimination**						**322603**
					OXI		
Staphylococcus simulans		74.80	0.34	VPT 4	1		
Kocuria kristinae		17.26	0.15	ARG 92	99		
Staphylococcus lugdunensis		4.94	0.05		1		
322610	**Low discrimination**						**322610**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		69.55	0.34	MLT 99	99	99	
Staphylococcus warneri		23.52	0.34		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		6.93	0.02		99	7	
322612	**Low discrimination**						**322612**
					NOV	PYR	
Staphylococcus simulans		55.27	0.35	VPT 4	1	70	
Staphylococcus saprophyticus subsp. saprophyticus		27.24	0.18	ARG 92	99	7	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		12.06	0.22		99	99	
322613	**Good identification**						**322613**
					NOV		
Staphylococcus simulans		98.40	0.46	VPT 4	1		
Staphylococcus saprophyticus subsp. bovis		1.60	0.09	ARG 92	99		
322640	**Low discrimination**						**322640**
					NOV	PYR	
Staphylococcus warneri		44.50	0.63		1	8	
Staphylococcus hominis subsp. hominis		40.50	0.57		1	1	
Staphylococcus saprophyticus subsp. bovis		5.02	0.34		99	99	
322641	**Low discrimination**						**322641**
					OXI		
Kocuria kristinae		48.57	0.47	URE 10	99		
Staphylococcus petrasii subsp. petrasii		43.18	0.47	NIT 7	1		
Staphylococcus lugdunensis		4.24	0.27		1		
322642	**Low discrimination**						**322642**
					NOV		
Staphylococcus hominis subsp. hominis		59.50	0.58	BGA 1	1		
Staphylococcus warneri		18.56	0.53		1		
Staphylococcus saprophyticus subsp. saprophyticus		14.62	0.31		99		
322643	**Low discrimination**						**322643**
					OXI		
Staphylococcus lugdunensis		48.40	0.47	ORN 99	1		
Staphylococcus petrasii subsp. petrasii		29.48	0.41		1		
Kocuria kristinae		15.39	0.35		99		
322650	**Low discrimination**						**322650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		84.73	0.75		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.45	0.43		99	7	
Staphylococcus warneri		5.50	0.60		1	8	
322651	**Good identification**						**322651**
					OXI	NOV	
Staphylococcus saprophyticus subsp. bovis		94.52	0.63		1	99	
Staphylococcus petrasii subsp. petrasii		3.01	0.27		1	1	
Kocuria kristinae		1.27	0.19		99	1	
322652	**Low discrimination**						**322652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		65.85	0.59	NIT 1	99	7	
Staphylococcus saprophyticus subsp. bovis		29.15	0.63		99	99	
Staphylococcus warneri		2.52	0.51		1	8	
322653	**Low discrimination**						**322653**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		65.90	0.50		99	99	
Staphylococcus simulans		18.36	0.39		1	70	
Staphylococcus saprophyticus subsp. saprophyticus		6.02	0.18		99	7	
322670	**Low discrimination**						**322670**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		85.87	0.34	XYL 1	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.56	0.02		99	7	
Staphylococcus warneri		5.57	0.19		1	8	
322671	**Low discrimination**						**322671**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		73.87	0.27	VPT 14			
Staphylococcus saprophyticus subsp. bovis		26.13	0.22	GLR 95			
322673	**Good identification**						**322673**
Staphylococcus xylosus		96.99	0.37	VPT 14			
Staphylococcus saprophyticus subsp. bovis		3.01	0.09	GLR 95			
322700	**Low discrimination**						**322700**
Kocuria varians		59.41	0.33	VPT 14			
Rothia mucilaginosa		15.79	0.10	SUC 12			
Staphylococcus warneri		13.24	0.27	TRE 7			
322710	**Good identification**						**322710**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		97.40	0.47	MLT 99	99	99	
Staphylococcus warneri		2.60	0.24		1	8	
322711	**Low discrimination**						**322711**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		89.91	0.34	MLT 99	1	99	
Staphylococcus intermedius		5.88	0.08		99	1	
Staphylococcus xylosus		4.21	0.03		1	99	
322712	**Low discrimination**						**322712**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		84.43	0.34	MLT 99	99	99	
Staphylococcus simulans		8.42	0.14		1	70	
Staphylococcus saprophyticus subsp. saprophyticus		4.15	-.04		99	7	
322733	**Good identification**						**322733**
Staphylococcus xylosus		100.0	0.34	VPT 14			
322740	**Low discrimination**						**322740**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		49.66	0.58	BGA 1	1	1	
Staphylococcus saprophyticus subsp. bovis		22.71	0.47		99	99	
Staphylococcus warneri		15.91	0.53		1	8	
322741	**Low discrimination**						**322741**
					OXI		
Staphylococcus petrasii subsp. petrasii		43.38	0.43	ARG 99	1		
Kocuria kristinae		24.40	0.38		99		
Rothia mucilaginosa		11.30	0.27		1		
322742	**Low discrimination**						**322742**
					NOV	PYR	
Staphylococcus hominis subsp. hominis		74.77	0.59	BGA 1	1	1	
Staphylococcus saprophyticus subsp. bovis		7.28	0.34		99	99	
Staphylococcus warneri		6.80	0.43		1	8	
322743	**Low discrimination**						**322743**
					OXI		
Staphylococcus lugdunensis		57.36	0.47	ORN 99	1		
Staphylococcus petrasii subsp. petrasii		23.29	0.37		1		
Kocuria kristinae		6.08	0.25		99		
322750	**Very good identification**						**322750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.90	0.88		99	99	
Staphylococcus warneri		0.51	0.50		1	8	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. hominis		0.33	0.41		1	1		
322751	**Very good identification**							**322751**
					NOV			
Staphylococcus saprophyticus subsp. bovis		98.58	0.75		99			
Staphylococcus xylosus		0.62	0.26		99			
Staphylococcus petrasii subsp. petrasii		0.52	0.23		1			
322752	**Good identification**							**322752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		93.19	0.75		99	99		
Staphylococcus saprophyticus subsp. saprophyticus		4.58	0.37		99	7		
Staphylococcus hominis subsp. hominis		1.44	0.42		1	1		
322753	**Low discrimination**							**322753**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		88.94	0.63		1	99		
Staphylococcus xylosus		6.35	0.36		1	99		
Staphylococcus intermedius		2.17	0.28		99	1		
322763	**Low discrimination**							**322763**
					COA	NOV	PYR	
Staphylococcus xylosus		81.50	0.33	VPT 14	1	99	65	
Staphylococcus lutrae		12.91	0.12	GLR 95	99	1	10	
Staphylococcus lugdunensis		3.97	0.05	MAN 94	1	1	96	
322770	**Good identification**							**322770**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		97.74	0.47	XYL 1	99	99		
Staphylococcus xylosus		1.76	0.07		99	65		
Staphylococcus warneri		0.50	0.09		1	8		
322771	**Low discrimination**							**322771**
Staphylococcus xylosus		87.72	0.48	VPT 14				
Staphylococcus saprophyticus subsp. bovis		12.28	0.34	GLR 95				
322772	**Low discrimination**							**322772**
					PYR			
Staphylococcus saprophyticus subsp. bovis		79.71	0.34	XYL 1	99			
Staphylococcus xylosus		16.37	0.16		65			
Staphylococcus saprophyticus subsp. saprophyticus		3.92	-.04		7			
322773	**Very good identification**							**322773**
					COA	NOV		
Staphylococcus xylosus		96.91	0.57	VPT 14	1	99		
Staphylococcus lutrae		1.90	0.18	GLR 95	99	1		
Staphylococcus saprophyticus subsp. bovis		1.19	0.22		1	99		
323000	**Good identification**							**323000**
Kocuria varians		100.0	0.29	VPT 14				
323100	**Good identification**							**323100**
Kocuria varians		100.0	0.33	VPT 14				
323240	**Good identification**							**323240**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		97.61	0.27	BGA 1	99			
Staphylococcus hominis subsp. hominis		2.40	0.05	ESL 8	1			
323340	**Acceptable identification**							**323340**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. novobiosepticus		94.56	0.37	BGA 1	99		
Staphylococcus epidermidis		2.89	0.14	ESL 8	1		
Rothia mucilaginosa		1.12	-.04		1		
323342	**Low discrimination**						**323342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		83.38	0.32	BGA 1	99		
Staphylococcus epidermidis		15.01	0.24	ESL 8	1		
Staphylococcus hominis subsp. hominis		1.61	0.08		1		
323343	**Low discrimination**						**323343**
					NOV		
Staphylococcus epidermidis		57.21	0.30	ESL 1	1		
Staphylococcus hominis subsp. novobiosepticus		42.79	0.19		99		
323453	**Good identification**						**323453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	GLR 99			
323550	**Good identification**						**323550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ESL 1			
323641	**Acceptable identification**						**323641**
					OXI		
Kocuria kristinae		91.57	0.35	URE 10	99		
Rothia mucilaginosa		5.14	0.05	NIT 7	1		
Staphylococcus petrasii subsp. petrasii		3.29	0.05		1		
323650	**Low discrimination**						**323650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		85.87	0.34	ESL 1	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.56	0.02		99	7	
Staphylococcus warneri		5.57	0.19		1	8	
323700	**Good identification**						**323700**
Rothia mucilaginosa		100.0	0.29	URE 1			
323701	**Good identification**						**323701**
					OXI		
Rothia mucilaginosa		98.47	0.39	URE 1	1		
Kocuria kristinae		1.53	0.06		99		
323713	**Acceptable identification**						**323713**
					OXI		
Staphylococcus lentus		93.10	0.30	VPT 7	99		
Staphylococcus xylosus		6.90	0.00	URE 1	1		
323733	**Low discrimination**						**323733**
					OXI		
Staphylococcus lentus		47.04	0.30	VPT 7	99		
Staphylococcus xylosus		40.08	0.22	URE 1	1		
Staphylococcus equorum		12.88	0.13		1		
323740	**Good identification**						**323740**
					NOV	PYR	
Rothia mucilaginosa		95.13	0.37	URE 1	1	75	
Staphylococcus hominis subsp. hominis		2.28	0.16		1	1	
Staphylococcus saprophyticus subsp. bovis		1.04	0.05		99	99	
323741	**Acceptable identification**						**323741**
					OXI		
Rothia mucilaginosa		93.96	0.47	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria kristinae		5.63	0.25		99		
Staphylococcus petrasii subsp. petrasii		0.41	0.02		1		
323750	**Good identification**						**323750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.58	0.47	ESL 1	99	99	
Rothia mucilaginosa		0.92	-.04		1	75	
Staphylococcus warneri		0.51	0.09		1	8	
323751	**Low discrimination**						**323751**
					OXI	PYR	
Staphylococcus saprophyticus subsp. bovis		64.43	0.34	ESL 1	1	99	
Staphylococcus lentus		16.57	0.25		99	6	
Staphylococcus xylosus		10.01	0.14		1	65	
323752	**Good identification**						**323752**
					PYR		
Staphylococcus saprophyticus subsp. bovis		95.32	0.34	ESL 1	99		
Staphylococcus saprophyticus subsp. saprophyticus		4.68	-.04		7		
323753	**Low discrimination**						**323753**
					OXI	PYR	
Staphylococcus lentus		56.85	0.36	VPT 7	99	6	
Staphylococcus xylosus		27.56	0.23	URE 1	1	65	
Staphylococcus saprophyticus subsp. bovis		15.59	0.22		1	99	
323771	**Low discrimination**						**323771**
					OXI		
Staphylococcus xylosus		69.51	0.36	VPT 14	1		
Staphylococcus equorum		18.81	0.25	GLR 95	1		
Staphylococcus lentus		10.01	0.25		99		
323773	**Low discrimination**						**323773**
					OXI		
Staphylococcus xylosus		69.23	0.45	VPT 14	1		
Staphylococcus equorum		16.93	0.34	GLR 95	1		
Staphylococcus lentus		12.42	0.36		99		
324050	**Good identification**						**324050**
Staphylococcus kloosii		100.0	0.27	VPT 10			
324052	**Low discrimination**						**324052**
Staphylococcus kloosii		54.90	0.28	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		45.10	-.03	TRE 99			
324150	**Good identification**						**324150**
Staphylococcus kloosii		100.0	0.33	VPT 10			
324152	**Good identification**						**324152**
Staphylococcus kloosii		100.0	0.33	VPT 10			
324241	**Acceptable identification**						**324241**
					OXI	PYR	
Staphylococcus epidermidis		90.69	0.29	NIT 97	1	1	
Kocuria kristinae		9.32	0.03		99	93	
324242	**Good identification**						**324242**
					NOV		
Staphylococcus epidermidis		97.54	0.34	NIT 97	1		
Staphylococcus saprophyticus subsp. saprophyticus		2.46	-.11		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
324243	**Good identification**						**324243**
	Staphylococcus epidermidis	100.0	0.40	NIT 97			
324340	**Good identification**						**324340**
					NOV		
	Staphylococcus epidermidis	97.63	0.39	NIT 97	1		
	Staphylococcus hominis subsp. novobiosepticus	2.37	-.02		99		
324341	**Good identification**						**324341**
	Staphylococcus epidermidis	100.0	0.45	NIT 97			
324342	**Good identification**						**324342**
					NOV		
	Staphylococcus epidermidis	99.59	0.50	NIT 97	1		
	Staphylococcus hominis subsp. novobiosepticus	0.41	-.07		99		
324343	**Very good identification**						**324343**
					COA	PYR	
	Staphylococcus epidermidis	99.59	0.55	NIT 97	1	1	
	Staphylococcus intermedius	0.41	0.04		99	90	
324410	**Good identification**						**324410**
	Staphylococcus kloosii	100.0	0.42	VPT 10			
324412	**Low discrimination**						**324412**
	Staphylococcus kloosii	86.38	0.42	VPT 10			
	Staphylococcus saprophyticus subsp. saprophyticus	13.62	-.03	MLT 95			
324413	**Good identification**						**324413**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	GLR 99			
324430	**Good identification**						**324430**
	Staphylococcus kloosii	100.0	0.29	VPT 10			
324432	**Good identification**						**324432**
	Staphylococcus kloosii	100.0	0.30	VPT 10			
324440	**Low discrimination**						**324440**
	Staphylococcus kloosii	60.69	0.27	VPT 10			
	Staphylococcus saprophyticus subsp. saprophyticus	39.31	-.06	MAN 99			
324443	**Good identification**						**324443**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.34	GLR 99			
324450	**Low discrimination**						**324450**
					PYR		
	Staphylococcus kloosii	85.68	0.68	VPT 10	52		
	Staphylococcus saprophyticus subsp. saprophyticus	13.45	0.23		7		
	Staphylococcus saprophyticus subsp. bovis	0.86	0.09		99		
324451	**Low discrimination**						**324451**
	Staphylococcus cohnii subsp. urealyticum	88.84	0.34	GLR 99			
	Staphylococcus kloosii	11.16	0.27	LAC 91			
324452	**Low discrimination**						**324452**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		54.68	0.69	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		44.91	0.38				
Staphylococcus cohnii subsp. urealyticum		0.42	0.13				
324453	**Very good identification**						**324453**
Staphylococcus cohnii subsp. urealyticum		97.61	0.54	GLR 99			
Staphylococcus kloosii		1.31	0.28				
Staphylococcus saprophyticus subsp. saprophyticus		1.08	-.03				
324470	**Very good identification**						**324470**
Staphylococcus kloosii		100.0	0.55	VPT 10			
324472	**Very good identification**						**324472**
Staphylococcus kloosii		96.79	0.56	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		3.21	-.03				
324510	**Good identification**						**324510**
Staphylococcus kloosii		100.0	0.47	VPT 10			
324512	**Good identification**						**324512**
Staphylococcus kloosii		100.0	0.48	VPT 10			
324513	**Good identification**						**324513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	GLR 99			
324530	**Good identification**						**324530**
Staphylococcus kloosii		100.0	0.35	VPT 10			
324532	**Good identification**						**324532**
Staphylococcus kloosii		100.0	0.36	VPT 10			
324540	**Good identification**						**324540**
Staphylococcus kloosii		100.0	0.33	VPT 10			
324542	**Low discrimination**						**324542**
Staphylococcus kloosii		86.82	0.33	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		13.18	-.12	MAN 99			
324550	**Very good identification**						**324550**
					PYR		
Staphylococcus kloosii		97.30	0.74	VPT 10	52		
Staphylococcus saprophyticus subsp. bovis		2.02	0.22		99		
Staphylococcus saprophyticus subsp. saprophyticus		0.68	0.01		7		
324552	**Very good identification**						**324552**
					PYR		
Staphylococcus kloosii		95.85	0.75	VPT 10	52		
Staphylococcus saprophyticus subsp. saprophyticus		3.53	0.16		7		
Staphylococcus saprophyticus subsp. bovis		0.46	0.09		99		
324553	**Acceptable identification**						**324553**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		91.87	0.47	GLR 99	99	64	
Staphylococcus kloosii		5.60	0.33		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus caprae		2.53	-.02		1	1		
324570	**Very good identification**							**324570**
Staphylococcus kloosii		100.0	0.61	VPT 10				
324572	**Very good identification**							**324572**
Staphylococcus kloosii		100.0	0.62	VPT 10				
324641	**Good identification**							**324641**
Kocuria kristinae		100.0	0.30	URE 10				
324642	**Good identification**							**324642**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		97.10	0.31	PHS 1	99			
Staphylococcus warneri		2.91	0.20	MAN 96	1			
324650	**Acceptable identification**							**324650**
					PYR			
Staphylococcus saprophyticus subsp. saprophyticus		90.66	0.43	PHS 1	7			
Staphylococcus kloosii		5.65	0.47		52			
Staphylococcus saprophyticus subsp. bovis		2.30	0.22		99			
324652	**Very good identification**							**324652**
					PYR			
Staphylococcus saprophyticus subsp. saprophyticus		98.63	0.59	PHS 1	7			
Staphylococcus kloosii		1.18	0.48		52			
Staphylococcus saprophyticus subsp. bovis		0.11	0.09		99			
324670	**Low discrimination**							**324670**
Staphylococcus kloosii		60.66	0.35	VPT 10				
Staphylococcus saprophyticus subsp. saprophyticus		39.34	0.02	SUC 9				
324710	**Good identification**							**324710**
Staphylococcus kloosii		100.0	0.27	VPT 10				
324712	**Low discrimination**							**324712**
Staphylococcus kloosii		58.08	0.28	VPT 10				
Staphylococcus saprophyticus subsp. saprophyticus		41.93	-.04	SUC 9				
324713	**Good identification**							**324713**
					COA			
Staphylococcus intermedius		96.26	0.32	VPT 1	99			
Staphylococcus felis		3.74	-.04	NIT 93	1			
324743	**Low discrimination**							**324743**
					OXI	COA	PYR	
Staphylococcus intermedius		74.22	0.27	VPT 1	1	99	90	
Staphylococcus epidermidis		13.59	0.14	NIT 93	1	1	1	
Kocuria kristinae		12.19	0.07		99	1	93	
324750	**Low discrimination**							**324750**
					PYR			
Staphylococcus kloosii		38.50	0.53	VPT 10	52			
Staphylococcus saprophyticus subsp. bovis		32.27	0.34	SUC 9	99			
Staphylococcus saprophyticus subsp. saprophyticus		27.69	0.22		7			
324752	**Low discrimination**							**324752**
					PYR			
Staphylococcus saprophyticus subsp. saprophyticus		75.92	0.37	PHS 1	7			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		20.18	0.54	GAL 8	52			
Staphylococcus saprophyticus subsp. bovis		3.90	0.22		99			
324753	**Low discrimination**							**324753**
					COA	PYR	NOV	
Staphylococcus intermedius		80.01	0.36	VPT 1	99	90	1	
Staphylococcus aureus subsp. aureus		5.68	-.02	NIT 93	99	10	4	
Staphylococcus cohnii subsp. urealyticum		3.65	0.16		1	53	99	
324770	**Good identification**							**324770**
Staphylococcus kloosii		100.0	0.41	VPT 10				
324772	**Low discrimination**							**324772**
Staphylococcus kloosii		86.81	0.41	VPT 10				
Staphylococcus saprophyticus subsp. saprophyticus		13.19	-.04	SUC 9				
324773	**Low discrimination**							**324773**
					COA	NOV		
Staphylococcus xylosus		63.98	0.27	VPT 14	1	99		
Staphylococcus lutrae		36.02	0.18	GLR 95	99	1		
325050	**Good identification**							**325050**
Staphylococcus kloosii		100.0	0.27	VPT 10				
325052	**Good identification**							**325052**
Staphylococcus kloosii		100.0	0.28	VPT 10				
325150	**Good identification**							**325150**
Staphylococcus kloosii		100.0	0.33	VPT 10				
325152	**Good identification**							**325152**
Staphylococcus kloosii		100.0	0.33	VPT 10				
325410	**Good identification**							**325410**
Staphylococcus kloosii		100.0	0.42	VPT 10				
325412	**Good identification**							**325412**
Staphylococcus kloosii		100.0	0.42	VPT 10				
325413	**Good identification**							**325413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	GLR 99				
325430	**Good identification**							**325430**
Staphylococcus kloosii		100.0	0.29	VPT 10				
325432	**Good identification**							**325432**
Staphylococcus kloosii		100.0	0.30	VPT 10				
325440	**Good identification**							**325440**
Staphylococcus kloosii		100.0	0.27	VPT 10				
325442	**Good identification**							**325442**
Staphylococcus kloosii		100.0	0.28	VPT 10				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
325450	**Very good identification**					**325450**
Staphylococcus kloosii		100.0	0.68	VPT 10		
325452	**Very good identification**					**325452**
Staphylococcus kloosii		98.89	0.69	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		0.82	-.03			
Staphylococcus cohnii subsp. urealyticum		0.29	0.05			
325453	**Good identification**					**325453**
Staphylococcus cohnii subsp. urealyticum		96.66	0.46	GLR 99		
Staphylococcus kloosii		3.34	0.28			
325470	**Very good identification**					**325470**
Staphylococcus kloosii		100.0	0.55	VPT 10		
325472	**Very good identification**					**325472**
Staphylococcus kloosii		100.0	0.56	VPT 10		
325510	**Good identification**					**325510**
Staphylococcus kloosii		100.0	0.47	VPT 10		
325512	**Good identification**					**325512**
Staphylococcus kloosii		100.0	0.48	VPT 10		
325530	**Good identification**					**325530**
Staphylococcus kloosii		100.0	0.35	VPT 10		
325532	**Good identification**					**325532**
Staphylococcus kloosii		100.0	0.36	VPT 10		
325540	**Good identification**					**325540**
Staphylococcus kloosii		100.0	0.33	VPT 10		
325542	**Good identification**					**325542**
Staphylococcus kloosii		100.0	0.33	VPT 10		
325550	**Very good identification**					**325550**
Staphylococcus kloosii		100.0	0.74	VPT 10		
325551	**Low discrimination**					**325551**
Staphylococcus kloosii		59.40	0.33	VPT 10		
Staphylococcus cohnii subsp. urealyticum		40.61	0.18	MNS 1		
325552	**Very good identification**					**325552**
Staphylococcus kloosii		100.0	0.75	VPT 10		
325553	**Low discrimination**					**325553**
Staphylococcus cohnii subsp. urealyticum		86.45	0.38	GLR 99		
Staphylococcus kloosii		13.55	0.33			
325570	**Very good identification**					**325570**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.61	VPT 10		
325572	**Very good identification**					**325572**
Staphylococcus kloosii		100.0	0.62	VPT 10		
325650	**Low discrimination**					**325650**
Staphylococcus kloosii		86.05	0.47	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		13.95	0.02	SUC 9		
325652	**Low discrimination**					**325652**
Staphylococcus kloosii		54.11	0.48	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		45.89	0.18	SUC 9		
325670	**Good identification**					**325670**
Staphylococcus kloosii		100.0	0.35	VPT 10		
325672	**Good identification**					**325672**
Staphylococcus kloosii		100.0	0.36	VPT 10		
325710	**Good identification**					**325710**
Staphylococcus kloosii		100.0	0.27	VPT 10		
325712	**Good identification**					**325712**
Staphylococcus kloosii		100.0	0.28	VPT 10		
325750	**Very good identification**					**325750**
Staphylococcus kloosii		100.0	0.53	VPT 10		
325752	**Very good identification**					**325752**
Staphylococcus kloosii		96.34	0.54	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		3.66	-.04	SUC 9		
325770	**Good identification**					**325770**
Staphylococcus kloosii		100.0	0.41	VPT 10		
325772	**Good identification**					**325772**
Staphylococcus kloosii		100.0	0.41	VPT 10		
326000	**Acceptable identification**					**326000**
					OXI	
Kocuria varians		90.74	0.29	VPT 14	13	
Dermacoccus nishinomiyaensis		9.26	0.01	PHS 1	93	
326001	**Good identification**					**326001**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.06	0.33	URE 1	1	
Kocuria varians		3.08	0.07	ARG 86	1	
Staphylococcus schleiferi subsp. coagulans		1.86	0.02		99	
326043	**Good identification**					**326043**
Staphylococcus epidermidis		100.0	0.30	SUC 99		
326100	**Good identification**					**326100**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COA	
Kocuria varians		96.61	0.33	VPT 14	1	
Staphylococcus schleiferi subsp. coagulans		3.39	0.02	PHS 1	99	
326101	**Low discrimination**					**326101**
					COA	
Staphylococcus schleiferi subsp. coagulans		64.92	0.43	ARG 99	99	
Staphylococcus schleiferi subsp. schleiferi		33.45	0.33	LAC 86	1	
Kocuria varians		1.63	0.11		1	
326103	**Very good identification**					**326103**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.77	0.59	ARG 99	99	
Staphylococcus epidermidis		0.23	0.04		1	
326111	**Good identification**					**326111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ARG 99		
326113	**Very good identification**					**326113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	ARG 99	99	
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1	
326140	**Low discrimination**					**326140**
Staphylococcus epidermidis		79.26	0.29	SUC 99		
Kocuria varians		20.75	0.19			
326141	**Good identification**					**326141**
					COA	
Staphylococcus epidermidis		97.80	0.35	SUC 99	1	
Staphylococcus schleiferi subsp. coagulans		2.20	0.02		99	
326142	**Good identification**					**326142**
Staphylococcus epidermidis		97.72	0.40	SUC 99		
Kocuria varians		2.28	0.08			
326143	**Good identification**					**326143**
					COA	
Staphylococcus epidermidis		95.81	0.45	SUC 99	1	
Staphylococcus schleiferi subsp. coagulans		4.19	0.18		99	
326203	**Acceptable identification**					**326203**
					COA	
Staphylococcus epidermidis		92.74	0.30	MLT 99	1	
Staphylococcus schleiferi subsp. coagulans		7.26	0.08		99	
326240	**Very good identification**					**326240**
					NOV	
Staphylococcus epidermidis		97.94	0.55		1	
Staphylococcus hominis subsp. novobiosepticus		1.25	0.08		99	
Staphylococcus hominis subsp. hominis		0.81	0.15		1	
326241	**Very good identification**					**326241**
					NOV	
Staphylococcus epidermidis		99.83	0.60		1	
Staphylococcus hominis subsp. novobiosepticus		0.17	-.04		99	
326242	**Very good identification**					**326242**
					NOV	
Staphylococcus epidermidis		99.48	0.65		1	
Staphylococcus hominis subsp. hominis		0.31	0.17		1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hominis subsp. novobiosepticus		0.22	0.03		99			
326243	**Very good identification**							**326243**
Staphylococcus epidermidis		100.0	0.71					
326253	**Low discrimination**							**326253**
Staphylococcus epidermidis		81.93	0.30	MAN 1				
Staphylococcus capitis subsp. ureolyticus		13.70	0.00					
Staphylococcus simulans		4.37	0.03					
326263	**Good identification**							**326263**
Staphylococcus epidermidis		100.0	0.30	XYL 1				
326300	**Low discrimination**							**326300**
Staphylococcus epidermidis		86.01	0.29	MLT 99				
Kocuria varians		13.99	0.15					
326301	**Low discrimination**							**326301**
					COA	PYR		
Staphylococcus epidermidis		56.73	0.35	MLT 99	1	1		
Staphylococcus schleiferi subsp. coagulans		39.97	0.32		99	20		
Staphylococcus intermedius		3.29	0.07		99	90		
326302	**Good identification**							**326302**
					COA			
Staphylococcus epidermidis		96.14	0.40	MLT 99	1			
Staphylococcus schleiferi subsp. coagulans		2.47	0.08		99			
Kocuria varians		1.39	0.04		1			
326303	**Low discrimination**							**326303**
					COA	PYR		
Staphylococcus schleiferi subsp. coagulans		54.47	0.49	ARG 99	99	20		
Staphylococcus epidermidis		39.85	0.45		1	1		
Staphylococcus intermedius		4.49	0.23		99	90		
326311	**Low discrimination**							**326311**
Staphylococcus schleiferi subsp. coagulans		80.56	0.32	ARG 99				
Staphylococcus intermedius		19.44	0.16	LAC 86				
326313	**Low discrimination**							**326313**
					COA			
Staphylococcus schleiferi subsp. coagulans		79.03	0.48	ARG 99	99			
Staphylococcus intermedius		19.07	0.32		99			
Staphylococcus felis		0.74	-.04		1			
326340	**Very good identification**							**326340**
					NOV			
Staphylococcus epidermidis		99.14	0.70		1			
Staphylococcus hominis subsp. novobiosepticus		0.67	0.18		99			
Staphylococcus hominis subsp. hominis		0.16	0.16		1			
326341	**Very good identification**							**326341**
					COA	NOV	PYR	
Staphylococcus epidermidis		99.82	0.76		1	1	1	
Staphylococcus hominis subsp. novobiosepticus		0.09	0.05		1	99	1	
Staphylococcus intermedius		0.09	0.11		99	1	90	
326342	**Very good identification**							**326342**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus epidermidis		99.83	0.81		1			
Staphylococcus hominis subsp. novobiosepticus		0.12	0.12		99			
Staphylococcus hominis subsp. hominis		0.06	0.18		1			
326343	**Very good identification**							**326343**
					COA	YEP	PYR	
Staphylococcus epidermidis		99.77	0.86		1	1	1	
Staphylococcus intermedius		0.17	0.27		99	1	90	
Staphylococcus chromogenes		0.03	0.10		1	95	45	
326350	**Low discrimination**							**326350**
					PYR	NOV		
Staphylococcus saprophyticus subsp. bovis		64.68	0.34	TRE 99	99	99		
Staphylococcus epidermidis		35.32	0.29		1	1		
326351	**Low discrimination**							**326351**
					COA	NOV	PYR	
Staphylococcus epidermidis		67.48	0.35	MAN 1	1	1	1	
Staphylococcus saprophyticus subsp. bovis		16.63	0.22		1	99	99	
Staphylococcus intermedius		15.89	0.20		99	1	90	
326352	**Low discrimination**							**326352**
					PYR	NOV		
Staphylococcus epidermidis		87.37	0.40	MAN 1	1	1		
Staphylococcus saprophyticus subsp. bovis		12.63	0.22		99	99		
326353	**Low discrimination**							**326353**
					COA	PYR		
Staphylococcus epidermidis		64.81	0.45	MAN 1	1	1		
Staphylococcus intermedius		29.60	0.36		99	90		
Staphylococcus aureus subsp. aureus		2.10	-.02		99	10		
326360	**Good identification**							**326360**
Staphylococcus epidermidis		100.0	0.29	XYL 1				
326361	**Good identification**							**326361**
Staphylococcus epidermidis		100.0	0.35	XYL 1				
326362	**Good identification**							**326362**
Staphylococcus epidermidis		100.0	0.40	XYL 1				
326363	**Acceptable identification**							**326363**
					COA			
Staphylococcus epidermidis		93.98	0.45	XYL 1	1			
Staphylococcus lutrae		6.02	0.12		99			
326401	**Good identification**							**326401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.43	URE 1				
326450	**Good identification**							**326450**
Staphylococcus saprophyticus subsp. bovis		99.00	0.50					
Staphylococcus kloosii		1.01	0.27					
326451	**Acceptable identification**							**326451**
Staphylococcus saprophyticus subsp. bovis		94.61	0.38					
Staphylococcus cohnii subsp. urealyticum		5.40	0.18					
326452	**Acceptable identification**							**326452**
					PYR			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		92.59	0.38		99		
Staphylococcus kloosii		4.07	0.28		52		
Staphylococcus saprophyticus subsp. saprophyticus		3.34	-.03		7		
326453	**Low discrimination**						**326453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		68.48	0.39	GLR 99	99		
Staphylococcus saprophyticus subsp. bovis		29.69	0.26		99		
Staphylococcus simulans		1.84	0.01		1		
326501	**Good identification**						**326501**
					COA		
Staphylococcus schleiferi subsp. schleiferi		99.35	0.43	URE 1	1		
Staphylococcus schleiferi subsp. coagulans		0.65	0.02	ARG 86	99		
326513	**Low discrimination**						**326513**
					COA		
Staphylococcus felis		69.06	0.31	VPT 1	1		
Staphylococcus condimenti		17.83	0.14	ARG 92	1		
Staphylococcus intermedius		4.89	0.14		99		
326550	**Very good identification**						**326550**
Staphylococcus saprophyticus subsp. bovis		99.51	0.63				
Staphylococcus kloosii		0.49	0.33				
326551	**Good identification**						**326551**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		98.44	0.50		1	99	
Staphylococcus cohnii subsp. urealyticum		0.60	0.10		1	99	
Staphylococcus intermedius		0.51	0.02		99	1	
326552	**Good identification**						**326552**
					NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		96.79	0.50		99	80	
Staphylococcus kloosii		2.07	0.33		99	99	
Staphylococcus caprae		1.15	0.00		1	1	
326553	**Low discrimination**						**326553**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		48.45	0.38		1	99	
Staphylococcus caprae		16.83	0.18		1	1	
Staphylococcus lutrae		12.50	0.18		99	1	
326563	**Very good identification**						**326563**
					COA	NOV	
Staphylococcus lutrae		99.91	0.53	VPT 1	99	1	
Staphylococcus xylosus		0.10	-.05		1	99	
326573	**Very good identification**						**326573**
					COA	NOV	
Staphylococcus lutrae		99.24	0.59	VPT 1	99	1	
Staphylococcus xylosus		0.76	0.20		1	99	
326613	**Low discrimination**						**326613**
					COA		
Staphylococcus simulans		49.78	0.36	VPT 4	1		
Staphylococcus felis		43.78	0.26	ARG 92	1		
Staphylococcus intermedius		5.11	0.14		99		
326643	**Low discrimination**						**326643**
					YEP		
Staphylococcus epidermidis		55.18	0.30	TRE 1	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		13.99	0.17		1		
Staphylococcus chromogenes		11.87	0.12		95		
326650	**Very good identification**						**326650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		99.19	0.63		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		0.40	0.02		99	7	
Staphylococcus warneri		0.26	0.19		1	8	
326651	**Good identification**						**326651**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		99.49	0.50		1	99	
Staphylococcus intermedius		0.52	0.02		99	1	
326652	**Low discrimination**						**326652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		88.53	0.50		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.08	0.18		99	7	
Staphylococcus simulans		2.46	0.18		1	70	
326653	**Low discrimination**						**326653**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		58.59	0.38		1	99	
Staphylococcus simulans		21.77	0.29		1	1	
Staphylococcus aureus subsp. aureus		7.97	0.05		99	4	
326701	**Low discrimination**						**326701**
					COA	YEP	
Staphylococcus intermedius		83.26	0.30	VPT 1	99	1	
Staphylococcus chromogenes		12.98	0.12	LAC 86	1	95	
Staphylococcus schleiferi subsp. schleiferi		3.76	0.02		1	1	
326703	**Low discrimination**						**326703**
					COA	YEP	
Staphylococcus intermedius		72.74	0.46	VPT 1	99	1	
Staphylococcus chromogenes		16.61	0.32		1	95	
Staphylococcus felis		8.52	0.20		1	1	
326710	**Good identification**						**326710**
					COA	NOV	
Staphylococcus saprophyticus subsp. bovis		97.91	0.34	MLT 99	1	99	
Staphylococcus intermedius		2.09	-.02		99	1	
326711	**Low discrimination**						**326711**
					COA	NOV	
Staphylococcus intermedius		86.77	0.39	VPT 1	99	1	
Staphylococcus saprophyticus subsp. bovis		10.26	0.22	LAC 86	1	99	
Staphylococcus felis		1.56	-.04		1	1	
326713	**Low discrimination**						**326713**
					COA		
Staphylococcus intermedius		71.62	0.55	VPT 1	99		
Staphylococcus felis		20.76	0.37		1		
Staphylococcus condimenti		3.96	0.18		1		
326740	**Low discrimination**						**326740**
					PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		54.20	0.34	MAN 99	99	99	
Staphylococcus epidermidis		29.60	0.29		1	1	
Staphylococcus hominis subsp. hominis		14.67	0.26		1	1	
326741	**Low discrimination**						**326741**
					COA	NOV	PYR

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		44.55	0.34	VPT 1	99	1	90	
Staphylococcus epidermidis		38.50	0.35	LAC 86	1	1	1	
Staphylococcus saprophyticus subsp. bovis		9.49	0.22		1	99	99	
326742	**Low discrimination**							**326742**
					PYR	NOV		
Staphylococcus epidermidis		72.98	0.40	TRE 1	1	1		
Staphylococcus hominis subsp. hominis		13.40	0.28		1	1		
Staphylococcus saprophyticus subsp. bovis		10.55	0.22		99	99		
326743	**Low discrimination**							**326743**
					COA	YEP	PYR	
Staphylococcus intermedius		60.02	0.50	VPT 1	99	1	90	
Staphylococcus epidermidis		26.74	0.45		1	1	1	
Staphylococcus chromogenes		9.14	0.32		1	95	45	
326750	**Very good identification**							**326750**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		99.91	0.75		1	99	99	
Staphylococcus hominis subsp. hominis		0.04	0.09		1	1	1	
Staphylococcus intermedius		0.03	0.02		99	1	90	
326751	**Low discrimination**							**326751**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		87.65	0.63		1	99		
Staphylococcus intermedius		11.24	0.43		99	1		
Staphylococcus xylosus		0.62	0.15		1	99		
326752	**Very good identification**							**326752**
					COA	PYR	NOV	
Staphylococcus saprophyticus subsp. bovis		98.31	0.63		1	99	99	
Staphylococcus intermedius		0.78	0.18		99	90	1	
Staphylococcus aureus subsp. aureus		0.41	-.02		99	10	4	
326753	**Low discrimination**							**326753**
					COA	PYR	NOV	
Staphylococcus intermedius		51.95	0.59	VPT 1	99	90	1	
Staphylococcus aureus subsp. aureus		27.48	0.39		99	10	4	
Staphylococcus saprophyticus subsp. bovis		16.49	0.50		1	99	99	
326763	**Very good identification**							**326763**
					COA	PYR	NOV	
Staphylococcus lutrae		97.74	0.53	VPT 1	99	10	1	
Staphylococcus xylosus		1.76	0.22		1	65	99	
Staphylococcus intermedius		0.35	0.09		99	90	1	
326770	**Good identification**							**326770**
Staphylococcus saprophyticus subsp. bovis		98.01	0.34	XYL 1				
Staphylococcus xylosus		2.00	-.04					
326771	**Low discrimination**							**326771**
					COA	NOV		
Staphylococcus xylosus		73.57	0.37	VPT 14	1	99		
Staphylococcus lutrae		14.57	0.18	GLR 95	99	1		
Staphylococcus saprophyticus subsp. bovis		9.13	0.22		1	99		
326773	**Low discrimination**							**326773**
					COA	PYR	NOV	
Staphylococcus lutrae		86.43	0.59	VPT 1	99	10	1	
Staphylococcus xylosus		12.55	0.46		1	65	99	
Staphylococcus intermedius		0.43	0.18		99	90	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
327101	**Good identification**					**327101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ARG 99		
327103	**Very good identification**					**327103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	ARG 99		
327111	**Good identification**					**327111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ARG 99		
327113	**Very good identification**					**327113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	ARG 99		
327243	**Good identification**					**327243**
Staphylococcus epidermidis		100.0	0.30	ESL 1		
327301	**Good identification**					**327301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	ARG 99		
327303	**Good identification**					**327303**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.24	0.48	ARG 99	99	
Staphylococcus epidermidis		0.76	0.04		1	
327311	**Good identification**					**327311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	ARG 99		
327313	**Good identification**					**327313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	ARG 99		
327340	**Acceptable identification**					**327340**
					NOV	
Staphylococcus epidermidis		94.50	0.29	ESL 1	1	
Staphylococcus hominis subsp. novobiosepticus		5.51	-.04		99	
327341	**Good identification**					**327341**
Staphylococcus epidermidis		100.0	0.35	ESL 1		
327342	**Good identification**					**327342**
Staphylococcus epidermidis		100.0	0.40	ESL 1		
327343	**Good identification**					**327343**
					COA	
Staphylococcus epidermidis		98.69	0.45	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		1.31	0.07		99	
327450	**Low discrimination**					**327450**
Staphylococcus kloosii		50.11	0.27	VPT 10		
Staphylococcus saprophyticus subsp. bovis		49.89	0.09	NIT 1		
327452	**Good identification**					**327452**
Staphylococcus kloosii		100.0	0.28	VPT 10		
327453	**Good identification**					**327453**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	GLR 99				
327552	**Low discrimination**							**327552**
Staphylococcus kloosii		67.87	0.33	VPT 10				
Staphylococcus saprophyticus subsp. bovis		32.13	0.09	NIT 1				
327750	**Good identification**							**327750**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ESL 1				
327763	**Low discrimination**							**327763**
					COA	NOV		
Staphylococcus gallinarum		77.60	0.29	VPT 1	1	99		
Staphylococcus lutrae		15.50	0.12	MAN 91	99	1		
Staphylococcus xylosus		6.90	0.09		1	99		
327771	**Low discrimination**							**327771**
Staphylococcus gallinarum		76.67	0.42	VPT 1				
Staphylococcus equorum		13.08	0.29					
Staphylococcus xylosus		9.09	0.24					
327773	**Low discrimination**							**327773**
Staphylococcus gallinarum		72.61	0.50	VPT 1				
Staphylococcus equorum		13.01	0.38					
Staphylococcus xylosus		10.00	0.34					
330452	**Good identification**							**330452**
Staphylococcus saprophyticus subsp. saprophyticus		100.0	0.38	ORN 1				
330641	**Low discrimination**							**330641**
					OXI			
Kocuria kristinae		60.57	0.30	URE 10	99			
Staphylococcus lugdunensis		39.43	0.27	ORN 1	1			
330642	**Acceptable identification**							**330642**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		94.60	0.31	ORN 1	99	7		
Staphylococcus warneri		2.83	0.20	MAN 96	1	8		
Staphylococcus lugdunensis		2.57	0.05		1	96		
330643	**Good identification**							**330643**
					OXI	NOV	PYR	
Staphylococcus lugdunensis		95.56	0.47	NIT 99	1	1	96	
Kocuria kristinae		4.08	0.17		99	1	93	
Staphylococcus saprophyticus subsp. saprophyticus		0.36	-.11		1	99	7	
330650	**Good identification**							**330650**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		98.49	0.43	ORN 1	99			
Staphylococcus warneri		1.51	0.27		1			
330652	**Very good identification**							**330652**
					NOV			
Staphylococcus saprophyticus subsp. saprophyticus		99.91	0.59	ORN 1	99			
Staphylococcus warneri		0.09	0.17		1			
330741	**Low discrimination**							**330741**
					OXI			
Staphylococcus lugdunensis		66.13	0.27	NIT 99	1			
Kocuria kristinae		33.87	0.20	LAC 90	99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
330743	**Good identification**						**330743**
					OXI		
Staphylococcus lugdunensis		98.60	0.47	NIT 99	1		
Kocuria kristinae		1.40	0.07		99		
330752	**Good identification**						**330752**
Staphylococcus saprophyticus subsp. saprophyticus		100.0	0.37	ORN 1			
332000	**Good identification**						**332000**
Kocuria varians		100.0	0.29	VPT 14			
332100	**Good identification**						**332100**
Kocuria varians		100.0	0.33	VPT 14			
332241	**Low discrimination**						**332241**
					NOV	PYR	
Staphylococcus lugdunensis		60.38	0.27	TRE 99	1	96	
Staphylococcus epidermidis		36.98	0.27	LAC 90	1	1	
Staphylococcus hominis subsp. novobiosepticus		2.64	-.04		99	1	
332242	**Low discrimination**						**332242**
					NOV	PYR	
Staphylococcus epidermidis		82.04	0.32	ORN 12	1	1	
Staphylococcus hominis subsp. novobiosepticus		7.41	0.03		99	1	
Staphylococcus lugdunensis		7.14	0.05		1	96	
332243	**Low discrimination**						**332243**
					PYR		
Staphylococcus lugdunensis		82.27	0.47	TRE 99	96		
Staphylococcus epidermidis		17.73	0.37		1		
332340	**Low discrimination**						**332340**
					NOV		
Staphylococcus epidermidis		75.89	0.37	ORN 12	1		
Staphylococcus hominis subsp. novobiosepticus		21.34	0.18		99		
Staphylococcus hominis subsp. hominis		1.63	0.06		1		
332341	**Low discrimination**						**332341**
					NOV	PYR	
Staphylococcus epidermidis		75.42	0.43	ORN 12	1	1	
Staphylococcus lugdunensis		21.73	0.27		1	96	
Staphylococcus hominis subsp. novobiosepticus		2.86	0.05		99	1	
332342	**Acceptable identification**						**332342**
					NOV	PYR	
Staphylococcus epidermidis		93.36	0.47	ORN 12	1	1	
Staphylococcus hominis subsp. novobiosepticus		4.46	0.12		99	1	
Staphylococcus lugdunensis		1.43	0.05		1	96	
332343	**Low discrimination**						**332343**
					NOV	PYR	
Staphylococcus epidermidis		54.79	0.53	ORN 12	1	1	
Staphylococcus lugdunensis		44.86	0.47		1	96	
Staphylococcus hominis subsp. novobiosepticus		0.35	0.00		99	1	
332441	**Good identification**						**332441**
Staphylococcus lugdunensis		100.0	0.27	SUC 99			
332443	**Good identification**						**332443**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lugdunensis		100.0	0.47	SUC 99			
332541	**Good identification**						**332541**
Staphylococcus lugdunensis		100.0	0.27	SUC 99			
332543	**Good identification**						**332543**
Staphylococcus lugdunensis		100.0	0.47	SUC 99			
332550	**Good identification**						**332550**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
332601	**Good identification**						**332601**
Staphylococcus lugdunensis		100.0	0.27	MLT 99			
332603	**Good identification**						**332603**
Staphylococcus lugdunensis		100.0	0.47	MLT 99			
332640	**Low discrimination**						**332640**
					PYR		
Staphylococcus lugdunensis		80.26	0.27	MNS 99	96		
Staphylococcus warneri		10.33	0.22	LAC 90	8		
Staphylococcus hominis subsp. hominis		9.40	0.16		1		
332641	**Very good identification**						**332641**
					OXI		
Staphylococcus lugdunensis		99.78	0.68	LAC 90	1		
Kocuria kristinae		0.12	0.06		99		
Staphylococcus petrasii subsp. petrasii		0.10	0.05		1		
332642	**Good identification**						**332642**
					PYR		
Staphylococcus lugdunensis		97.68	0.47	MNS 99	96		
Staphylococcus hominis subsp. hominis		1.49	0.17		1		
Staphylococcus warneri		0.47	0.12		8		
332643	**Very good identification**						**332643**
					OXI		
Staphylococcus lugdunensis		99.99	0.88		1		
Staphylococcus petrasii subsp. petrasii		0.01	0.00		1		
Kocuria kristinae		0.00	-.06		99		
332650	**Low discrimination**						**332650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		85.87	0.34	ORN 1	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.56	0.02		99	7	
Staphylococcus warneri		5.57	0.19		1	8	
332651	**Low discrimination**						**332651**
					NOV		
Staphylococcus lugdunensis		73.58	0.27	MAN 1	1		
Staphylococcus saprophyticus subsp. bovis		26.42	0.22	LAC 90	99		
332653	**Good identification**						**332653**
					NOV		
Staphylococcus lugdunensis		99.01	0.47	MAN 1	1		
Staphylococcus saprophyticus subsp. bovis		0.99	0.09		99		
332661	**Good identification**						**332661**
Staphylococcus lugdunensis		100.0	0.27	XYL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
332663	**Good identification**						**332663**
	Staphylococcus lugdunensis	100.0	0.47	XYL 1			
332701	**Good identification**						**332701**
	Staphylococcus lugdunensis	100.0	0.27	MLT 99			
332703	**Good identification**						**332703**
	Staphylococcus lugdunensis	100.0	0.47	MLT 99			
332740	**Low discrimination**						**332740**
					NOV	PYR	
	Staphylococcus lugdunensis	81.59	0.27	MNS 99	1	96	
	Staphylococcus hominis subsp. hominis	10.36	0.16	LAC 90	1	1	
	Staphylococcus saprophyticus subsp. bovis	4.74	0.05		99	99	
332741	**Very good identification**						**332741**
					OXI		
	Staphylococcus lugdunensis	99.86	0.68	LAC 90	1		
	Staphylococcus petrasii subsp. petrasii	0.07	0.02		1		
	Kocuria kristinae	0.04	-.04		99		
332742	**Good identification**						**332742**
					PYR		
	Staphylococcus lugdunensis	97.73	0.47	MNS 99	96		
	Staphylococcus hominis subsp. hominis	1.62	0.18		1		
	Staphylococcus epidermidis	0.65	0.06		1		
332743	**Very good identification**						**332743**
					PYR		
	Staphylococcus lugdunensis	99.98	0.88		96		
	Staphylococcus epidermidis	0.01	0.12		1		
	Staphylococcus petrasii subsp. petrasii	0.00	-.04		99		
332750	**Good identification**						**332750**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	99.49	0.47	ORN 1	99	99	
	Staphylococcus warneri	0.51	0.09		1	8	
332751	**Low discrimination**						**332751**
					NOV		
	Staphylococcus saprophyticus subsp. bovis	58.96	0.34	ORN 1	99		
	Staphylococcus lugdunensis	41.04	0.27		1		
332752	**Low discrimination**						**332752**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	89.90	0.34	ORN 1	99	99	
	Staphylococcus lugdunensis	5.69	0.05		1	96	
	Staphylococcus saprophyticus subsp. saprophyticus	4.42	-.04		99	7	
332753	**Good identification**						**332753**
					NOV		
	Staphylococcus lugdunensis	96.16	0.47	MAN 1	1		
	Staphylococcus saprophyticus subsp. bovis	3.84	0.22		99		
332761	**Good identification**						**332761**
	Staphylococcus lugdunensis	100.0	0.27	XYL 1			
332763	**Good identification**						**332763**
	Staphylococcus lugdunensis	100.0	0.47	XYL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
333641	**Good identification**					**333641**
					OXI	
Staphylococcus lugdunensis		97.19	0.27	ESL 1	1	
Kocuria kristinae		2.81	-.06	LAC 90	99	
333643	**Good identification**					**333643**
Staphylococcus lugdunensis		100.0	0.47	ESL 1		
333741	**Low discrimination**					**333741**
Staphylococcus lugdunensis		86.15	0.27	ESL 1		
Rothia mucilaginosa		13.85	0.05	LAC 90		
333743	**Good identification**					**333743**
Staphylococcus lugdunensis		100.0	0.47	ESL 1		
334341	**Good identification**					**334341**
Staphylococcus epidermidis		100.0	0.27	ORN 12		
334342	**Good identification**					**334342**
Staphylococcus epidermidis		100.0	0.32	ORN 12		
334343	**Good identification**					**334343**
Staphylococcus epidermidis		100.0	0.37	ORN 12		
334450	**Good identification**					**334450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
334452	**Low discrimination**					**334452**
Staphylococcus kloosii		54.90	0.28	VPT 10		
Staphylococcus saprophyticus subsp. saprophyticus		45.10	-.03	ORN 1		
334550	**Good identification**					**334550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
334552	**Good identification**					**334552**
Staphylococcus kloosii		100.0	0.33	VPT 10		
335450	**Good identification**					**335450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
335452	**Good identification**					**335452**
Staphylococcus kloosii		100.0	0.28	VPT 10		
335550	**Good identification**					**335550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
335552	**Good identification**					**335552**
Staphylococcus kloosii		100.0	0.33	VPT 10		
336143	**Good identification**					**336143**
Staphylococcus epidermidis		100.0	0.27	ORN 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
336240	**Good identification**						**336240**
Staphylococcus epidermidis		100.0	0.37	ORN 12			
336241	**Good identification**						**336241**
Staphylococcus epidermidis		100.0	0.43	ORN 12			
336242	**Good identification**						**336242**
Staphylococcus epidermidis		100.0	0.47	ORN 12			
336243	**Good identification**						**336243**
					PYR		
Staphylococcus epidermidis		99.18	0.53	ORN 12	1		
Staphylococcus lugdunensis		0.82	0.05		96		
336303	**Acceptable identification**						**336303**
					COA		
Staphylococcus epidermidis		90.81	0.27	ORN 12	1		
Staphylococcus schleiferi subsp. coagulans		9.19	0.08	MLT 99	99		
336340	**Good identification**						**336340**
Staphylococcus epidermidis		100.0	0.53	ORN 12			
336341	**Very good identification**						**336341**
Staphylococcus epidermidis		100.0	0.58	ORN 12			
336342	**Very good identification**						**336342**
Staphylococcus epidermidis		100.0	0.63	ORN 12			
336343	**Very good identification**						**336343**
					PYR		
Staphylococcus epidermidis		99.85	0.68	ORN 12	1		
Staphylococcus lugdunensis		0.15	0.05		96		
336353	**Good identification**						**336353**
Staphylococcus epidermidis		100.0	0.27	ORN 12			
336363	**Good identification**						**336363**
Staphylococcus epidermidis		100.0	0.27	ORN 12			
336641	**Good identification**						**336641**
					PYR		
Staphylococcus lugdunensis		96.61	0.27	PHS 1	96		
Staphylococcus epidermidis		3.39	0.02	LAC 90	1		
336643	**Good identification**						**336643**
					PYR		
Staphylococcus lugdunensis		98.78	0.47	PHS 1	96		
Staphylococcus epidermidis		1.22	0.12		1		
336741	**Low discrimination**						**336741**
					PYR		
Staphylococcus lugdunensis		83.43	0.27	PHS 1	96		
Staphylococcus epidermidis		16.57	0.17	LAC 90	1		
336743	**Acceptable identification**						**336743**
					COA	PYR	
Staphylococcus lugdunensis		92.46	0.47	PHS 1	1	96	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		6.46	0.27		1	1	
Staphylococcus intermedius		1.08	0.09		99	90	
336750	**Good identification**						**336750**
Staphylococcus saprophyticus subsp. bovis		100.0	0.34	ORN 1			
337343	**Good identification**						**337343**
Staphylococcus epidermidis		100.0	0.27	ORN 12			
340000	**Low discrimination**						**340000**
Dermacoccus nishinomiyaensis		54.80	0.33	VPT 5			
Micrococcus sp.		17.27	0.13	GLR 1			
Kocuria rosea		16.40	0.21				
340013	**Good identification**						**340013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99			
340040	**Acceptable identification**						**340040**
					OXI		
Staphylococcus warneri		90.71	0.43	SUC 95	1		
Kocuria varians		5.79	0.11	TRE 95	13		
Micrococcus sp.		3.51	-.10		89		
340042	**Good identification**						**340042**
Staphylococcus warneri		100.0	0.33	SUC 95			
340050	**Low discrimination**						**340050**
					OXI	NOV	
Staphylococcus warneri		80.85	0.40	SUC 95	1	1	
Macrococcus equipersicus		9.40	0.11	TRE 95	99	99	
Staphylococcus pasteuri		5.82	0.02		1	1	
340052	**Low discrimination**						**340052**
					NOV		
Staphylococcus warneri		78.09	0.30	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		8.00	0.06	TRE 95	99		
Staphylococcus saprophyticus subsp. saprophyticus		7.02	-.12		99		
340053	**Good identification**						**340053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	TRE 99			
340100	**Low discrimination**						**340100**
					OXI		
Kocuria varians		58.41	0.29	VPT 14	13		
Dermacoccus nishinomiyaensis		41.59	0.18	GLR 1	93		
340140	**Low discrimination**						**340140**
					NOV		
Staphylococcus warneri		69.28	0.33	SUC 95	1		
Kocuria varians		21.00	0.15	TRE 95	1		
Staphylococcus hominis subsp. novobiosepticus		9.72	-.02		99		
340150	**Low discrimination**						**340150**
					OXI	NOV	
Staphylococcus warneri		86.37	0.30	SUC 95	1	1	
Macrococcus equipersicus		13.63	0.04	TRE 95	99	99	
340153	**Good identification**						**340153**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	TRE 99				
340200	**Low discrimination**							**340200**
					OXI	YEP		
Staphylococcus warneri		72.18	0.43	TRE 95	1	43		
Staphylococcus pasteuri		9.87	0.11	MLT 95	1	86		
Dermacoccus nishinomiyaensis		7.51	0.08		93	1		
340202	**Low discrimination**							**340202**
Staphylococcus warneri		85.65	0.33	TRE 95				
Staphylococcus pasteuri		14.35	0.03	MLT 95				
340210	**Low discrimination**							**340210**
Staphylococcus pasteuri		53.69	0.27	TRE 99				
Staphylococcus warneri		46.31	0.40	MLT 86				
340240	**Low discrimination**							**340240**
					NOV	YEP		
Staphylococcus warneri		88.00	0.69	TRE 95	1	43		
Staphylococcus hominis subsp. novobiosepticus		6.77	0.30		99	1		
Staphylococcus pasteuri		3.89	0.27		1	86		
340241	**Low discrimination**							**340241**
					OXI	NOV	PYR	
Staphylococcus warneri		41.37	0.35	TRE 95	1	1	8	
Staphylococcus hominis subsp. novobiosepticus		38.99	0.17	MNS 2	1	99	1	
Kocuria kristinae		19.64	0.13		99	1	93	
340242	**Low discrimination**							**340242**
					NOV	YEP		
Staphylococcus warneri		84.49	0.60	TRE 95	1	43		
Staphylococcus hominis subsp. novobiosepticus		10.50	0.24		99	1		
Staphylococcus pasteuri		4.58	0.19		1	86		
340250	**Low discrimination**							**340250**
					NOV			
Staphylococcus warneri		72.58	0.66	TRE 95	1			
Staphylococcus pasteuri		27.21	0.43		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.20	-.07		99			
340251	**Low discrimination**							**340251**
Staphylococcus warneri		84.35	0.32	TRE 95				
Staphylococcus pasteuri		15.65	0.02	MNS 2				
340252	**Low discrimination**							**340252**
					NOV			
Staphylococcus warneri		66.35	0.57	TRE 95	1			
Staphylococcus pasteuri		30.48	0.35		1			
Staphylococcus saprophyticus subsp. saprophyticus		3.18	0.09		99			
340260	**Good identification**							**340260**
Staphylococcus warneri		100.0	0.28	TRE 95				
340300	**Low discrimination**							**340300**
					NOV			
Staphylococcus warneri		75.27	0.33	TRE 95	1			
Kocuria varians		14.17	0.11	MLT 95	1			
Staphylococcus hominis subsp. novobiosepticus		10.56	-.02		99			
340310	**Good identification**							**340310**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		100.0	0.30	TRE 95				
340340	**Low discrimination**							**340340**
					NOV			
Staphylococcus warneri		57.60	0.59	TRE 95	1			
Staphylococcus hominis subsp. novobiosepticus		42.11	0.39		99			
Staphylococcus hominis subsp. hominis		0.29	0.06		1			
340341	**Low discrimination**							**340341**
					OXI	NOV	PYR	
Staphylococcus hominis subsp. novobiosepticus		84.06	0.27	GLR 1	1	99	1	
Staphylococcus warneri		9.39	0.24	NIT 90	1	1	8	
Kocuria kristinae		4.70	0.03		99	1	93	
340342	**Low discrimination**							**340342**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		53.21	0.34	GLR 1	99			
Staphylococcus warneri		45.06	0.49	NIT 90	1			
Staphylococcus epidermidis		0.93	0.05		1			
340350	**Very good identification**							**340350**
					NOV	YEP		
Staphylococcus warneri		97.83	0.56	TRE 95	1	43		
Staphylococcus pasteuri		1.17	0.02		1	86		
Staphylococcus hominis subsp. novobiosepticus		1.00	-.02		99	1		
340352	**Good identification**							**340352**
					NOV	YEP		
Staphylococcus warneri		96.98	0.46	TRE 95	1	43		
Staphylococcus hominis subsp. novobiosepticus		1.60	-.07		99	1		
Staphylococcus pasteuri		1.43	-.06		1	86		
340400	**Low discrimination**							**340400**
					NOV	YEP		
Staphylococcus warneri		63.72	0.43	SUC 95	1	43		
Staphylococcus cohnii subsp. cohnii		26.18	0.30	MLT 95	99	12		
Staphylococcus pasteuri		8.71	0.11		1	86		
340401	**Low discrimination**							**340401**
Staphylococcus cohnii subsp. cohnii		51.55	0.31	URE 1				
Staphylococcus cohnii subsp. urealyticum		48.45	0.31					
340402	**Low discrimination**							**340402**
					NOV			
Staphylococcus warneri		77.18	0.33	SUC 95	1			
Staphylococcus pasteuri		12.93	0.03	MLT 95	1			
Staphylococcus cohnii subsp. urealyticum		9.89	0.11		99			
340403	**Good identification**							**340403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	MAN 91				
340410	**Low discrimination**							**340410**
					NOV	YEP		
Staphylococcus pasteuri		38.46	0.27	SUC 99	1	86		
Staphylococcus warneri		33.17	0.40	MLT 86	1	43		
Staphylococcus cohnii subsp. cohnii		23.95	0.32		99	12		
340411	**Low discrimination**							**340411**
Staphylococcus cohnii subsp. urealyticum		88.19	0.52	LAC 91				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. cohnii		11.81	0.33				
340412	**Low discrimination**						**340412**
					NOV		
Staphylococcus cohnii subsp. urealyticum		41.62	0.31	MNS 99	99		
Staphylococcus pasteuri		33.04	0.19		1		
Staphylococcus warneri		23.25	0.30		1		
340413	**Very good identification**						**340413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.73				
340433	**Good identification**						**340433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	XYL 1			
340440	**Low discrimination**						**340440**
					NOV	YEP	
Staphylococcus warneri		88.68	0.69	SUC 95	1	43	
Staphylococcus cohnii subsp. cohnii		6.07	0.41		99	12	
Staphylococcus pasteuri		3.92	0.27		1	86	
340441	**Low discrimination**						**340441**
					NOV		
Staphylococcus cohnii subsp. urealyticum		56.40	0.47	MAN 91	99		
Staphylococcus cohnii subsp. cohnii		33.53	0.41	LAC 91	99		
Staphylococcus warneri		9.23	0.35		1		
340442	**Good identification**						**340442**
					NOV		
Staphylococcus warneri		90.37	0.60	SUC 95	1		
Staphylococcus pasteuri		4.89	0.19		1		
Staphylococcus cohnii subsp. urealyticum		3.46	0.26		99		
340443	**Very good identification**						**340443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.46	0.67	MAN 91	99		
Staphylococcus warneri		0.54	0.25		1		
340450	**Low discrimination**						**340450**
					NOV	YEP	
Staphylococcus warneri		64.15	0.66	SUC 95	1	43	
Staphylococcus pasteuri		24.05	0.43		1	86	
Staphylococcus cohnii subsp. cohnii		7.72	0.43		99	12	
340451	**Good identification**						**340451**
					NOV		
Staphylococcus cohnii subsp. urealyticum		91.85	0.67	LAC 91	99		
Staphylococcus cohnii subsp. cohnii		6.87	0.43		99		
Staphylococcus warneri		1.08	0.32		1		
340452	**Low discrimination**						**340452**
					NOV		
Staphylococcus warneri		40.35	0.57	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		21.54	0.47		99		
Staphylococcus saprophyticus subsp. saprophyticus		18.90	0.29		99		
340453	**Very good identification**						**340453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.94	0.88		99		
Staphylococcus warneri		0.04	0.22		1		
Staphylococcus pasteuri		0.01	-.06		1		
340460	**Good identification**						**340460**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		100.0	0.28	SUC 95			
340473	**Good identification**						**340473**
Staphylococcus cohnii subsp. urealyticum		97.31	0.47	XYL 1			
Staphylococcus xylosus		2.69	0.06				
340500	**Low discrimination**						**340500**
					NOV		
Staphylococcus warneri		80.10	0.33	SUC 95	1		
Staphylococcus cohnii subsp. cohnii		11.58	0.11	MLT 95	99		
Kocuria varians		8.33	0.05		1		
340503	**Good identification**						**340503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	MAN 91			
340510	**Low discrimination**						**340510**
					NOV		
Staphylococcus warneri		79.74	0.30	SUC 95	1		
Staphylococcus cohnii subsp. cohnii		20.26	0.13	MLT 95	99		
340511	**Good identification**						**340511**
Staphylococcus cohnii subsp. urealyticum		96.64	0.44	LAC 91			
Staphylococcus cohnii subsp. cohnii		3.36	0.13				
340513	**Very good identification**						**340513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.65				
340540	**Very good identification**						**340540**
					NOV		
Staphylococcus warneri		97.65	0.59	SUC 95	1		
Staphylococcus cohnii subsp. cohnii		2.35	0.21		99		
340541	**Low discrimination**						**340541**
					NOV		
Staphylococcus cohnii subsp. urealyticum		78.45	0.39	MAN 91	99		
Staphylococcus cohnii subsp. cohnii		12.09	0.22	LAC 91	99		
Staphylococcus warneri		9.46	0.24		1		
340542	**Good identification**						**340542**
					NOV		
Staphylococcus warneri		95.07	0.49	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		4.93	0.19		99		
340543	**Very good identification**						**340543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.60	0.60	MAN 91	99		
Staphylococcus warneri		0.40	0.14		1		
340550	**Low discrimination**						**340550**
					NOV	PYR	
Staphylococcus warneri		85.26	0.56	SUC 95	1	8	
Staphylococcus cohnii subsp. cohnii		3.61	0.23		99	1	
Staphylococcus saprophyticus subsp. bovis		3.56	0.18		99	99	
340551	**Very good identification**						**340551**
					NOV		
Staphylococcus cohnii subsp. urealyticum		96.67	0.60	LAC 91	99		
Staphylococcus cohnii subsp. cohnii		1.88	0.24		99		
Staphylococcus warneri		0.83	0.21		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
340552	**Low discrimination**					**340552**
					NOV	
Staphylococcus warneri		49.93	0.46	SUC 95	1	
Staphylococcus cohnii subsp. urealyticum		36.17	0.39		99	
Staphylococcus saprophyticus subsp. saprophyticus		6.44	0.07		99	
340553	**Very good identification**					**340553**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.91	0.80		99	
Staphylococcus xylosus		0.06	0.05		99	
Staphylococcus warneri		0.03	0.12		1	
340573	**Low discrimination**					**340573**
Staphylococcus cohnii subsp. urealyticum		60.51	0.39	XYL 1		
Staphylococcus xylosus		39.49	0.26			
340600	**Low discrimination**					**340600**
					PYR	
Staphylococcus warneri		57.94	0.69	MLT 95	8	
Staphylococcus pasteuri		41.27	0.52		1	
Staphylococcus petrasii subsp. croceilyticus		0.80	0.05		90	
340602	**Low discrimination**					**340602**
Staphylococcus warneri		53.40	0.60	MLT 95		
Staphylococcus pasteuri		46.60	0.44			
340610	**Low discrimination**					**340610**
					NOV	
Staphylococcus pasteuri		85.77	0.68	MLT 86	1	
Staphylococcus warneri		14.20	0.66		1	
Staphylococcus saprophyticus subsp. saprophyticus		0.04	-.07		99	
340611	**Low discrimination**					**340611**
					NOV	
Staphylococcus pasteuri		67.11	0.27	MLT 86	1	
Staphylococcus warneri		22.44	0.32	MNS 1	1	
Staphylococcus cohnii subsp. urealyticum		10.46	0.21		99	
340612	**Low discrimination**					**340612**
					NOV	
Staphylococcus pasteuri		87.60	0.60	MLT 86	1	
Staphylococcus warneri		11.83	0.57		1	
Staphylococcus saprophyticus subsp. saprophyticus		0.57	0.09		99	
340613	**Low discrimination**					**340613**
					NOV	
Staphylococcus cohnii subsp. urealyticum		73.97	0.41	SUC 3	99	
Staphylococcus pasteuri		19.18	0.19		1	
Staphylococcus warneri		5.23	0.22		1	
340620	**Low discrimination**					**340620**
Staphylococcus warneri		58.40	0.28	XYL 1		
Staphylococcus pasteuri		41.60	0.11	MLT 95		
340630	**Low discrimination**					**340630**
Staphylococcus pasteuri		85.80	0.27	XYL 1		
Staphylococcus warneri		14.20	0.25	MLT 86		
340640	**Low discrimination**					**340640**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR			
Staphylococcus warneri		76.60	0.96		8			
Staphylococcus pasteuri		17.64	0.68		1			
Staphylococcus petrasii subsp. croceilyticus		5.48	0.47		90			
340641	**Low discrimination**							**340641**
					OXI	PYR		
Staphylococcus warneri		61.03	0.61	MNS 2	1	8		
Kocuria kristinae		28.96	0.39		99	93		
Staphylococcus pasteuri		6.96	0.27		1	1		
340642	**Low discrimination**							**340642**
					NOV			
Staphylococcus warneri		77.36	0.86		1			
Staphylococcus pasteuri		21.83	0.60		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.58	0.21		99			
340643	**Low discrimination**							**340643**
					OXI	NOV	PYR	
Staphylococcus warneri		55.63	0.51	MNS 2	1	1	8	
Kocuria kristinae		19.80	0.27		99	1	93	
Staphylococcus cohnii subsp. urealyticum		16.79	0.36		1	99	53	
340650	**Low discrimination**							**340650**
					NOV			
Staphylococcus pasteuri		65.78	0.84		1			
Staphylococcus warneri		33.67	0.93		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.49	0.34		99			
340651	**Low discrimination**							**340651**
					NOV			
Staphylococcus warneri		46.38	0.58	MNS 2	1			
Staphylococcus pasteuri		44.84	0.43		1			
Staphylococcus cohnii subsp. urealyticum		6.45	0.36		99			
340652	**Low discrimination**							**340652**
					NOV			
Staphylococcus pasteuri		65.68	0.76		1			
Staphylococcus warneri		27.44	0.83		1			
Staphylococcus saprophyticus subsp. saprophyticus		6.84	0.50		99			
340653	**Low discrimination**							**340653**
					NOV			
Staphylococcus cohnii subsp. urealyticum		63.80	0.57	SUC 3	99			
Staphylococcus pasteuri		17.93	0.35		1			
Staphylococcus warneri		15.14	0.48		1			
340660	**Low discrimination**							**340660**
					PYR			
Staphylococcus warneri		76.82	0.55	XYL 1	8			
Staphylococcus pasteuri		17.69	0.27		1			
Staphylococcus petrasii subsp. croceilyticus		5.50	0.05		90			
340662	**Low discrimination**							**340662**
Staphylococcus warneri		77.99	0.45	XYL 1				
Staphylococcus pasteuri		22.01	0.19					
340670	**Low discrimination**							**340670**
					NOV			
Staphylococcus pasteuri		65.81	0.43	XYL 1	1			
Staphylococcus warneri		33.69	0.52		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus saprophyticus subsp. saprophyticus		0.49	-.07		99			
340672	**Low discrimination**							**340672**
					NOV			
Staphylococcus pasteuri		65.71	0.35	XYL 1	1			
Staphylococcus warneri		27.45	0.42		1			
Staphylococcus saprophyticus subsp. saprophyticus		6.85	0.09		99			
340673	**Acceptable identification**							**340673**
					NOV			
Staphylococcus xylosus		92.99	0.32	VPT 14	99			
Staphylococcus cohnii subsp. urealyticum		5.47	0.16	NIT 89	99			
Staphylococcus pasteuri		1.54	-.06	GAL 91	1			
340700	**Very good identification**							**340700**
Staphylococcus warneri		97.77	0.59	MLT 95				
Staphylococcus pasteuri		2.23	0.11					
340702	**Good identification**							**340702**
Staphylococcus warneri		97.28	0.49	MLT 95				
Staphylococcus pasteuri		2.72	0.03					
340710	**Low discrimination**							**340710**
Staphylococcus warneri		83.80	0.56	MLT 95				
Staphylococcus pasteuri		16.20	0.27					
340712	**Low discrimination**							**340712**
Staphylococcus warneri		80.85	0.46	MLT 95				
Staphylococcus pasteuri		19.15	0.19					
340713	**Low discrimination**							**340713**
					NOV			
Staphylococcus cohnii subsp. urealyticum		83.17	0.34	SUC 3	99			
Staphylococcus xylosus		12.49	0.08		99			
Staphylococcus warneri		4.34	0.12		1			
340733	**Good identification**							**340733**
Staphylococcus xylosus		100.0	0.30	VPT 14				
340740	**Very good identification**							**340740**
					PYR			
Staphylococcus warneri		98.92	0.85		8			
Staphylococcus pasteuri		0.73	0.27		1			
Staphylococcus petrasii subsp. croceilyticus		0.23	0.05		90			
340741	**Low discrimination**							**340741**
					OXI	NOV	PYR	
Staphylococcus warneri		65.13	0.51	MNS 2	1	1	8	
Kocuria kristinae		32.63	0.30		99	1	93	
Staphylococcus cohnii subsp. urealyticum		0.88	0.08		1	99	53	
340742	**Very good identification**							**340742**
Staphylococcus warneri		98.55	0.76					
Staphylococcus pasteuri		0.89	0.19					
Staphylococcus hominis subsp. hominis		0.29	0.18					
340743	**Low discrimination**							**340743**
					OXI	NOV	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		53.23	0.41	MNS 2	1	1	8	
Staphylococcus cohnii subsp. urealyticum		21.81	0.29		1	99	53	
Kocuria kristinae		20.00	0.17		99	1	93	
340750	**Good identification**							**340750**
					NOV	PYR		
Staphylococcus warneri		93.00	0.82		1	8		
Staphylococcus pasteuri		5.81	0.43		1	1		
Staphylococcus saprophyticus subsp. bovis		0.82	0.31		99	99		
340751	**Low discrimination**							**340751**
					NOV			
Staphylococcus warneri		62.81	0.48	MNS 2	1			
Staphylococcus xylosus		14.82	0.22		99			
Staphylococcus cohnii subsp. urealyticum		11.85	0.29		99			
340752	**Low discrimination**							**340752**
					NOV			
Staphylococcus warneri		86.73	0.73		1			
Staphylococcus pasteuri		6.64	0.35		1			
Staphylococcus saprophyticus subsp. saprophyticus		5.96	0.28		99			
340753	**Low discrimination**							**340753**
					NOV			
Staphylococcus cohnii subsp. urealyticum		64.61	0.49	SUC 3	99			
Staphylococcus xylosus		22.75	0.31		99			
Staphylococcus warneri		11.30	0.38		1			
340760	**Good identification**							**340760**
Staphylococcus warneri		100.0	0.44	XYL 1				
340762	**Good identification**							**340762**
Staphylococcus warneri		100.0	0.34	XYL 1				
340763	**Good identification**							**340763**
Staphylococcus xylosus		100.0	0.28	VPT 14				
340770	**Low discrimination**							**340770**
					NOV			
Staphylococcus warneri		89.46	0.41	XYL 1	1			
Staphylococcus pasteuri		5.59	0.02		1			
Staphylococcus xylosus		4.95	0.02		99			
340771	**Good identification**							**340771**
Staphylococcus xylosus		100.0	0.43	VPT 14				
340772	**Low discrimination**							**340772**
					NOV			
Staphylococcus warneri		58.30	0.32	XYL 1	1			
Staphylococcus xylosus		27.56	0.12		99			
Staphylococcus arlettae		9.68	0.12		99			
340773	**Good identification**							**340773**
Staphylococcus xylosus		99.29	0.53	VPT 14				
Staphylococcus arlettae		0.47	0.14	NIT 89				
Staphylococcus cohnii subsp. urealyticum		0.25	0.08					
341053	**Good identification**							**341053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	TRE 99		
341153	**Good identification**					**341153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
341240	**Low discrimination**					**341240**
					NOV	
Staphylococcus warneri		60.16	0.28	ESL 1	1	
Staphylococcus hominis subsp. novobiosepticus		39.84	0.08	TRE 95	99	
341403	**Good identification**					**341403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91		
341411	**Good identification**					**341411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	LAC 91		
341413	**Very good identification**					**341413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.64			
341440	**Good identification**					**341440**
Staphylococcus warneri		100.0	0.28	ESL 1		
341441	**Good identification**					**341441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	MAN 91		
341443	**Very good identification**					**341443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	MAN 91		
341451	**Very good identification**					**341451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	LAC 91		
341452	**Low discrimination**					**341452**
					NOV	
Staphylococcus cohnii subsp. urealyticum		85.12	0.39	MNS 99	99	
Staphylococcus kloosii		6.90	0.28		99	
Staphylococcus warneri		4.14	0.16		1	
341453	**Very good identification**					**341453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.80			
341473	**Good identification**					**341473**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	XYL 1		
341503	**Good identification**					**341503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91		
341511	**Good identification**					**341511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	LAC 91		
341513	**Very good identification**					**341513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.57			
341541	**Good identification**					**341541**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	MAN 91				
341543	**Good identification**							**341543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	MAN 91				
341550	**Low discrimination**							**341550**
					OXI	NOV		
Staphylococcus kloosii		46.20	0.33	VPT 10	1	99		
Macrococcus equipersicus		25.64	0.10	PHS 99	99	99		
Staphylococcus warneri		14.67	0.15		1	1		
341551	**Good identification**							**341551**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	LAC 91				
341552	**Low discrimination**							**341552**
Staphylococcus cohnii subsp. urealyticum		73.15	0.31	MNS 99				
Staphylococcus kloosii		26.85	0.34					
341553	**Very good identification**							**341553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.72					
341573	**Low discrimination**							**341573**
Staphylococcus cohnii subsp. urealyticum		70.45	0.31	XYL 1				
Staphylococcus xylosus		29.55	0.14					
341600	**Low discrimination**							**341600**
Staphylococcus warneri		58.40	0.28	ESL 1				
Staphylococcus pasteuri		41.60	0.11	MLT 95				
341610	**Low discrimination**							**341610**
Staphylococcus pasteuri		85.80	0.27	ESL 1				
Staphylococcus warneri		14.20	0.25	MLT 86				
341613	**Good identification**							**341613**
Staphylococcus cohnii subsp. urealyticum		100.0	0.33	SUC 3				
341640	**Low discrimination**							**341640**
					PYR			
Staphylococcus warneri		76.82	0.55	ESL 1	8			
Staphylococcus pasteuri		17.69	0.27		1			
Staphylococcus petrasii subsp. croceilyticus		5.50	0.05		90			
341641	**Low discrimination**							**341641**
					OXI	NOV	PYR	
Kocuria kristinae		89.47	0.27	URE 10	99	1	93	
Staphylococcus warneri		7.62	0.20	GLR 1	1	1	8	
Staphylococcus cohnii subsp. urealyticum		2.92	0.07		1	99	53	
341642	**Low discrimination**							**341642**
Staphylococcus warneri		77.99	0.45	ESL 1				
Staphylococcus pasteuri		22.01	0.19					
341643	**Low discrimination**							**341643**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. urealyticum		54.23	0.28	SUC 3	1	99	53	
Kocuria kristinae		41.11	0.15	MAN 91	99	1	93	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		4.67	0.10		1 1 8	
341650	**Low discrimination**					**341650**
					NOV	
Staphylococcus pasteuri		65.81	0.43	ESL 1	1	
Staphylococcus warneri		33.69	0.52		1	
Staphylococcus saprophyticus subsp. saprophyticus		0.49	-.07		99	
341651	**Low discrimination**					**341651**
					NOV	
Staphylococcus cohnii subsp. urealyticum		66.94	0.28	SUC 3	99	
Staphylococcus warneri		12.51	0.17	LAC 91	1	
Staphylococcus pasteuri		12.10	0.02		1	
341652	**Low discrimination**					**341652**
					NOV	
Staphylococcus pasteuri		65.11	0.35	ESL 1	1	
Staphylococcus warneri		27.20	0.42		1	
Staphylococcus saprophyticus subsp. saprophyticus		6.78	0.09		99	
341653	**Good identification**					**341653**
					NOV	
Staphylococcus cohnii subsp. urealyticum		98.34	0.49	SUC 3	99	
Staphylococcus xylosus		0.94	-.02		99	
Staphylococcus pasteuri		0.72	-.06		1	
341740	**Good identification**					**341740**
Staphylococcus warneri		100.0	0.44	ESL 1		
341742	**Good identification**					**341742**
Staphylococcus warneri		100.0	0.34	ESL 1		
341750	**Acceptable identification**					**341750**
Staphylococcus warneri		94.12	0.41	ESL 1		
Staphylococcus pasteuri		5.88	0.02			
341752	**Acceptable identification**					**341752**
Staphylococcus warneri		92.89	0.32	ESL 1		
Staphylococcus pasteuri		7.11	-.06			
341753	**Low discrimination**					**341753**
Staphylococcus cohnii subsp. urealyticum		81.54	0.41	SUC 3		
Staphylococcus xylosus		18.46	0.19			
341771	**Low discrimination**					**341771**
Staphylococcus xylosus		69.40	0.31	VPT 14		
Staphylococcus equorum		30.60	0.25	NIT 89		
341773	**Low discrimination**					**341773**
					YEP	
Staphylococcus xylosus		70.56	0.40	VPT 14	40	
Staphylococcus equorum		28.11	0.34	NIT 89	1	
Staphylococcus arlettae		1.33	0.14		99	
342000	**Low discrimination**					**342000**
					OXI	
Kocuria rosea		64.79	0.40	URE 1	35	
Kocuria varians		20.95	0.39		13	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Dermacoccus nishinomiyaensis		13.53	0.28		93		
342002	**Acceptable identification**						**342002**
Kocuria varians		90.03	0.28	VPT 14			
Kocuria rosea		9.97	-.01	GLR 1			
342020	**Low discrimination**						**342020**
Kocuria rosea		79.05	0.33	URE 1			
Kocuria varians		20.95	0.30				
342040	**Low discrimination**						**342040**
					NOV		
Staphylococcus warneri		45.33	0.35	SUC 95	1		
Kocuria varians		32.95	0.26	TRE 95	1		
Staphylococcus hominis subsp. novobiosepticus		14.06	0.08		99		
342050	**Low discrimination**						**342050**
					NOV	PYR	
Staphylococcus warneri		64.21	0.33	SUC 95	1	8	
Staphylococcus pasteuri		26.39	0.10	TRE 95	1	1	
Staphylococcus saprophyticus subsp. bovis		9.40	0.06		99	99	
342053	**Acceptable identification**						**342053**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.19	0.31	TRE 99	99		
Staphylococcus capitis subsp. ureolyticus		6.81	-.11		1		
342100	**Acceptable identification**						**342100**
					OXI		
Kocuria varians		90.67	0.43	VPT 14	13		
Dermacoccus nishinomiyaensis		7.44	0.13	GLR 1	93		
Kocuria rosea		1.89	-.01		35		
342102	**Good identification**						**342102**
Kocuria varians		100.0	0.31	VPT 14			
342120	**Good identification**						**342120**
Kocuria varians		100.0	0.33	VPT 14			
342140	**Low discrimination**						**342140**
					NOV		
Kocuria varians		45.29	0.29	VPT 14	1		
Staphylococcus hominis subsp. novobiosepticus		38.64	0.18	GLR 1	99		
Staphylococcus warneri		13.12	0.25		1		
342210	**Low discrimination**						**342210**
					YEP		
Staphylococcus pasteuri		84.94	0.35	TRE 99	86		
Staphylococcus warneri		12.82	0.33	MLT 86	43		
Kocuria rosea		2.24	0.01		1		
342212	**Low discrimination**						**342212**
Staphylococcus pasteuri		89.03	0.27	TRE 99			
Staphylococcus warneri		10.97	0.23	MLT 86			
342240	**Low discrimination**						**342240**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		48.16	0.49	GLR 1	99		
Staphylococcus warneri		29.81	0.62		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		10.17	0.47		1		
342241	**Acceptable identification**						**342241**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		90.12	0.37	GLR 1	99		
Staphylococcus warneri		4.55	0.27		1		
Staphylococcus epidermidis		3.77	0.16		1		
342242	**Low discrimination**						**342242**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		50.12	0.44	GLR 1	99		
Staphylococcus hominis subsp. hominis		23.08	0.48		1		
Staphylococcus warneri		19.20	0.52		1		
342243	**Low discrimination**						**342243**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		75.03	0.31	GLR 1	99		
Staphylococcus epidermidis		18.47	0.27		1		
Staphylococcus hominis subsp. hominis		2.82	0.13		1		
342250	**Low discrimination**						**342250**
Staphylococcus pasteuri		65.59	0.51	TRE 99			
Staphylococcus warneri		30.62	0.59				
Staphylococcus hominis subsp. hominis		2.16	0.30				
342252	**Low discrimination**						**342252**
Staphylococcus pasteuri		66.75	0.43	TRE 99			
Staphylococcus warneri		25.43	0.49				
Staphylococcus hominis subsp. hominis		6.31	0.31				
342340	**Low discrimination**						**342340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		86.63	0.59	GLR 1	99		
Staphylococcus hominis subsp. hominis		6.61	0.47		1		
Staphylococcus warneri		5.64	0.51		1		
342341	**Acceptable identification**						**342341**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		91.69	0.47	GLR 1	99		
Staphylococcus epidermidis		7.25	0.32		1		
Staphylococcus hominis subsp. hominis		0.57	0.12		1		
342342	**Low discrimination**						**342342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		78.79	0.53	GLR 1	99		
Staphylococcus hominis subsp. hominis		13.10	0.49		1		
Staphylococcus epidermidis		4.93	0.36		1		
342343	**Low discrimination**						**342343**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		67.43	0.41	GLR 1	99		
Staphylococcus epidermidis		31.35	0.42		1		
Staphylococcus hominis subsp. hominis		0.92	0.14		1		
342350	**Low discrimination**						**342350**
					NOV	PYR	
Staphylococcus warneri		52.24	0.49	TRE 95	1	8	
Staphylococcus saprophyticus subsp. bovis		20.38	0.31		99	99	
Staphylococcus hominis subsp. hominis		12.62	0.30		1	1	
342352	**Low discrimination**						**342352**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus warneri		38.86	0.39	TRE 95	1		
Staphylococcus hominis subsp. hominis		33.06	0.32		1		
Staphylococcus hominis subsp. novobiosepticus		13.44	0.12		99		
342371	**Good identification**						**342371**
Staphylococcus xylosus		100.0	0.34	VPT 14			
342373	**Good identification**						**342373**
Staphylococcus xylosus		100.0	0.43	VPT 14			
342400	**Low discrimination**						**342400**
Staphylococcus warneri		45.34	0.35	SUC 95			
Staphylococcus pasteuri		35.40	0.19	MLT 95			
Kocuria varians		11.30	0.16				
342403	**Good identification**						**342403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91			
342410	**Low discrimination**						**342410**
					NOV	PYR	
Staphylococcus pasteuri		83.88	0.35	SUC 99	1	1	
Staphylococcus warneri		12.66	0.33	MLT 86	1	8	
Staphylococcus saprophyticus subsp. bovis		1.85	0.06		99	99	
342411	**Acceptable identification**						**342411**
Staphylococcus cohnii subsp. urealyticum		94.60	0.36	LAC 91			
Staphylococcus cohnii subsp. cohnii		5.40	0.10				
342412	**Low discrimination**						**342412**
					NOV		
Staphylococcus pasteuri		82.37	0.27	SUC 99	1		
Staphylococcus warneri		10.15	0.23	MLT 86	1		
Staphylococcus cohnii subsp. urealyticum		7.48	0.16		99		
342413	**Very good identification**						**342413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.62	0.57		99		
Staphylococcus simulans		0.38	0.02		1		
342440	**Low discrimination**						**342440**
					PYR		
Staphylococcus warneri		70.61	0.62	SUC 95	8		
Staphylococcus pasteuri		17.83	0.35		1		
Staphylococcus petrasii subsp. croceilyticus		9.05	0.18		90		
342441	**Low discrimination**						**342441**
					NOV		
Staphylococcus cohnii subsp. urealyticum		60.58	0.31	MAN 91	99		
Staphylococcus warneri		24.06	0.27	LAC 91	1		
Staphylococcus cohnii subsp. cohnii		15.36	0.18		99		
342442	**Low discrimination**						**342442**
Staphylococcus warneri		73.34	0.52	SUC 95			
Staphylococcus pasteuri		22.69	0.27				
Staphylococcus hominis subsp. hominis		2.82	0.17				
342443	**Good identification**						**342443**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV			
Staphylococcus cohnii subsp. urealyticum		98.71	0.52	MAN 91	99			
Staphylococcus warneri		1.29	0.17		1			
342450	**Low discrimination**							**342450**
					NOV	PYR		
Staphylococcus pasteuri		54.46	0.51	SUC 99	1	1		
Staphylococcus warneri		25.43	0.59		1	8		
Staphylococcus saprophyticus subsp. bovis		19.39	0.47		99	99		
342451	**Low discrimination**							**342451**
					PYR			
Staphylococcus cohnii subsp. urealyticum		73.74	0.52	LAC 91	53			
Staphylococcus saprophyticus subsp. bovis		19.59	0.34		99			
Staphylococcus cohnii subsp. cohnii		2.35	0.20		1			
342452	**Low discrimination**							**342452**
					NOV	PYR		
Staphylococcus pasteuri		59.28	0.43	SUC 99	1	1		
Staphylococcus warneri		22.59	0.49		1	8		
Staphylococcus saprophyticus subsp. bovis		12.92	0.34		99	99		
342453	**Very good identification**							**342453**
					NOV	PYR		
Staphylococcus cohnii subsp. urealyticum		99.03	0.73		99	53		
Staphylococcus saprophyticus subsp. bovis		0.65	0.22		99	99		
Staphylococcus pasteuri		0.12	0.02		1	1		
342503	**Good identification**							**342503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.29	MAN 91				
342511	**Low discrimination**							**342511**
Staphylococcus cohnii subsp. urealyticum		87.57	0.29	LAC 91				
Staphylococcus saprophyticus subsp. bovis		12.43	0.06					
342513	**Good identification**							**342513**
Staphylococcus cohnii subsp. urealyticum		98.89	0.49					
Staphylococcus xylosus		1.11	0.00					
342540	**Low discrimination**							**342540**
					OXI	NOV	PYR	
Staphylococcus warneri		80.34	0.51	SUC 95	1	1	8	
Macrococcus caseolyticus		9.88	0.14		99	99	80	
Staphylococcus saprophyticus subsp. bovis		5.67	0.18		1	99	99	
342542	**Low discrimination**							**342542**
					OXI	NOV		
Staphylococcus warneri		57.53	0.42	SUC 95	1	1		
Macrococcus caseolyticus		31.84	0.18		99	99		
Staphylococcus hominis subsp. hominis		7.59	0.18		1	1		
342543	**Good identification**							**342543**
Staphylococcus cohnii subsp. urealyticum		98.33	0.44	MAN 91				
Staphylococcus xylosus		1.67	-.01					
342550	**Good identification**							**342550**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		90.04	0.59		99	99		
Staphylococcus warneri		9.32	0.49		1	8		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		0.64	0.10		1	1	
342551	**Low discrimination**						**342551**
Staphylococcus saprophyticus subsp. bovis		68.99	0.47				
Staphylococcus cohnii subsp. urealyticum		27.83	0.44				
Staphylococcus xylosus		2.60	0.14				
342552	**Low discrimination**						**342552**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		83.97	0.47		99	99	
Staphylococcus warneri		11.60	0.39		1	8	
Staphylococcus cohnii subsp. urealyticum		3.46	0.24		99	53	
342553	**Good identification**						**342553**
Staphylococcus cohnii subsp. urealyticum		91.95	0.65				
Staphylococcus saprophyticus subsp. bovis		5.64	0.34				
Staphylococcus xylosus		2.42	0.23				
342571	**Good identification**						**342571**
Staphylococcus xylosus		97.72	0.36	VPT 14			
Staphylococcus saprophyticus subsp. bovis		2.28	0.06	SUC 95			
342573	**Good identification**						**342573**
Staphylococcus xylosus		96.77	0.45	VPT 14			
Staphylococcus cohnii subsp. urealyticum		3.23	0.24	SUC 95			
342600	**Low discrimination**						**342600**
					PYR		
Staphylococcus pasteuri		78.13	0.60	MAN 86	1		
Staphylococcus warneri		19.20	0.62	MLT 86	8		
Staphylococcus petrasii subsp. croceilyticus		2.46	0.18		90		
342602	**Low discrimination**						**342602**
Staphylococcus pasteuri		82.50	0.52	MAN 86			
Staphylococcus warneri		16.55	0.52	MLT 86			
Staphylococcus hominis subsp. hominis		0.64	0.17				
342610	**Very good identification**						**342610**
					NOV	PYR	
Staphylococcus pasteuri		97.10	0.76	MLT 86	1	1	
Staphylococcus warneri		2.81	0.59		1	8	
Staphylococcus saprophyticus subsp. bovis		0.09	0.18		99	99	
342611	**Acceptable identification**						**342611**
					NOV	PYR	
Staphylococcus pasteuri		90.62	0.35	MLT 86	1	1	
Staphylococcus warneri		5.31	0.24	MNS 1	1	8	
Staphylococcus saprophyticus subsp. bovis		2.00	0.06		99	99	
342612	**Very good identification**						**342612**
					YEP		
Staphylococcus pasteuri		97.40	0.68	MLT 86	86		
Staphylococcus warneri		2.30	0.49		43		
Staphylococcus simulans		0.24	0.19		1		
342613	**Low discrimination**						**342613**
					NOV	YEP	
Staphylococcus pasteuri		44.70	0.27	MLT 86	1	86	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		37.11	0.30	MNS 1	1	1	
Staphylococcus cohnii subsp. urealyticum		12.43	0.26		99	64	
342630	**Good identification**						**342630**
Staphylococcus pasteuri		97.18	0.35	XYL 1			
Staphylococcus warneri		2.82	0.18	MLT 86			
342632	**Good identification**						**342632**
Staphylococcus pasteuri		100.0	0.27	XYL 1			
342633	**Good identification**						**342633**
Staphylococcus xylosus		100.0	0.28	VPT 14			
342640	**Low discrimination**						**342640**
					PYR		
Staphylococcus pasteuri		43.20	0.76	MAN 86	1		
Staphylococcus warneri		32.84	0.88		8		
Staphylococcus petrasii subsp. croceilyticus		21.93	0.59		90		
342641	**Low discrimination**						**342641**
					PYR		
Staphylococcus warneri		44.15	0.53	MNS 2	8		
Staphylococcus pasteuri		28.74	0.35		1		
Staphylococcus petrasii subsp. croceilyticus		14.59	0.18		90		
342642	**Low discrimination**						**342642**
Staphylococcus pasteuri		56.79	0.68	MAN 86			
Staphylococcus warneri		35.23	0.78				
Staphylococcus hominis subsp. hominis		7.06	0.58				
342643	**Low discrimination**						**342643**
					PYR		
Staphylococcus warneri		33.30	0.43	MNS 2	8		
Staphylococcus pasteuri		26.56	0.27		1		
Staphylococcus lugdunensis		17.63	0.24		96		
342650	**Good identification**						**342650**
					NOV	PYR	
Staphylococcus pasteuri		90.41	0.92		1	1	
Staphylococcus warneri		8.10	0.85		1	8	
Staphylococcus saprophyticus subsp. bovis		1.30	0.59		99	99	
342651	**Low discrimination**						**342651**
					NOV	PYR	
Staphylococcus pasteuri		63.39	0.51	MNS 1	1	1	
Staphylococcus saprophyticus subsp. bovis		22.57	0.47		99	99	
Staphylococcus warneri		11.48	0.50		1	8	
342652	**Good identification**						**342652**
					NOV	PYR	
Staphylococcus pasteuri		92.10	0.84		1	1	
Staphylococcus warneri		6.74	0.75		1	8	
Staphylococcus saprophyticus subsp. bovis		0.81	0.47		99	99	
342653	**Low discrimination**						**342653**
					NOV	PYR	
Staphylococcus pasteuri		53.64	0.43	MNS 1	1	1	
Staphylococcus cohnii subsp. urealyticum		13.76	0.41		99	53	
Staphylococcus saprophyticus subsp. bovis		11.69	0.34		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
342660	**Low discrimination**						**342660**
					PYR		
Staphylococcus pasteuri		43.27	0.35	MAN 86	1		
Staphylococcus warneri		32.89	0.47	XYL 1	8		
Staphylococcus petrasii subsp. croceilyticus		21.97	0.18		90		
342662	**Low discrimination**						**342662**
Staphylococcus pasteuri		57.32	0.27	MAN 86			
Staphylococcus warneri		35.56	0.37	XYL 1			
Staphylococcus hominis subsp. hominis		7.12	0.17				
342670	**Acceptable identification**						**342670**
					NOV	PYR	
Staphylococcus pasteuri		90.34	0.51	XYL 1	1	1	
Staphylococcus warneri		8.10	0.44		1	8	
Staphylococcus saprophyticus subsp. bovis		1.30	0.18		99	99	
342671	**Acceptable identification**						**342671**
					NOV	PYR	
Staphylococcus xylosus		94.91	0.41	VPT 14	99	65	
Staphylococcus pasteuri		3.31	0.10	GAL 91	1	1	
Staphylococcus saprophyticus subsp. bovis		1.18	0.06		99	99	
342672	**Acceptable identification**						**342672**
					NOV		
Staphylococcus pasteuri		90.72	0.43	XYL 1	1		
Staphylococcus warneri		6.63	0.34		1		
Staphylococcus xylosus		1.85	0.10		99		
342673	**Good identification**						**342673**
					NOV		
Staphylococcus xylosus		99.50	0.51	VPT 14	99		
Staphylococcus pasteuri		0.50	0.02	GAL 91	1		
342700	**Low discrimination**						**342700**
Staphylococcus warneri		85.03	0.51	MLT 95			
Staphylococcus pasteuri		11.07	0.19				
Staphylococcus hominis subsp. hominis		3.18	0.16				
342702	**Low discrimination**						**342702**
Staphylococcus warneri		77.44	0.42	MLT 95			
Staphylococcus pasteuri		12.35	0.11				
Staphylococcus hominis subsp. hominis		10.21	0.18				
342710	**Low discrimination**						**342710**
					NOV	PYR	
Staphylococcus pasteuri		44.26	0.35	GAL 1	1	1	
Staphylococcus warneri		40.09	0.49	MLT 86	1	8	
Staphylococcus saprophyticus subsp. bovis		15.64	0.31		99	99	
342712	**Low discrimination**						**342712**
					NOV	PYR	
Staphylococcus pasteuri		51.14	0.27	GAL 1	1	1	
Staphylococcus warneri		37.80	0.39	MLT 86	1	8	
Staphylococcus saprophyticus subsp. bovis		11.06	0.18		99	99	
342713	**Low discrimination**						**342713**
					NOV		
Staphylococcus xylosus		76.98	0.26	VPT 14	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		11.18	0.18	XYL 92	99		
Staphylococcus simulans		6.77	0.08	MLT 93	1		
342731	**Good identification**						**342731**
Staphylococcus xylosus		100.0	0.39	VPT 14			
342733	**Good identification**						**342733**
Staphylococcus xylosus		100.0	0.48	VPT 14			
342740	**Low discrimination**						**342740**
Staphylococcus warneri		77.34	0.78				
Staphylococcus hominis subsp. hominis		15.09	0.58				
Staphylococcus pasteuri		3.25	0.35				
342741	**Low discrimination**						**342741**
					NOV	PYR	
Staphylococcus warneri		47.71	0.43	MNS 2	1	8	
Staphylococcus petrasii subsp. petrasii		9.95	0.14		1	99	
Staphylococcus xylosus		9.82	0.16		99	65	
342742	**Low discrimination**						**342742**
Staphylococcus warneri		55.75	0.68				
Staphylococcus hominis subsp. hominis		38.31	0.59				
Staphylococcus pasteuri		2.87	0.27				
342750	**Low discrimination**						**342750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		59.16	0.72		99	99	
Staphylococcus warneri		29.10	0.75		1	8	
Staphylococcus pasteuri		10.39	0.51		1	1	
342751	**Low discrimination**						**342751**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		81.16	0.59		99	99	
Staphylococcus xylosus		14.52	0.40		99	65	
Staphylococcus warneri		3.26	0.40		1	8	
342752	**Low discrimination**						**342752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		48.86	0.59		99	99	
Staphylococcus warneri		32.04	0.65		1	8	
Staphylococcus pasteuri		14.02	0.43		1	1	
342753	**Low discrimination**						**342753**
Staphylococcus xylosus		62.82	0.50	VPT 14			
Staphylococcus saprophyticus subsp. bovis		30.86	0.47	XYL 92			
Staphylococcus cohnii subsp. urealyticum		3.89	0.34				
342760	**Low discrimination**						**342760**
					NOV		
Staphylococcus warneri		80.42	0.37	XYL 1	1		
Staphylococcus hominis subsp. hominis		15.69	0.16		1		
Staphylococcus xylosus		3.89	-.04		99		
342761	**Good identification**						**342761**
Staphylococcus xylosus		100.0	0.37	VPT 14			
342762	**Low discrimination**						**342762**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus warneri		47.63	0.27	XYL 1	1		
Staphylococcus hominis subsp. hominis		32.73	0.18		1		
Staphylococcus xylosus		19.65	0.06		99		
342763	**Good identification**						**342763**
Staphylococcus xylosus		100.0	0.47	VPT 14			
342770	**Low discrimination**						**342770**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		45.84	0.31	XYL 1	99	99	
Staphylococcus xylosus		23.57	0.21		99	65	
Staphylococcus warneri		22.55	0.34		1	8	
342771	**Very good identification**						**342771**
					PYR		
Staphylococcus xylosus		99.36	0.62	VPT 14	65		
Staphylococcus saprophyticus subsp. bovis		0.49	0.18		99		
Staphylococcus equorum		0.16	0.06		5		
342772	**Low discrimination**						**342772**
					NOV	PYR	
Staphylococcus xylosus		75.08	0.30	VPT 14	99	65	
Staphylococcus saprophyticus subsp. bovis		12.83	0.18	MNS 99	99	99	
Staphylococcus warneri		8.41	0.24		1	8	
342773	**Very good identification**						**342773**
					PYR		
Staphylococcus xylosus		99.81	0.71	VPT 14	65		
Staphylococcus equorum		0.14	0.14		5		
Staphylococcus saprophyticus subsp. bovis		0.04	0.06		99		
343240	**Acceptable identification**						**343240**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		91.21	0.27	GLR 1	99		
Staphylococcus warneri		6.56	0.21	ESL 8	1		
Staphylococcus hominis subsp. hominis		2.24	0.05		1		
343340	**Good identification**						**343340**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.38	0.37	GLR 1	99		
Staphylococcus hominis subsp. hominis		0.87	0.06	ESL 8	1		
Staphylococcus warneri		0.75	0.10		1		
343342	**Good identification**						**343342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		98.11	0.32	GLR 1	99		
Staphylococcus hominis subsp. hominis		1.90	0.08	ESL 8	1		
343373	**Low discrimination**						**343373**
Staphylococcus xylosus		87.93	0.31	VPT 14			
Staphylococcus equorum		12.07	0.14	TRE 96			
343403	**Good identification**						**343403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	MAN 91			
343411	**Good identification**						**343411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	LAC 91			
343413	**Good identification**						**343413**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.49				
343443	**Good identification**						**343443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91			
343451	**Good identification**						**343451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	LAC 91			
343453	**Very good identification**						**343453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.64				
343513	**Good identification**						**343513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.41				
343543	**Good identification**						**343543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91			
343551	**Low discrimination**						**343551**
Staphylococcus cohnii subsp. urealyticum		88.93	0.36	LAC 91			
Staphylococcus saprophyticus subsp. bovis		5.73	0.06				
Staphylococcus xylosus		5.34	0.02				
343553	**Very good identification**						**343553**
Staphylococcus cohnii subsp. urealyticum		98.34	0.57				
Staphylococcus xylosus		1.66	0.11				
343573	**Low discrimination**						**343573**
Staphylococcus xylosus		86.16	0.33	VPT 14			
Staphylococcus equorum		9.36	0.14	SUC 95			
Staphylococcus cohnii subsp. urealyticum		4.48	0.15				
343610	**Good identification**						**343610**
Staphylococcus pasteuri		97.18	0.35	ESL 1			
Staphylococcus warneri		2.82	0.18	MLT 86			
343612	**Good identification**						**343612**
Staphylococcus pasteuri		100.0	0.27	ESL 1			
343640	**Low discrimination**						**343640**
					PYR		
Staphylococcus pasteuri		43.27	0.35	ESL 1	1		
Staphylococcus warneri		32.89	0.47	MAN 86	8		
Staphylococcus petrasii subsp. croceilyticus		21.97	0.18		90		
343642	**Low discrimination**						**343642**
Staphylococcus pasteuri		57.32	0.27	ESL 1			
Staphylococcus warneri		35.56	0.37	MAN 86			
Staphylococcus hominis subsp. hominis		7.12	0.17				
343650	**Acceptable identification**						**343650**
					NOV	PYR	
Staphylococcus pasteuri		90.58	0.51	ESL 1	1	1	
Staphylococcus warneri		8.12	0.44		1	8	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. bovis		1.30	0.18		99	99	
343652	**Acceptable identification**						**343652**
					NOV	PYR	
Staphylococcus pasteuri		92.43	0.43	ESL 1	1	1	
Staphylococcus warneri		6.76	0.34		1	8	
Staphylococcus saprophyticus subsp. bovis		0.81	0.06		99	99	
343653	**Low discrimination**						**343653**
					NOV		
Staphylococcus cohnii subsp. urealyticum		64.89	0.33	SUC 3	99		
Staphylococcus xylosus		28.54	0.17		99		
Staphylococcus pasteuri		6.57	0.02		1		
343671	**Low discrimination**						**343671**
Staphylococcus xylosus		61.47	0.29	VPT 14			
Staphylococcus equorum		38.53	0.26	GAL 91			
343673	**Low discrimination**						**343673**
Staphylococcus xylosus		63.85	0.38	VPT 14			
Staphylococcus equorum		36.15	0.35	GAL 91			
343731	**Low discrimination**						**343731**
Staphylococcus xylosus		54.84	0.27	VPT 14			
Staphylococcus equorum		45.16	0.26	MLT 93			
343733	**Low discrimination**						**343733**
Staphylococcus xylosus		57.34	0.36	VPT 14			
Staphylococcus equorum		42.66	0.35	MLT 93			
343740	**Low discrimination**						**343740**
Staphylococcus warneri		62.70	0.37	ESL 1			
Rothia mucilaginosa		20.66	0.08				
Staphylococcus hominis subsp. hominis		12.24	0.16				
343742	**Low discrimination**						**343742**
Staphylococcus warneri		59.27	0.27	ESL 1			
Staphylococcus hominis subsp. hominis		40.73	0.18				
343750	**Low discrimination**						**343750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		59.97	0.31	ESL 1	99	99	
Staphylococcus warneri		29.50	0.34		1	8	
Staphylococcus pasteuri		10.53	0.10		1	1	
343751	**Low discrimination**						**343751**
					PYR		
Staphylococcus xylosus		75.42	0.28	VPT 14	65		
Staphylococcus saprophyticus subsp. bovis		17.04	0.18	XYL 92	99		
Staphylococcus equorum		5.49	0.06		5		
343753	**Low discrimination**						**343753**
Staphylococcus xylosus		84.61	0.37	VPT 14			
Staphylococcus cohnii subsp. urealyticum		8.15	0.25	XYL 92			
Staphylococcus equorum		5.57	0.14				
343763	**Acceptable identification**						**343763**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		90.27	0.34	VPT 14		
Staphylococcus equorum		8.09	0.14	MAN 94		
Staphylococcus gallinarum		1.64	-.03			
343771	**Low discrimination**					**343771**
Staphylococcus xylosus		60.96	0.50	VPT 14		
Staphylococcus equorum		38.21	0.47			
Staphylococcus gallinarum		0.83	0.10			
343773	**Low discrimination**					**343773**
Staphylococcus xylosus		63.37	0.59	VPT 14		
Staphylococcus equorum		35.88	0.55			
Staphylococcus gallinarum		0.74	0.18			
344013	**Good identification**					**344013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
344043	**Good identification**					**344043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	TRE 99		
344050	**Good identification**					**344050**
Staphylococcus kloosii		100.0	0.27	VPT 10		
344051	**Good identification**					**344051**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	TRE 99		
344052	**Low discrimination**					**344052**
Staphylococcus kloosii		67.53	0.28	VPT 10		
Staphylococcus cohnii subsp. urealyticum		32.47	0.10	TRE 99		
344053	**Good identification**					**344053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	TRE 99		
344113	**Good identification**					**344113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	TRE 99		
344150	**Good identification**					**344150**
Staphylococcus kloosii		100.0	0.33	VPT 10		
344152	**Good identification**					**344152**
Staphylococcus kloosii		100.0	0.34	VPT 10		
344153	**Good identification**					**344153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	TRE 99		
344240	**Good identification**					**344240**
Staphylococcus warneri		100.0	0.28	PHS 1		
344401	**Acceptable identification**					**344401**
Staphylococcus cohnii subsp. urealyticum		93.45	0.35	MAN 91		
Staphylococcus cohnii subsp. cohnii		6.56	0.10	LAC 91		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
344403	**Very good identification**						**344403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	MAN 91			
344410	**Low discrimination**						**344410**
					YEP		
Staphylococcus kloosii		76.21	0.42	VPT 10	99		
Staphylococcus cohnii subsp. urealyticum		13.17	0.14	MLT 95	64		
Staphylococcus cohnii subsp. cohnii		10.62	0.12		12		
344411	**Very good identification**						**344411**
Staphylococcus cohnii subsp. urealyticum		99.13	0.55	LAC 91			
Staphylococcus cohnii subsp. cohnii		0.88	0.12				
344412	**Low discrimination**						**344412**
Staphylococcus cohnii subsp. urealyticum		61.72	0.35	MNS 99			
Staphylococcus kloosii		38.28	0.43				
344413	**Very good identification**						**344413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.76				
344430	**Good identification**						**344430**
Staphylococcus kloosii		100.0	0.30	VPT 10			
344432	**Good identification**						**344432**
Staphylococcus kloosii		100.0	0.30	VPT 10			
344433	**Good identification**						**344433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	XYL 1			
344440	**Low discrimination**						**344440**
					NOV	YEP	
Staphylococcus warneri		44.87	0.28	PHS 1	1	43	
Staphylococcus cohnii subsp. cohnii		30.08	0.20	SUC 95	99	12	
Staphylococcus kloosii		16.65	0.27		99	99	
344441	**Good identification**						**344441**
Staphylococcus cohnii subsp. urealyticum		96.23	0.50	MAN 91			
Staphylococcus cohnii subsp. cohnii		3.77	0.21	LAC 91			
344442	**Low discrimination**						**344442**
					NOV		
Staphylococcus cohnii subsp. urealyticum		72.02	0.30	MAN 91	99		
Staphylococcus kloosii		15.30	0.28	MNS 99	99		
Staphylococcus warneri		12.69	0.18		1		
344443	**Very good identification**						**344443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.71	MAN 91			
344450	**Good identification**						**344450**
					YEP		
Staphylococcus kloosii		90.76	0.68	VPT 10	99		
Staphylococcus cohnii subsp. urealyticum		4.68	0.30		64		
Staphylococcus cohnii subsp. cohnii		2.11	0.22		12		
344451	**Very good identification**						**344451**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		99.31	0.71	LAC 91	64	
Staphylococcus cohnii subsp. cohnii		0.49	0.23		12	
Staphylococcus kloosii		0.20	0.27		99	
344452	**Low discrimination**					**344452**
					NOV	
Staphylococcus kloosii		67.00	0.69	VPT 10	99	
Staphylococcus cohnii subsp. urealyticum		32.22	0.51		99	
Staphylococcus warneri		0.41	0.16		1	
344453	**Very good identification**					**344453**
Staphylococcus cohnii subsp. urealyticum		99.98	0.92			
Staphylococcus kloosii		0.02	0.28			
344463	**Good identification**					**344463**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	MAN 91		
344470	**Very good identification**					**344470**
Staphylococcus kloosii		100.0	0.56	VPT 10		
344471	**Good identification**					**344471**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	XYL 1		
344472	**Very good identification**					**344472**
Staphylococcus kloosii		98.09	0.57	VPT 10		
Staphylococcus cohnii subsp. urealyticum		1.91	0.10			
344473	**Good identification**					**344473**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	XYL 1		
344501	**Good identification**					**344501**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91		
344503	**Good identification**					**344503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	MAN 91		
344510	**Good identification**					**344510**
Staphylococcus kloosii		96.33	0.48	VPT 10		
Staphylococcus cohnii subsp. urealyticum		3.67	0.07	MLT 95		
344511	**Good identification**					**344511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	LAC 91		
344512	**Low discrimination**					**344512**
Staphylococcus kloosii		73.75	0.49	VPT 10		
Staphylococcus cohnii subsp. urealyticum		26.25	0.28	MLT 95		
344513	**Very good identification**					**344513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.69			
344530	**Good identification**					**344530**
Staphylococcus kloosii		100.0	0.35	VPT 10		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
344532	**Good identification**						**344532**
Staphylococcus kloosii		100.0	0.36	VPT 10			
344533	**Good identification**						**344533**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	XYL 1			
344540	**Low discrimination**						**344540**
					NOV		
Staphylococcus kloosii		69.52	0.33	VPT 10	99		
Staphylococcus warneri		30.48	0.18	MAN 99	1		
344541	**Good identification**						**344541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91			
344543	**Very good identification**						**344543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.63	MAN 91			
344550	**Very good identification**						**344550**
Staphylococcus kloosii		98.24	0.74	VPT 10			
Staphylococcus cohnii subsp. urealyticum		1.12	0.22				
Staphylococcus saprophyticus subsp. bovis		0.33	0.06				
344551	**Very good identification**						**344551**
Staphylococcus cohnii subsp. urealyticum		99.11	0.63	LAC 91			
Staphylococcus kloosii		0.89	0.33				
344552	**Good identification**						**344552**
Staphylococcus kloosii		90.40	0.75	VPT 10			
Staphylococcus cohnii subsp. urealyticum		9.60	0.43				
344553	**Very good identification**						**344553**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		99.86	0.84		99	64	
Staphylococcus kloosii		0.10	0.34		99	99	
Staphylococcus caprae		0.04	-.02		1	1	
344570	**Very good identification**						**344570**
Staphylococcus kloosii		100.0	0.62	VPT 10			
344572	**Very good identification**						**344572**
Staphylococcus kloosii		100.0	0.62	VPT 10			
344573	**Low discrimination**						**344573**
Staphylococcus cohnii subsp. urealyticum		87.22	0.43	XYL 1			
Staphylococcus xylosus		10.70	0.15				
Staphylococcus kloosii		2.08	0.21				
344600	**Low discrimination**						**344600**
Staphylococcus warneri		58.40	0.28	PHS 1			
Staphylococcus pasteuri		41.60	0.11	MLT 95			
344610	**Low discrimination**						**344610**
					NOV		
Staphylococcus pasteuri		82.70	0.27	PHS 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		13.69	0.25	MLT 86	1	
Staphylococcus kloosii		3.61	0.21		99	
344613	**Good identification**					**344613**
Staphylococcus cohnii subsp. urealyticum		100.0	0.45	SUC 3		
344640	**Low discrimination**					**344640**
					PYR	
Staphylococcus warneri		76.82	0.55	PHS 1	8	
Staphylococcus pasteuri		17.69	0.27		1	
Staphylococcus petrasii subsp. croceilyticus		5.50	0.05		90	
344642	**Low discrimination**					**344642**
Staphylococcus warneri		77.99	0.45	PHS 1		
Staphylococcus pasteuri		22.01	0.19			
344643	**Good identification**					**344643**
					NOV	
Staphylococcus cohnii subsp. urealyticum		97.82	0.40	SUC 3	99	
Staphylococcus warneri		2.18	0.10	MAN 91	1	
344650	**Low discrimination**					**344650**
					NOV	
Staphylococcus pasteuri		60.12	0.43	PHS 1	1	
Staphylococcus warneri		30.78	0.52		1	
Staphylococcus kloosii		8.13	0.48		99	
344651	**Acceptable identification**					**344651**
					NOV	
Staphylococcus cohnii subsp. urealyticum		91.30	0.40	SUC 3	99	
Staphylococcus warneri		4.42	0.17	LAC 91	1	
Staphylococcus pasteuri		4.28	0.02		1	
344652	**Low discrimination**					**344652**
					NOV	
Staphylococcus pasteuri		51.70	0.35	PHS 1	1	
Staphylococcus warneri		21.60	0.42		1	
Staphylococcus kloosii		18.53	0.48		99	
344653	**Very good identification**					**344653**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.53	0.61	SUC 3	99	
Staphylococcus xylosus		0.28	-.01		99	
Staphylococcus pasteuri		0.19	-.06		1	
344670	**Low discrimination**					**344670**
					NOV	
Staphylococcus kloosii		68.88	0.35	VPT 10	99	
Staphylococcus pasteuri		20.59	0.02	SUC 9	1	
Staphylococcus warneri		10.54	0.11		1	
344672	**Low discrimination**					**344672**
					NOV	
Staphylococcus kloosii		89.87	0.36	VPT 10	99	
Staphylococcus pasteuri		10.13	-.06	SUC 9	1	
344710	**Low discrimination**					**344710**
					NOV	
Staphylococcus kloosii		61.88	0.27	VPT 10	99	
Staphylococcus warneri		38.12	0.15	SUC 9	1	
344712	**Good identification**					**344712**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.28	VPT 10			
344713	**Good identification**						**344713**
Staphylococcus cohnii subsp. urealyticum		97.25	0.38	SUC 3			
Staphylococcus xylosus		2.75	-.04				
344740	**Good identification**						**344740**
Staphylococcus warneri		100.0	0.44	PHS 1			
344742	**Good identification**						**344742**
Staphylococcus warneri		100.0	0.34	PHS 1			
344743	**Good identification**						**344743**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	SUC 3			
344750	**Low discrimination**						**344750**
					NOV	PYR	
Staphylococcus kloosii		55.90	0.53	VPT 10	99	52	
Staphylococcus warneri		34.44	0.41	SUC 9	1	8	
Staphylococcus saprophyticus subsp. bovis		7.50	0.18		99	99	
344751	**Low discrimination**						**344751**
Staphylococcus cohnii subsp. urealyticum		75.16	0.32	SUC 3			
Staphylococcus xylosus		17.68	0.10	LAC 91			
Staphylococcus saprophyticus subsp. bovis		7.16	0.06				
344752	**Low discrimination**						**344752**
					NOV		
Staphylococcus kloosii		78.85	0.54	VPT 10	99		
Staphylococcus warneri		14.95	0.32	SUC 9	1		
Staphylococcus cohnii subsp. urealyticum		2.62	0.12		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
344753	**Good identification**					**344753**
					NOV	
Staphylococcus cohnii subsp. urealyticum		93.35	0.53	SUC 3	99	
Staphylococcus xylosus		6.18	0.20		99	
Staphylococcus caprae		0.47	-.11		1	
344770	**Good identification**					**344770**
Staphylococcus kloosii		100.0	0.41	VPT 10		
344771	**Good identification**					**344771**
Staphylococcus xylosus		100.0	0.32	VPT 14		
344772	**Low discrimination**					**344772**
Staphylococcus kloosii		77.53	0.42	VPT 10		
Staphylococcus arlettae		14.63	0.15	SUC 9		
Staphylococcus xylosus		7.84	0.00			
344773	**Good identification**					**344773**
Staphylococcus xylosus		96.32	0.41	VPT 14		
Staphylococcus arlettae		2.41	0.18	NIT 89		
Staphylococcus cohnii subsp. urealyticum		1.28	0.12			
345013	**Good identification**					**345013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99		
345050	**Good identification**					**345050**
Staphylococcus kloosii		100.0	0.27	VPT 10		
345052	**Good identification**					**345052**
Staphylococcus kloosii		100.0	0.28	VPT 10		
345053	**Good identification**					**345053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	TRE 99		
345150	**Good identification**					**345150**
Staphylococcus kloosii		100.0	0.33	VPT 10		
345152	**Good identification**					**345152**
Staphylococcus kloosii		100.0	0.34	VPT 10		
345153	**Good identification**					**345153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
345403	**Good identification**					**345403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	MAN 91		
345410	**Acceptable identification**					**345410**
Staphylococcus kloosii		93.71	0.42	VPT 10		
Staphylococcus cohnii subsp. urealyticum		6.30	0.06	MLT 95		
345411	**Good identification**					**345411**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	LAC 91		
345412	**Low discrimination**					**345412**
Staphylococcus kloosii		61.47	0.43	VPT 10		
Staphylococcus cohnii subsp. urealyticum		38.54	0.27	MLT 95		
345413	**Very good identification**					**345413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.68			
345430	**Good identification**					**345430**
Staphylococcus kloosii		100.0	0.30	VPT 10		
345432	**Good identification**					**345432**
Staphylococcus kloosii		100.0	0.30	VPT 10		
345433	**Good identification**					**345433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	XYL 1		
345440	**Good identification**					**345440**
Staphylococcus kloosii		100.0	0.27	VPT 10		
345441	**Good identification**					**345441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	MAN 91		
345443	**Very good identification**					**345443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.63	MAN 91		
345450	**Very good identification**					**345450**
Staphylococcus kloosii		98.04	0.68	VPT 10		
Staphylococcus cohnii subsp. urealyticum		1.96	0.21			
345451	**Very good identification**					**345451**
Staphylococcus cohnii subsp. urealyticum		99.49	0.63	LAC 91		
Staphylococcus kloosii		0.51	0.27			
345452	**Low discrimination**					**345452**
Staphylococcus kloosii		84.25	0.69	VPT 10		
Staphylococcus cohnii subsp. urealyticum		15.75	0.42			
345453	**Very good identification**					**345453**
Staphylococcus cohnii subsp. urealyticum		99.95	0.83			
Staphylococcus kloosii		0.06	0.28			
345470	**Very good identification**					**345470**
Staphylococcus kloosii		100.0	0.56	VPT 10		
345472	**Very good identification**					**345472**
Staphylococcus kloosii		100.0	0.57	VPT 10		
345473	**Good identification**					**345473**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	XYL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
345503	**Good identification**					**345503**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	MAN 91		
345510	**Good identification**					**345510**
	Staphylococcus kloosii	100.0	0.48	VPT 10		
345511	**Good identification**					**345511**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	LAC 91		
345512	**Low discrimination**					**345512**
	Staphylococcus kloosii	87.84	0.49	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	12.16	0.19	MLT 95		
345513	**Very good identification**					**345513**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.60			
345530	**Good identification**					**345530**
	Staphylococcus kloosii	100.0	0.35	VPT 10		
345532	**Good identification**					**345532**
	Staphylococcus kloosii	100.0	0.36	VPT 10		
345540	**Good identification**					**345540**
	Staphylococcus kloosii	100.0	0.33	VPT 10		
345541	**Good identification**					**345541**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.34	MAN 91		
345542	**Low discrimination**					**345542**
	Staphylococcus kloosii	71.21	0.34	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	28.79	0.14	MAN 99		
345543	**Very good identification**					**345543**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.55	MAN 91		
345550	**Very good identification**					**345550**
	Staphylococcus kloosii	99.56	0.74	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	0.44	0.14			
345551	**Very good identification**					**345551**
	Staphylococcus cohnii subsp. urealyticum	97.75	0.55	LAC 91		
	Staphylococcus kloosii	2.26	0.33			
345552	**Very good identification**					**345552**
	Staphylococcus kloosii	96.04	0.75	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	3.97	0.35			
345553	**Very good identification**					**345553**
	Staphylococcus cohnii subsp. urealyticum	99.75	0.76			
	Staphylococcus kloosii	0.25	0.34			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
345570	**Very good identification**					**345570**
Staphylococcus kloosii		100.0	0.62	VPT 10		
345572	**Very good identification**					**345572**
Staphylococcus kloosii		100.0	0.62	VPT 10		
345573	**Low discrimination**					**345573**
Staphylococcus cohnii subsp. urealyticum		87.72	0.35	XYL 1		
Staphylococcus xylosus		6.92	0.03			
Staphylococcus kloosii		5.37	0.21			
345613	**Good identification**					**345613**
Staphylococcus cohnii subsp. urealyticum		100.0	0.37	SUC 3		
345643	**Good identification**					**345643**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	SUC 3		
345650	**Low discrimination**					**345650**
					NOV	
Staphylococcus kloosii		89.85	0.48	VPT 10	99	
Staphylococcus pasteuri		6.71	0.02	SUC 9	1	
Staphylococcus warneri		3.44	0.11		1	
345651	**Good identification**					**345651**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	SUC 3		
345652	**Acceptable identification**					**345652**
					NOV	
Staphylococcus kloosii		92.03	0.48	VPT 10	99	
Staphylococcus cohnii subsp. urealyticum		5.38	0.11	SUC 9	99	
Staphylococcus pasteuri		2.59	-.06		1	
345653	**Good identification**					**345653**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52	SUC 3		
345670	**Good identification**					**345670**
Staphylococcus kloosii		100.0	0.35	VPT 10		
345672	**Good identification**					**345672**
Staphylococcus kloosii		100.0	0.36	VPT 10		
345710	**Good identification**					**345710**
Staphylococcus kloosii		100.0	0.27	VPT 10		
345712	**Good identification**					**345712**
Staphylococcus kloosii		100.0	0.28	VPT 10		
345713	**Good identification**					**345713**
Staphylococcus cohnii subsp. urealyticum		100.0	0.29	SUC 3		
345750	**Very good identification**					**345750**
Staphylococcus kloosii		100.0	0.53	VPT 10		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
345752	**Very good identification**					**345752**
Staphylococcus kloosii		100.0	0.54	VPT 10		
345753	**Good identification**					**345753**
Staphylococcus cohnii subsp. urealyticum		95.92	0.45	SUC 3		
Staphylococcus xylosus		4.08	0.07			
345770	**Good identification**					**345770**
Staphylococcus kloosii		100.0	0.41	VPT 10		
345771	**Low discrimination**					**345771**
Staphylococcus equorum		62.77	0.29	VPT 1		
Staphylococcus xylosus		26.77	0.20	NIT 92		
Staphylococcus gallinarum		10.46	0.10			
345772	**Low discrimination**					**345772**
Staphylococcus kloosii		84.12	0.42	VPT 10		
Staphylococcus arlettae		15.88	0.15	SUC 9		
345773	**Low discrimination**					**345773**
Staphylococcus equorum		59.60	0.37	VPT 1		
Staphylococcus xylosus		28.13	0.29	NIT 92		
Staphylococcus gallinarum		9.46	0.18			
346053	**Good identification**					**346053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
346153	**Low discrimination**					**346153**
					NOV	
Staphylococcus cohnii subsp. urealyticum		81.02	0.28	TRE 99	99	
Staphylococcus caprae		18.98	-.02		1	
346241	**Good identification**					**346241**
					NOV	
Staphylococcus epidermidis		95.92	0.32	GLR 1	1	
Staphylococcus hominis subsp. novobiosepticus		4.09	-.04		99	
346242	**Low discrimination**					**346242**
					NOV	
Staphylococcus epidermidis		86.94	0.36	GLR 1	1	
Staphylococcus hominis subsp. hominis		6.59	0.17		1	
Staphylococcus hominis subsp. novobiosepticus		4.68	0.03		99	
346243	**Good identification**					**346243**
Staphylococcus epidermidis		100.0	0.42	GLR 1		
346340	**Low discrimination**					**346340**
					NOV	
Staphylococcus epidermidis		82.14	0.42	GLR 1	1	
Staphylococcus hominis subsp. novobiosepticus		13.76	0.18		99	
Staphylococcus hominis subsp. hominis		3.21	0.16		1	
346341	**Good identification**					**346341**
					NOV	
Staphylococcus epidermidis		97.80	0.47	GLR 1	1	
Staphylococcus hominis subsp. novobiosepticus		2.21	0.05		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
346342	**Good identification**						**346342**
					NOV		
Staphylococcus epidermidis		95.88	0.52	GLR 1	1		
Staphylococcus hominis subsp. novobiosepticus		2.73	0.12		99		
Staphylococcus hominis subsp. hominis		1.39	0.18		1		
346343	**Very good identification**						**346343**
					NOV		
Staphylococcus epidermidis		99.62	0.58	GLR 1	1		
Staphylococcus hominis subsp. novobiosepticus		0.38	0.00		99		
346373	**Good identification**						**346373**
Staphylococcus xylosus		100.0	0.32	VPT 14			
346403	**Good identification**						**346403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	MAN 91			
346411	**Good identification**						**346411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	LAC 91			
346413	**Very good identification**						**346413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.61				
346441	**Good identification**						**346441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91			
346443	**Very good identification**						**346443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	MAN 91			
346450	**Low discrimination**						**346450**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		77.71	0.34		99	99	
Staphylococcus pasteuri		8.82	0.10		1	1	
Staphylococcus kloosii		4.92	0.27		99	52	
346451	**Very good identification**						**346451**
Staphylococcus cohnii subsp. urealyticum		95.76	0.55	LAC 91			
Staphylococcus saprophyticus subsp. bovis		4.24	0.22				
346452	**Low discrimination**						**346452**
Staphylococcus cohnii subsp. urealyticum		61.24	0.35	MNS 99			
Staphylococcus saprophyticus subsp. bovis		26.55	0.22				
Staphylococcus kloosii		7.29	0.28				
346453	**Very good identification**						**346453**
Staphylococcus cohnii subsp. urealyticum		99.89	0.76				
Staphylococcus saprophyticus subsp. bovis		0.11	0.10				
346473	**Low discrimination**						**346473**
Staphylococcus cohnii subsp. urealyticum		80.75	0.35	XYL 1			
Staphylococcus xylosus		19.26	0.13				
346503	**Good identification**						**346503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
346511	**Good identification**						**346511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	LAC 91			
346513	**Very good identification**						**346513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.53				
346541	**Good identification**						**346541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91			
346543	**Good identification**						**346543**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.88	0.48	MAN 91	99		
Staphylococcus caprae		1.12	-.09		1		
346550	**Good identification**						**346550**
Staphylococcus saprophyticus subsp. bovis		96.45	0.47				
Staphylococcus kloosii		2.97	0.33				
Staphylococcus cohnii subsp. urealyticum		0.59	0.07				
346551	**Low discrimination**						**346551**
Staphylococcus cohnii subsp. urealyticum		69.34	0.48	LAC 91			
Staphylococcus saprophyticus subsp. bovis		28.66	0.34				
Staphylococcus xylosus		1.22	0.03				
346552	**Low discrimination**						**346552**
Staphylococcus saprophyticus subsp. bovis		68.90	0.34				
Staphylococcus cohnii subsp. urealyticum		17.03	0.28				
Staphylococcus kloosii		9.18	0.34				
346553	**Very good identification**						**346553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		96.48	0.69		99		
Staphylococcus caprae		2.06	0.18		1		
Staphylococcus saprophyticus subsp. bovis		0.99	0.22		99		
346573	**Low discrimination**						**346573**
Staphylococcus xylosus		84.91	0.34	VPT 14			
Staphylococcus cohnii subsp. urealyticum		15.09	0.28	SUC 95			
346610	**Good identification**						**346610**
					NOV	PYR	
Staphylococcus pasteuri		95.14	0.35	PHS 1	1	1	
Staphylococcus warneri		2.76	0.18	MLT 86	1	8	
Staphylococcus saprophyticus subsp. bovis		2.10	0.06		99	99	
346612	**Acceptable identification**						**346612**
					YEP		
Staphylococcus pasteuri		92.37	0.27	PHS 1	86		
Staphylococcus simulans		7.63	0.10	MLT 86	1		
346613	**Low discrimination**						**346613**
					NOV		
Staphylococcus cohnii subsp. urealyticum		60.12	0.30	SUC 3	99		
Staphylococcus simulans		39.88	0.20		1		
346640	**Low discrimination**						**346640**
					PYR		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		41.28	0.35	PHS 1	1		
Staphylococcus warneri		31.38	0.47	MAN 86	8		
Staphylococcus petrasii subsp. croceilyticus		20.96	0.18		90		
346642	**Low discrimination**						**346642**
Staphylococcus pasteuri		49.98	0.27	PHS 1			
Staphylococcus warneri		31.01	0.37	MAN 86			
Staphylococcus hominis subsp. hominis		19.02	0.27				
346650	**Low discrimination**						**346650**
					NOV	PYR	
Staphylococcus pasteuri		69.02	0.51	PHS 1	1	1	
Staphylococcus saprophyticus subsp. bovis		24.57	0.47		99	99	
Staphylococcus warneri		6.19	0.44		1	8	
346651	**Low discrimination**						**346651**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		72.02	0.34		99	99	
Staphylococcus cohnii subsp. urealyticum		12.58	0.24		99	53	
Staphylococcus pasteuri		8.18	0.10		1	1	
346652	**Low discrimination**						**346652**
					NOV	PYR	
Staphylococcus pasteuri		76.55	0.43	PHS 1	1	1	
Staphylococcus saprophyticus subsp. bovis		16.68	0.34		99	99	
Staphylococcus warneri		5.60	0.34		1	8	
346653	**Low discrimination**						**346653**
Staphylococcus cohnii subsp. urealyticum		73.55	0.45	SUC 3			
Staphylococcus saprophyticus subsp. bovis		10.41	0.22				
Staphylococcus xylosus		9.46	0.18				
346671	**Good identification**						**346671**
Staphylococcus xylosus		100.0	0.30	VPT 14			
346673	**Good identification**						**346673**
Staphylococcus xylosus		99.25	0.39	VPT 14			
Staphylococcus equorum		0.76	-.03	GAL 91			
346703	**Good identification**						**346703**
					YEP		
Staphylococcus hyicus		98.28	0.31	VPT 1	1		
Staphylococcus chromogenes		1.72	-.02	ARG 96	95		
346731	**Good identification**						**346731**
Staphylococcus xylosus		100.0	0.28	VPT 14			
346733	**Good identification**						**346733**
Staphylococcus xylosus		99.01	0.37	VPT 14			
Staphylococcus equorum		0.99	-.03	MLT 93			
346740	**Low discrimination**						**346740**
					NOV	PYR	
Staphylococcus warneri		49.65	0.37	PHS 1	1	8	
Staphylococcus hominis subsp. hominis		29.66	0.26		1	1	
Staphylococcus saprophyticus subsp. bovis		18.27	0.18		99	99	
346742	**Low discrimination**						**346742**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		54.75	0.28	GLR 1			
Staphylococcus warneri		26.02	0.27	PHS 3			
Staphylococcus epidermidis		12.05	0.11				
346750	**Very good identification**						**346750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		97.14	0.59		99	99	
Staphylococcus warneri		1.93	0.34		1	8	
Staphylococcus pasteuri		0.69	0.10		1	1	
346751	**Low discrimination**						**346751**
Staphylococcus saprophyticus subsp. bovis		81.94	0.47				
Staphylococcus xylosus		16.53	0.29				
Staphylococcus cohnii subsp. urealyticum		1.53	0.17				
346752	**Acceptable identification**						**346752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		94.17	0.47		99	99	
Staphylococcus warneri		2.50	0.24		1	8	
Staphylococcus pasteuri		1.09	0.02		1	1	
346753	**Low discrimination**						**346753**
					COA	NOV	
Staphylococcus xylosus		44.56	0.38	VPT 14	1	99	
Staphylococcus saprophyticus subsp. bovis		19.40	0.34	XYL 92	1	99	
Staphylococcus aureus subsp. aureus		17.63	0.18		99	4	
346763	**Good identification**						**346763**
Staphylococcus xylosus		98.42	0.36	VPT 14			
Staphylococcus gallinarum		1.58	-.03	MAN 94			
346771	**Good identification**						**346771**
Staphylococcus xylosus		97.58	0.51	VPT 14			
Staphylococcus gallinarum		1.18	0.10				
Staphylococcus equorum		0.82	0.09				
346773	**Very good identification**						**346773**
Staphylococcus xylosus		98.23	0.60	VPT 14			
Staphylococcus gallinarum		1.02	0.18				
Staphylococcus equorum		0.75	0.18				
347053	**Good identification**						**347053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99			
347403	**Good identification**						**347403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91			
347411	**Good identification**						**347411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	LAC 91			
347413	**Good identification**						**347413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52				
347443	**Good identification**						**347443**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Staphylococcus cohnii subsp. urealyticum	100.0	0.47	MAN 91		
347450	**Low discrimination**					**347450**
	Staphylococcus kloosii	74.07	0.27	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	25.93	0.06	NIT 1		
347451	**Good identification**					**347451**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.47	LAC 91		
347452	**Low discrimination**					**347452**
	Staphylococcus cohnii subsp. urealyticum	76.56	0.27	MNS 99		
	Staphylococcus kloosii	23.44	0.28			
347453	**Very good identification**					**347453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.68			
347473	**Low discrimination**					**347473**
	Staphylococcus cohnii subsp. urealyticum	80.52	0.27	XYL 1		
	Staphylococcus xylosus	12.34	0.01			
	Staphylococcus equorum	7.13	-.03			
347513	**Good identification**					**347513**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.45			
347543	**Good identification**					**347543**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	MAN 91		
347550	**Low discrimination**					**347550**
	Staphylococcus kloosii	75.27	0.33	VPT 10		
	Staphylococcus saprophyticus subsp. bovis	24.73	0.06	NIT 1		
347551	**Good identification**					**347551**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	LAC 91		
347552	**Low discrimination**					**347552**
	Staphylococcus kloosii	58.10	0.34	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	41.90	0.19	NIT 1		
347553	**Very good identification**					**347553**
	Staphylococcus cohnii subsp. urealyticum	99.68	0.60			
	Staphylococcus xylosus	0.32	-.01			
347653	**Low discrimination**					**347653**
	Staphylococcus cohnii subsp. urealyticum	89.96	0.37	SUC 3		
	Staphylococcus xylosus	7.44	0.05			
	Staphylococcus equorum	2.60	-.03			
347671	**Low discrimination**					**347671**
	Staphylococcus equorum	69.13	0.30	VPT 1		
	Staphylococcus xylosus	20.74	0.18	GAL 91		
	Staphylococcus gallinarum	10.13	0.10			
347673	**Low discrimination**					**347673**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus equorum		67.96	0.38	VPT 1		
Staphylococcus xylosus		22.57	0.27	GAL 91		
Staphylococcus gallinarum		9.48	0.18			
347731	**Low discrimination**					**347731**
Staphylococcus equorum		72.74	0.30	VPT 1		
Staphylococcus xylosus		16.61	0.15	MLT 91		
Staphylococcus gallinarum		10.66	0.10			
347733	**Low discrimination**					**347733**
Staphylococcus equorum		71.83	0.38	VPT 1		
Staphylococcus xylosus		18.15	0.25	MLT 91		
Staphylococcus gallinarum		10.02	0.18			
347761	**Low discrimination**					**347761**
Staphylococcus gallinarum		82.91	0.30	VPT 1		
Staphylococcus xylosus		11.19	0.14	MAN 91		
Staphylococcus equorum		5.90	0.09			
347763	**Low discrimination**					**347763**
Staphylococcus gallinarum		81.19	0.38	VPT 1		
Staphylococcus xylosus		12.74	0.23	MAN 91		
Staphylococcus equorum		6.07	0.18			
347771	**Low discrimination**					**347771**
Staphylococcus gallinarum		52.46	0.51	VPT 1		
Staphylococcus equorum		36.57	0.51			
Staphylococcus xylosus		10.97	0.38			
347773	**Low discrimination**					**347773**
Staphylococcus gallinarum		50.62	0.59	VPT 1		
Staphylococcus equorum		37.07	0.59			
Staphylococcus xylosus		12.31	0.48			
350240	**Good identification**					**350240**
Staphylococcus warneri		100.0	0.28	ORN 1		
350413	**Good identification**					**350413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
350440	**Good identification**					**350440**
Staphylococcus warneri		100.0	0.28	ORN 1		
350453	**Good identification**					**350453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	ORN 1		
350553	**Good identification**					**350553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ORN 1		
350600	**Low discrimination**					**350600**
Staphylococcus warneri		58.40	0.28	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus pasteuri		41.60	0.11	MLT 95		
350610	**Low discrimination**					**350610**
Staphylococcus pasteuri		85.80	0.27	ORN 1		
Staphylococcus warneri		14.20	0.25	MLT 86		
350640	**Low discrimination**					**350640**
					PYR	
Staphylococcus warneri		76.82	0.55	ORN 1	8	
Staphylococcus pasteuri		17.69	0.27		1	
Staphylococcus petrasii subsp. croceilyticus		5.50	0.05		90	
350642	**Low discrimination**					**350642**
Staphylococcus warneri		77.99	0.45	ORN 1		
Staphylococcus pasteuri		22.01	0.19			
350650	**Low discrimination**					**350650**
					NOV	
Staphylococcus pasteuri		65.81	0.43	ORN 1	1	
Staphylococcus warneri		33.69	0.52		1	
Staphylococcus saprophyticus subsp. saprophyticus		0.49	-.07		99	
350652	**Low discrimination**					**350652**
					NOV	
Staphylococcus pasteuri		65.71	0.35	ORN 1	1	
Staphylococcus warneri		27.45	0.42		1	
Staphylococcus saprophyticus subsp. saprophyticus		6.85	0.09		99	
350740	**Good identification**					**350740**
Staphylococcus warneri		100.0	0.44	ORN 1		
350742	**Good identification**					**350742**
Staphylococcus warneri		100.0	0.34	ORN 1		
350750	**Acceptable identification**					**350750**
Staphylococcus warneri		94.12	0.41	ORN 1		
Staphylococcus pasteuri		5.88	0.02			
350752	**Acceptable identification**					**350752**
Staphylococcus warneri		92.89	0.32	ORN 1		
Staphylococcus pasteuri		7.11	-.06			
351453	**Good identification**					**351453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ORN 1		
351553	**Good identification**					**351553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
352453	**Good identification**					**352453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
352610	**Good identification**					**352610**
Staphylococcus pasteuri		97.18	0.35	ORN 1		
Staphylococcus warneri		2.82	0.18	MLT 86		
352612	**Good identification**					**352612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		100.0	0.27	ORN 1			
352640	**Low discrimination**						**352640**
					PYR		
Staphylococcus pasteuri		42.71	0.35	ORN 1	1		
Staphylococcus warneri		32.47	0.47	MAN 86	8		
Staphylococcus petrasii subsp. croceilyticus		21.68	0.18		90		
352641	**Good identification**						**352641**
					PYR		
Staphylococcus lugdunensis		99.48	0.46	GLR 2	96		
Staphylococcus warneri		0.52	0.12	LAC 90	8		
352642	**Low discrimination**						**352642**
					PYR		
Staphylococcus pasteuri		41.52	0.27	ORN 1	1		
Staphylococcus lugdunensis		27.56	0.24	MAN 86	96		
Staphylococcus warneri		25.76	0.37		8		
352643	**Very good identification**						**352643**
Staphylococcus lugdunensis		100.0	0.65	GLR 2			
352650	**Acceptable identification**						**352650**
					NOV	PYR	
Staphylococcus pasteuri		90.58	0.51	ORN 1	1	1	
Staphylococcus warneri		8.12	0.44		1	8	
Staphylococcus saprophyticus subsp. bovis		1.30	0.18		99	99	
352652	**Acceptable identification**						**352652**
					NOV	PYR	
Staphylococcus pasteuri		92.43	0.43	ORN 1	1	1	
Staphylococcus warneri		6.76	0.34		1	8	
Staphylococcus saprophyticus subsp. bovis		0.81	0.06		99	99	
352740	**Low discrimination**						**352740**
					PYR		
Staphylococcus warneri		75.72	0.37	ORN 1	8		
Staphylococcus hominis subsp. hominis		14.78	0.16		1		
Staphylococcus lugdunensis		9.50	0.05		96		
352741	**Good identification**						**352741**
Staphylococcus lugdunensis		100.0	0.46	GLR 2			
352743	**Very good identification**						**352743**
Staphylococcus lugdunensis		100.0	0.65	GLR 2			
352750	**Low discrimination**						**352750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		59.97	0.31	ORN 1	99	99	
Staphylococcus warneri		29.50	0.34		1	8	
Staphylococcus pasteuri		10.53	0.10		1	1	
352773	**Good identification**						**352773**
Staphylococcus xylosus		100.0	0.30	VPT 14			
354413	**Good identification**						**354413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1			
354443	**Good identification**						**354443**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
354450	**Good identification**					**354450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
354451	**Good identification**					**354451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
354452	**Low discrimination**					**354452**
Staphylococcus kloosii		67.53	0.28	VPT 10		
Staphylococcus cohnii subsp. urealyticum		32.47	0.10	ORN 1		
354453	**Good identification**					**354453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	ORN 1		
354513	**Good identification**					**354513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ORN 1		
354550	**Good identification**					**354550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
354552	**Good identification**					**354552**
Staphylococcus kloosii		100.0	0.34	VPT 10		
354553	**Good identification**					**354553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ORN 1		
355413	**Good identification**					**355413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1		
355450	**Good identification**					**355450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
355452	**Good identification**					**355452**
Staphylococcus kloosii		100.0	0.28	VPT 10		
355453	**Good identification**					**355453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	ORN 1		
355550	**Good identification**					**355550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
355552	**Good identification**					**355552**
Staphylococcus kloosii		100.0	0.34	VPT 10		
355553	**Good identification**					**355553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1		
356341	**Good identification**					**356341**
Staphylococcus epidermidis		100.0	0.29	ORN 12		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
356342	**Good identification**					**356342**
	Staphylococcus epidermidis	100.0	0.34	ORN 12		
356343	**Good identification**					**356343**
	Staphylococcus epidermidis	100.0	0.40	ORN 12		
356453	**Good identification**					**356453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.35	ORN 1		
356553	**Good identification**					**356553**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.28	ORN 1		
357453	**Good identification**					**357453**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.27	ORN 1		
360013	**Good identification**					**360013**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.35	TRE 99		
360040	**Acceptable identification**					**360040**
	Staphylococcus warneri	92.16	0.31	SUC 95		
	Kocuria varians	7.84	0.02	TRE 95		
360043	**Good identification**					**360043**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.30	TRE 99		
360050	**Low discrimination**					**360050**
					NOV	
	Staphylococcus warneri	89.29	0.28	SUC 95	1	
	Staphylococcus saprophyticus subsp. saprophyticus	10.71	-.12	TRE 95	99	
360051	**Good identification**					**360051**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.30	TRE 99		
360053	**Good identification**					**360053**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.51	TRE 99		
360113	**Good identification**					**360113**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.28	TRE 99		
360153	**Good identification**					**360153**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.43	TRE 99		
360200	**Good identification**					**360200**
	Staphylococcus warneri	100.0	0.31	TRE 95		
360210	**Good identification**					**360210**
	Staphylococcus warneri	100.0	0.28	TRE 95		
360240	**Very good identification**					**360240**
	Staphylococcus warneri	100.0	0.57	TRE 95		
360242	**Good identification**					**360242**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					NOV	
Staphylococcus warneri		96.83	0.47	TRE 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		3.17	-.04		99	
360250	**Good identification**					**360250**
					NOV	
Staphylococcus warneri		92.68	0.54	TRE 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		5.91	0.09		99	
Staphylococcus pasteuri		1.40	0.02		1	
360340	**Good identification**					**360340**
					NOV	
Staphylococcus warneri		97.13	0.47	TRE 95	1	
Staphylococcus hominis subsp. novobiosepticus		2.87	-.02		99	
360342	**Good identification**					**360342**
					NOV	
Staphylococcus warneri		95.45	0.37	TRE 95	1	
Staphylococcus hominis subsp. novobiosepticus		4.55	-.07		99	
360350	**Good identification**					**360350**
Staphylococcus warneri		100.0	0.44	TRE 95		
360352	**Low discrimination**					**360352**
					NOV	
Staphylococcus warneri		77.00	0.34	TRE 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		23.00	0.02		99	
360371	**Good identification**					**360371**
Staphylococcus xylosus		100.0	0.27	VPT 14		
360373	**Good identification**					**360373**
Staphylococcus xylosus		100.0	0.37	VPT 14		
360400	**Good identification**					**360400**
Staphylococcus warneri		100.0	0.31	SUC 95		
360401	**Good identification**					**360401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91		
360403	**Very good identification**					**360403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	MAN 91		
360410	**Low discrimination**					**360410**
					NOV	
Staphylococcus warneri		67.85	0.28	SUC 95	1	
Staphylococcus cohnii subsp. urealyticum		24.02	0.14	MLT 95	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.14	-.12		99	
360411	**Very good identification**					**360411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	LAC 91		
360412	**Low discrimination**					**360412**
					NOV	
Staphylococcus cohnii subsp. urealyticum		77.95	0.35	MNS 99	99	
Staphylococcus saprophyticus subsp. saprophyticus		14.79	0.03		99	
Staphylococcus warneri		7.26	0.18		1	
360413	**Very good identification**					**360413**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Staphylococcus cohnii subsp. urealyticum	100.0	0.76			
360433	**Good identification**					**360433**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.35	XYL 1		
360440	**Very good identification**					**360440**
					NOV	
	Staphylococcus warneri	97.43	0.57	SUC 95	1	
	Staphylococcus saprophyticus subsp. saprophyticus	1.84	0.01		99	
	Staphylococcus cohnii subsp. urealyticum	0.74	0.09		99	
360441	**Good identification**					**360441**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	97.35	0.50	MAN 91	99	
	Staphylococcus warneri	2.65	0.22	LAC 91	1	
360442	**Low discrimination**					**360442**
					NOV	
	Staphylococcus warneri	64.52	0.47	SUC 95	1	
	Staphylococcus saprophyticus subsp. saprophyticus	20.68	0.16		99	
	Staphylococcus cohnii subsp. urealyticum	14.80	0.30		99	
360443	**Very good identification**					**360443**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	99.91	0.71	MAN 91	99	
	Staphylococcus warneri	0.09	0.12		1	
360450	**Low discrimination**					**360450**
					NOV	
	Staphylococcus warneri	55.99	0.54	SUC 95	1	
	Staphylococcus saprophyticus subsp. saprophyticus	34.98	0.29		99	
	Staphylococcus cohnii subsp. urealyticum	5.91	0.30		99	
360451	**Very good identification**					**360451**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	99.75	0.71	LAC 91	99	
	Staphylococcus warneri	0.20	0.19		1	
	Staphylococcus saprophyticus subsp. saprophyticus	0.06	-.12		99	
360452	**Low discrimination**					**360452**
					NOV	
	Staphylococcus saprophyticus subsp. saprophyticus	71.23	0.45	GLR 2	99	
	Staphylococcus cohnii subsp. urealyticum	21.48	0.51	SUC 91	99	
	Staphylococcus warneri	6.71	0.44		1	
360453	**Very good identification**					**360453**
					NOV	
	Staphylococcus cohnii subsp. urealyticum	99.95	0.92		99	
	Staphylococcus saprophyticus subsp. saprophyticus	0.03	0.03		99	
	Staphylococcus warneri	0.01	0.09		1	
360463	**Good identification**					**360463**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.30	MAN 91		
360471	**Low discrimination**					**360471**
	Staphylococcus cohnii subsp. urealyticum	79.24	0.30	XYL 1		
	Staphylococcus xylosus	20.76	0.09	LAC 91		
360473	**Acceptable identification**					**360473**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		93.13	0.51	XYL 1			
Staphylococcus xylosus		6.87	0.18				
360501	**Good identification**						**360501**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MAN 91			
360503	**Good identification**						**360503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	MAN 91			
360511	**Good identification**						**360511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48	LAC 91			
360512	**Good identification**						**360512**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	MNS 99			
360513	**Very good identification**						**360513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.69				
360533	**Low discrimination**						**360533**
Staphylococcus cohnii subsp. urealyticum		57.40	0.28	XYL 1			
Staphylococcus xylosus		42.60	0.16				
360540	**Good identification**						**360540**
Staphylococcus warneri		100.0	0.47	SUC 95			
360541	**Good identification**						**360541**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.03	0.43	MAN 91	99		
Staphylococcus warneri		1.97	0.12	LAC 91	1		
360542	**Low discrimination**						**360542**
					NOV		
Staphylococcus warneri		71.45	0.37	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		22.24	0.22		99		
Staphylococcus saprophyticus subsp. saprophyticus		6.31	-.06		99		
360543	**Very good identification**						**360543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.63	MAN 91			
360550	**Low discrimination**						**360550**
					NOV	PYR	
Staphylococcus warneri		59.05	0.44	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		14.81	0.22		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		10.16	0.07		99	7	
360551	**Very good identification**						**360551**
Staphylococcus cohnii subsp. urealyticum		98.88	0.63	LAC 91			
Staphylococcus xylosus		0.54	0.08				
Staphylococcus saprophyticus subsp. bovis		0.44	0.09				
360552	**Low discrimination**						**360552**
					NOV		
Staphylococcus cohnii subsp. urealyticum		48.99	0.43	MNS 99	99		
Staphylococcus saprophyticus subsp. saprophyticus		32.96	0.23		99		
Staphylococcus warneri		11.27	0.34		1		
360553	**Very good identification**						**360553**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		99.85	0.84					
Staphylococcus xylosus		0.15	0.17					
360571	**Low discrimination**							**360571**
Staphylococcus xylosus		86.08	0.29	VPT 14				
Staphylococcus cohnii subsp. urealyticum		13.92	0.22	NIT 89				
360573	**Low discrimination**							**360573**
Staphylococcus xylosus		63.51	0.39	VPT 14				
Staphylococcus cohnii subsp. urealyticum		36.49	0.43	NIT 89				
360600	**Very good identification**							**360600**
Staphylococcus warneri		97.20	0.57	MLT 95				
Staphylococcus pasteuri		2.80	0.11					
360602	**Acceptable identification**							**360602**
					NOV			
Staphylococcus warneri		93.63	0.47	MLT 95	1			
Staphylococcus pasteuri		3.30	0.03		1			
Staphylococcus saprophyticus subsp. saprophyticus		3.06	-.04		99			
360610	**Low discrimination**							**360610**
					NOV			
Staphylococcus warneri		76.46	0.54	MLT 95	1			
Staphylococcus pasteuri		18.67	0.27		1			
Staphylococcus saprophyticus subsp. saprophyticus		4.88	0.09		99			
360613	**Low discrimination**							**360613**
					NOV			
Staphylococcus cohnii subsp. urealyticum		86.17	0.45	SUC 3	99			
Staphylococcus simulans		11.35	0.21		1			
Staphylococcus xylosus		1.46	-.01		99			
360640	**Very good identification**							**360640**
					NOV			
Staphylococcus warneri		97.83	0.83		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.98	0.21		99			
Staphylococcus pasteuri		0.91	0.27		1			
360641	**Low discrimination**							**360641**
					OXI	NOV	PYR	
Staphylococcus warneri		59.08	0.48	MNS 2	1	1	8	
Kocuria kristinae		37.39	0.30		99	1	93	
Staphylococcus cohnii subsp. urealyticum		3.53	0.19		1	99	53	
360642	**Low discrimination**							**360642**
					NOV			
Staphylococcus warneri		84.61	0.73		1			
Staphylococcus saprophyticus subsp. saprophyticus		14.43	0.37		99			
Staphylococcus pasteuri		0.97	0.19		1			
360643	**Low discrimination**							**360643**
					OXI	NOV	PYR	
Staphylococcus cohnii subsp. urealyticum		52.28	0.40	SUC 3	1	99	53	
Staphylococcus warneri		28.86	0.39	MAN 91	1	1	8	
Kocuria kristinae		13.70	0.17		99	1	93	
360650	**Low discrimination**							**360650**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		70.64	0.80		1	
Staphylococcus saprophyticus subsp. saprophyticus		23.49	0.50		99	
Staphylococcus pasteuri		5.58	0.43		1	
360651	**Low discrimination**					**360651**
					NOV	
Staphylococcus warneri		44.59	0.46	MNS 2	1	
Staphylococcus cohnii subsp. urealyticum		37.18	0.40		99	
Staphylococcus saprophyticus subsp. saprophyticus		7.34	0.09		99	
360652	**Low discrimination**					**360652**
					NOV	
Staphylococcus saprophyticus subsp. saprophyticus		83.63	0.65	GLR 2	99	
Staphylococcus warneri		14.80	0.71		1	
Staphylococcus pasteuri		1.43	0.35		1	
360653	**Low discrimination**					**360653**
Staphylococcus cohnii subsp. urealyticum		83.18	0.61	SUC 3		
Staphylococcus saprophyticus subsp. saprophyticus		9.20	0.24			
Staphylococcus xylosus		3.31	0.23			
360660	**Good identification**					**360660**
Staphylococcus warneri		100.0	0.42	XYL 1		
360662	**Low discrimination**					**360662**
					NOV	
Staphylococcus warneri		85.43	0.32	XYL 1	1	
Staphylococcus saprophyticus subsp. saprophyticus		14.57	-.04		99	
360670	**Low discrimination**					**360670**
					NOV	
Staphylococcus warneri		70.85	0.39	XYL 1	1	
Staphylococcus saprophyticus subsp. saprophyticus		23.56	0.09		99	
Staphylococcus pasteuri		5.59	0.02		1	
360671	**Good identification**					**360671**
Staphylococcus xylosus		100.0	0.35	VPT 14		
360673	**Good identification**					**360673**
Staphylococcus xylosus		97.84	0.44	VPT 14		
Staphylococcus cohnii subsp. urealyticum		2.16	0.20	NIT 89		
360700	**Good identification**					**360700**
Staphylococcus warneri		100.0	0.47	MLT 95		
360702	**Good identification**					**360702**
Staphylococcus warneri		100.0	0.37	MLT 95		
360710	**Good identification**					**360710**
Staphylococcus warneri		100.0	0.44	MLT 95		
360712	**Low discrimination**					**360712**
					NOV	
Staphylococcus warneri		77.00	0.34	MLT 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		23.00	0.02		99	
360713	**Low discrimination**					**360713**
					NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus cohnii subsp. urealyticum		70.07	0.38	SUC 3	99			
Staphylococcus xylosus		28.06	0.20		99			
Staphylococcus simulans		1.87	-.01		1			
360731	**Good identification**							**360731**
Staphylococcus xylosus		100.0	0.33	VPT 14				
360733	**Good identification**							**360733**
Staphylococcus xylosus		100.0	0.42	VPT 14				
360740	**Very good identification**							**360740**
					NOV			
Staphylococcus warneri		99.73	0.73		1			
Staphylococcus saprophyticus subsp. saprophyticus		0.28	-.01		99			
360741	**Low discrimination**							**360741**
					OXI	NOV	PYR	
Staphylococcus warneri		51.99	0.38	MNS 2	1	1	8	
Kocuria kristinae		34.73	0.20		99	1	93	
Staphylococcus xylosus		9.07	0.09		1	99	65	
360742	**Very good identification**							**360742**
					NOV			
Staphylococcus warneri		95.52	0.63		1			
Staphylococcus saprophyticus subsp. saprophyticus		4.49	0.15		99			
360743	**Low discrimination**							**360743**
					NOV			
Staphylococcus cohnii subsp. urealyticum		45.11	0.32	SUC 3	99			
Staphylococcus xylosus		27.34	0.19	MAN 91	99			
Staphylococcus warneri		18.35	0.28		1			
360750	**Low discrimination**							**360750**
					NOV	PYR		
Staphylococcus warneri		87.19	0.70		1	8		
Staphylococcus saprophyticus subsp. saprophyticus		7.98	0.28		99	7		
Staphylococcus saprophyticus subsp. bovis		4.60	0.34		99	99		
360751	**Low discrimination**							**360751**
					NOV			
Staphylococcus xylosus		57.60	0.34	VPT 14	99			
Staphylococcus cohnii subsp. urealyticum		17.27	0.32	NIT 89	99			
Staphylococcus warneri		15.26	0.35	XYL 92	1			
360752	**Low discrimination**							**360752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		59.57	0.43	GLR 2	99	7		
Staphylococcus warneri		38.27	0.60	GAL 8	1	8		
Staphylococcus saprophyticus subsp. bovis		1.52	0.22		99	99		
360753	**Low discrimination**							**360753**
					NOV			
Staphylococcus cohnii subsp. urealyticum		49.88	0.53	SUC 3	99			
Staphylococcus xylosus		46.84	0.43		99			
Staphylococcus warneri		1.45	0.25		1			
360760	**Good identification**							**360760**
Staphylococcus warneri		100.0	0.32	XYL 1				
360761	**Good identification**							**360761**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		100.0	0.31	VPT 14		
360763	**Good identification**					**360763**
Staphylococcus xylosus		100.0	0.41	VPT 14		
360770	**Low discrimination**					**360770**
					NOV	
Staphylococcus warneri		53.03	0.29	XYL 1	1	
Staphylococcus xylosus		46.98	0.15		99	
360771	**Very good identification**					**360771**
Staphylococcus xylosus		100.0	0.56	VPT 14		
360773	**Very good identification**					**360773**
Staphylococcus xylosus		99.79	0.65	VPT 14		
Staphylococcus arlettae		0.12	0.14	NIT 89		
Staphylococcus cohnii subsp. urealyticum		0.09	0.12			
361013	**Good identification**					**361013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99		
361053	**Good identification**					**361053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	TRE 99		
361153	**Good identification**					**361153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	TRE 99		
361403	**Good identification**					**361403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	MAN 91		
361411	**Good identification**					**361411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	LAC 91		
361412	**Good identification**					**361412**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MNS 99		
361413	**Very good identification**					**361413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.68			
361433	**Good identification**					**361433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	XYL 1		
361441	**Good identification**					**361441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	MAN 91		
361443	**Very good identification**					**361443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.63	MAN 91		
361451	**Very good identification**					**361451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.63	LAC 91		
361452	**Low discrimination**					**361452**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		87.71	0.42	MNS 99		
Staphylococcus saprophyticus subsp. saprophyticus		7.55	0.03			
Staphylococcus kloosii		4.74	0.28			
361453	**Very good identification**					**361453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.83			
361473	**Good identification**					**361473**
Staphylococcus cohnii subsp. urealyticum		95.47	0.42	XYL 1		
Staphylococcus xylosus		4.53	0.06			
361503	**Good identification**					**361503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	MAN 91		
361511	**Good identification**					**361511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	LAC 91		
361513	**Very good identification**					**361513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.60			
361541	**Good identification**					**361541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	MAN 91		
361543	**Very good identification**					**361543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.55	MAN 91		
361550	**Low discrimination**					**361550**
Staphylococcus kloosii		69.55	0.33	VPT 10		
Staphylococcus cohnii subsp. urealyticum		30.45	0.14	PHS 99		
361551	**Very good identification**					**361551**
Staphylococcus cohnii subsp. urealyticum		99.65	0.55	LAC 91		
Staphylococcus xylosus		0.35	-.05			
361552	**Low discrimination**					**361552**
Staphylococcus cohnii subsp. urealyticum		80.34	0.35	MNS 99		
Staphylococcus kloosii		19.66	0.34			
361553	**Very good identification**					**361553**
Staphylococcus cohnii subsp. urealyticum		99.90	0.76			
Staphylococcus xylosus		0.10	0.05			
361613	**Good identification**					**361613**
Staphylococcus cohnii subsp. urealyticum		100.0	0.37	SUC 3		
361640	**Good identification**					**361640**
Staphylococcus warneri		100.0	0.42	ESL 1		
361642	**Low discrimination**					**361642**
					NOV	
Staphylococcus warneri		85.43	0.32	ESL 1	1	
Staphylococcus saprophyticus subsp. saprophyticus		14.57	-.04		99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
361643	**Low discrimination**						**361643**
					OXI	NOV	
Staphylococcus cohnii subsp. urealyticum		85.58	0.31	SUC 3	1	99	
Kocuria kristinae		14.42	0.05	MAN 91	99	1	
361650	**Low discrimination**						**361650**
					NOV		
Staphylococcus warneri		70.85	0.39	ESL 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		23.56	0.09		99		
Staphylococcus pasteuri		5.59	0.02		1		
361651	**Acceptable identification**						**361651**
Staphylococcus cohnii subsp. urealyticum		91.67	0.31	SUC 3			
Staphylococcus xylosus		8.33	0.01	LAC 91			
361653	**Good identification**						**361653**
Staphylococcus cohnii subsp. urealyticum		97.51	0.52	SUC 3			
Staphylococcus xylosus		2.50	0.10				
361673	**Low discrimination**						**361673**
Staphylococcus xylosus		89.62	0.32	VPT 14			
Staphylococcus equorum		7.30	0.11	NIT 89			
Staphylococcus cohnii subsp. urealyticum		3.08	0.11	GAL 91			
361713	**Low discrimination**						**361713**
Staphylococcus cohnii subsp. urealyticum		79.53	0.29	SUC 3			
Staphylococcus xylosus		20.48	0.08				
361733	**Acceptable identification**						**361733**
Staphylococcus xylosus		90.33	0.30	VPT 14			
Staphylococcus equorum		9.67	0.11	NIT 89			
361740	**Good identification**						**361740**
Staphylococcus warneri		100.0	0.32	ESL 1			
361750	**Good identification**						**361750**
Staphylococcus warneri		100.0	0.29	ESL 1			
361753	**Low discrimination**						**361753**
Staphylococcus cohnii subsp. urealyticum		62.36	0.45	SUC 3			
Staphylococcus xylosus		37.64	0.31				
361763	**Good identification**						**361763**
Staphylococcus xylosus		100.0	0.28	VPT 14			
361771	**Acceptable identification**						**361771**
Staphylococcus xylosus		91.73	0.43	VPT 14			
Staphylococcus equorum		8.27	0.23	NIT 89			
361773	**Acceptable identification**						**361773**
					YEP		
Staphylococcus xylosus		92.07	0.53	VPT 14	40		
Staphylococcus equorum		7.50	0.32	NIT 89	1		
Staphylococcus arlettae		0.43	0.14		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
362000	**Acceptable identification**						**362000**
	Kocuria varians	91.43	0.29	VPT 14			
	Kocuria rosea	8.57	-.01	GLR 1			
362053	**Good identification**						**362053**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.35	TRE 99			
362100	**Good identification**						**362100**
	Kocuria varians	100.0	0.33	VPT 14			
362153	**Low discrimination**						**362153**
	Staphylococcus cohnii subsp. urealyticum	77.58	0.28	TRE 99			
	Staphylococcus xylosus	22.43	0.07				
362173	**Good identification**						**362173**
	Staphylococcus xylosus	100.0	0.29	VPT 14			
362240	**Acceptable identification**						**362240**
					NOV		
	Staphylococcus warneri	92.67	0.49	TRE 95	1		
	Staphylococcus hominis subsp. novobiosepticus	6.05	0.08		99		
	Staphylococcus hominis subsp. hominis	1.28	0.05		1		
362242	**Low discrimination**						**362242**
					NOV		
	Staphylococcus warneri	80.61	0.40	TRE 95	1		
	Staphylococcus hominis subsp. novobiosepticus	8.50	0.03		99		
	Staphylococcus epidermidis	6.97	0.09		1		
362250	**Low discrimination**						**362250**
					NOV	PYR	
	Staphylococcus warneri	78.66	0.46	TRE 95	1	8	
	Staphylococcus saprophyticus subsp. bovis	14.54	0.22		99	99	
	Staphylococcus pasteuri	6.81	0.10		1	1	
362252	**Low discrimination**						**362252**
					NOV	PYR	
	Staphylococcus warneri	74.92	0.37	TRE 95	1	8	
	Staphylococcus saprophyticus subsp. bovis	10.38	0.09		99	99	
	Staphylococcus pasteuri	7.94	0.02		1	1	
362273	**Good identification**						**362273**
	Staphylococcus xylosus	100.0	0.35	VPT 14			
362333	**Good identification**						**362333**
	Staphylococcus xylosus	100.0	0.32	VPT 14			
362340	**Low discrimination**						**362340**
					NOV		
	Staphylococcus warneri	53.65	0.39	TRE 95	1		
	Staphylococcus hominis subsp. novobiosepticus	33.27	0.18		99		
	Staphylococcus epidermidis	8.77	0.14		1		
362343	**Low discrimination**						**362343**
					NOV		
	Staphylococcus epidermidis	80.35	0.30	GLR 1	1		
	Staphylococcus xylosus	12.67	0.09		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. novobiosepticus		6.98	0.00		99		
362350	**Low discrimination**						**362350**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		70.07	0.34	TRE 99	99	99	
Staphylococcus warneri		29.93	0.36		1	8	
362353	**Good identification**						**362353**
Staphylococcus xylosus		95.73	0.34	VPT 14			
Staphylococcus saprophyticus subsp. bovis		4.27	0.09	TRE 96			
362363	**Good identification**						**362363**
Staphylococcus xylosus		100.0	0.31	VPT 14			
362371	**Good identification**						**362371**
Staphylococcus xylosus		100.0	0.46	VPT 14			
362373	**Very good identification**						**362373**
Staphylococcus xylosus		100.0	0.55	VPT 14			
362403	**Acceptable identification**						**362403**
					NOV		
Staphylococcus cohnii subsp. urealyticum		94.49	0.40	MAN 91	99		
Staphylococcus simulans		5.51	0.09		1		
362411	**Good identification**						**362411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	LAC 91			
362413	**Very good identification**						**362413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.74	0.61		99		
Staphylococcus simulans		2.26	0.21		1		
362440	**Good identification**						**362440**
					NOV	PYR	
Staphylococcus warneri		96.76	0.49	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		3.24	0.09		99	99	
362441	**Acceptable identification**						**362441**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.79	0.35	MAN 91	99		
Staphylococcus warneri		6.21	0.15	LAC 91	1		
362442	**Acceptable identification**						**362442**
					NOV		
Staphylococcus warneri		91.37	0.40	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		8.63	0.14		99		
362443	**Very good identification**						**362443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.54	0.55	MAN 91	99		
Staphylococcus simulans		0.46	0.02		1		
362450	**Low discrimination**						**362450**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		79.98	0.50		99	99	
Staphylococcus warneri		17.49	0.46		1	8	
Staphylococcus pasteuri		1.51	0.10		1	1	
362451	**Low discrimination**						**362451**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		78.07	0.55	LAC 91			
Staphylococcus saprophyticus subsp. bovis		20.74	0.38				
Staphylococcus xylosus		0.82	0.06				
362452	**Low discrimination**						**362452**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		55.54	0.38		99	99	
Staphylococcus cohnii subsp. urealyticum		21.35	0.35		99	53	
Staphylococcus warneri		16.19	0.37		1	8	
362453	**Very good identification**						**362453**
Staphylococcus cohnii subsp. urealyticum		98.88	0.76				
Staphylococcus saprophyticus subsp. bovis		0.65	0.26				
Staphylococcus xylosus		0.29	0.15				
362471	**Acceptable identification**						**362471**
Staphylococcus xylosus		92.32	0.27	VPT 14			
Staphylococcus cohnii subsp. urealyticum		7.68	0.14	GAL 91			
362473	**Low discrimination**						**362473**
Staphylococcus xylosus		77.18	0.37	VPT 14			
Staphylococcus cohnii subsp. urealyticum		22.82	0.35	GAL 91			
362503	**Good identification**						**362503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91			
362511	**Low discrimination**						**362511**
Staphylococcus cohnii subsp. urealyticum		80.12	0.32	LAC 91			
Staphylococcus saprophyticus subsp. bovis		11.37	0.09				
Staphylococcus xylosus		8.51	0.03				
362513	**Very good identification**						**362513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		96.66	0.53		99		
Staphylococcus xylosus		2.89	0.13		99		
Staphylococcus simulans		0.45	-.01		1		
362533	**Good identification**						**362533**
Staphylococcus xylosus		97.15	0.34	VPT 14			
Staphylococcus cohnii subsp. urealyticum		2.86	0.12	SUC 95			
362540	**Low discrimination**						**362540**
					NOV	PYR	
Staphylococcus warneri		70.24	0.39	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		29.77	0.22		99	99	
362541	**Low discrimination**						**362541**
Staphylococcus cohnii subsp. urealyticum		70.73	0.27	MAN 91			
Staphylococcus saprophyticus subsp. bovis		17.91	0.09	LAC 91			
Staphylococcus xylosus		11.36	0.02				
362542	**Low discrimination**						**362542**
					NOV	PYR	
Staphylococcus warneri		69.16	0.29	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		21.98	0.09		99	99	
Staphylococcus cohnii subsp. urealyticum		8.86	0.07		99	53	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
362543	**Good identification**						**362543**
Staphylococcus cohnii subsp. urealyticum		95.67	0.48	MAN 91			
Staphylococcus xylosus		4.33	0.11				
362550	**Very good identification**						**362550**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		98.21	0.63		99	99	
Staphylococcus warneri		1.70	0.36		1	8	
Staphylococcus cohnii subsp. urealyticum		0.10	0.07		99	53	
362551	**Low discrimination**						**362551**
Staphylococcus saprophyticus subsp. bovis		66.50	0.50				
Staphylococcus cohnii subsp. urealyticum		26.82	0.48				
Staphylococcus xylosus		6.68	0.26				
362552	**Acceptable identification**						**362552**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		93.97	0.50		99	99	
Staphylococcus cohnii subsp. urealyticum		3.87	0.28		99	53	
Staphylococcus warneri		2.16	0.26		1	8	
362553	**Low discrimination**						**362553**
Staphylococcus cohnii subsp. urealyticum		88.39	0.69				
Staphylococcus xylosus		6.19	0.36				
Staphylococcus saprophyticus subsp. bovis		5.42	0.38				
362563	**Good identification**						**362563**
Staphylococcus xylosus		98.10	0.33	VPT 14			
Staphylococcus cohnii subsp. urealyticum		1.91	0.07	SUC 95			
362571	**Good identification**						**362571**
Staphylococcus xylosus		98.79	0.48	VPT 14			
Staphylococcus saprophyticus subsp. bovis		0.86	0.09	SUC 95			
Staphylococcus cohnii subsp. urealyticum		0.35	0.07				
362573	**Very good identification**						**362573**
Staphylococcus xylosus		98.76	0.57	VPT 14			
Staphylococcus cohnii subsp. urealyticum		1.24	0.28	SUC 95			
362600	**Low discrimination**						**362600**
Staphylococcus warneri		85.88	0.49	MLT 95			
Staphylococcus pasteuri		14.12	0.19				
362602	**Low discrimination**						**362602**
					YEP		
Staphylococcus warneri		52.86	0.40	MLT 95	43		
Staphylococcus simulans		36.49	0.27		1		
Staphylococcus pasteuri		10.65	0.11		86		
362603	**Good identification**						**362603**
					NOV		
Staphylococcus simulans		97.84	0.37	VPT 4	1		
Staphylococcus cohnii subsp. urealyticum		2.16	0.09	ARG 92	99		
362610	**Low discrimination**						**362610**
					NOV	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus pasteuri		54.07	0.35	BGA 1	1	1		
Staphylococcus warneri		38.77	0.46	MLT 86	1	8		
Staphylococcus saprophyticus subsp. bovis		7.17	0.22		99	99		
362612	**Low discrimination**							**362612**
					YEP			
Staphylococcus simulans		57.25	0.39	VPT 4	1			
Staphylococcus pasteuri		25.66	0.27	ARG 92	86			
Staphylococcus warneri		15.01	0.37		43			
362613	**Acceptable identification**							**362613**
					NOV			
Staphylococcus simulans		90.97	0.50	VPT 4	1			
Staphylococcus cohnii subsp. urealyticum		5.08	0.30	ARG 92	99			
Staphylococcus xylosus		3.95	0.18		99			
362631	**Good identification**							**362631**
Staphylococcus xylosus		100.0	0.31	VPT 14				
362633	**Good identification**							**362633**
					NOV			
Staphylococcus xylosus		98.02	0.40	VPT 14	99			
Staphylococcus simulans		1.98	0.09	GAL 91	1			
362640	**Good identification**							**362640**
					PYR			
Staphylococcus warneri		91.63	0.76		8			
Staphylococcus pasteuri		4.87	0.35		1			
Staphylococcus petrasii subsp. croceilyticus		2.47	0.18		90			
362641	**Low discrimination**							**362641**
					OXI	NOV	PYR	
Staphylococcus warneri		71.47	0.41	MNS 2	1	1	8	
Kocuria kristinae		7.94	0.06		99	1	93	
Staphylococcus xylosus		7.35	0.07		1	99	65	
362642	**Good identification**							**362642**
					YEP			
Staphylococcus warneri		90.74	0.66		43			
Staphylococcus pasteuri		5.91	0.27		86			
Staphylococcus simulans		1.48	0.19		1			
362643	**Low discrimination**							**362643**
					NOV			
Staphylococcus simulans		44.08	0.30	VPT 4	1			
Staphylococcus warneri		16.46	0.31	ARG 92	1			
Staphylococcus xylosus		14.46	0.17		99			
362650	**Low discrimination**							**362650**
					NOV	PYR		
Staphylococcus warneri		41.27	0.73		1	8		
Staphylococcus saprophyticus subsp. bovis		39.74	0.63		99	99		
Staphylococcus pasteuri		18.61	0.51		1	1		
362651	**Low discrimination**							**362651**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		75.43	0.50		99	99		
Staphylococcus xylosus		14.23	0.32		99	65		
Staphylococcus warneri		6.40	0.38		1	8		
362652	**Low discrimination**							**362652**
					NOV	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		39.93	0.63		1	8	
Staphylococcus saprophyticus subsp. bovis		28.84	0.50		99	99	
Staphylococcus pasteuri		22.06	0.43		1	1	
362653	**Low discrimination**						**362653**
					NOV		
Staphylococcus xylosus		34.98	0.41	VPT 14	99		
Staphylococcus simulans		27.22	0.43	GAL 91	1		
Staphylococcus cohnii subsp. urealyticum		19.17	0.45	XYL 92	99		
362660	**Good identification**						**362660**
Staphylococcus warneri		100.0	0.35	XYL 1			
362661	**Good identification**						**362661**
Staphylococcus xylosus		100.0	0.29	VPT 14			
362663	**Good identification**						**362663**
Staphylococcus xylosus		100.0	0.39	VPT 14			
362670	**Low discrimination**						**362670**
					NOV	PYR	
Staphylococcus warneri		34.06	0.32	XYL 1	1	8	
Staphylococcus saprophyticus subsp. bovis		32.80	0.22		99	99	
Staphylococcus xylosus		17.79	0.13		99	65	
362671	**Very good identification**						**362671**
Staphylococcus xylosus		99.54	0.54	VPT 14			
Staphylococcus saprophyticus subsp. bovis		0.46	0.09	GAL 91			
362673	**Very good identification**						**362673**
					NOV		
Staphylococcus xylosus		99.93	0.63	VPT 14	99		
Staphylococcus simulans		0.07	0.02	GAL 91	1		
362700	**Good identification**						**362700**
Staphylococcus warneri		100.0	0.39	MLT 95			
362702	**Low discrimination**						**362702**
Staphylococcus warneri		84.03	0.29	MLT 95			
Staphylococcus simulans		15.97	0.05				
362710	**Low discrimination**						**362710**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		70.07	0.34	MLT 99	99	99	
Staphylococcus warneri		29.93	0.36		1	8	
362711	**Low discrimination**						**362711**
Staphylococcus xylosus		78.04	0.30	VPT 14			
Staphylococcus saprophyticus subsp. bovis		21.96	0.22	XYL 92			
362713	**Low discrimination**						**362713**
					NOV		
Staphylococcus xylosus		78.30	0.39	VPT 14	99		
Staphylococcus simulans		15.50	0.28	XYL 92	1		
Staphylococcus cohnii subsp. urealyticum		4.26	0.22	MLT 93	99		
362721	**Good identification**						**362721**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		100.0	0.27	VPT 14			
362723	**Good identification**						**362723**
Staphylococcus xylosus		100.0	0.36	VPT 14			
362731	**Good identification**						**362731**
Staphylococcus xylosus		100.0	0.51	VPT 14			
362733	**Very good identification**						**362733**
Staphylococcus xylosus		100.0	0.61	VPT 14			
362740	**Good identification**						**362740**
					NOV	PYR	
Staphylococcus warneri		90.60	0.65		1	8	
Staphylococcus saprophyticus subsp. bovis		8.08	0.34		99	99	
Staphylococcus hominis subsp. hominis		0.71	0.16		1	1	
362741	**Low discrimination**						**362741**
					NOV	PYR	
Staphylococcus xylosus		57.66	0.28	VPT 14	99	65	
Staphylococcus saprophyticus subsp. bovis		19.13	0.22	MAN 94	99	99	
Staphylococcus warneri		17.51	0.31	XYL 92	1	8	
362742	**Low discrimination**						**362742**
					NOV	PYR	
Staphylococcus warneri		89.95	0.56		1	8	
Staphylococcus saprophyticus subsp. bovis		6.02	0.22		99	99	
Staphylococcus hominis subsp. hominis		2.50	0.18		1	1	
362743	**Low discrimination**						**362743**
					NOV		
Staphylococcus xylosus		83.44	0.37	VPT 14	99		
Staphylococcus lugdunensis		4.98	0.12	MAN 94	1		
Staphylococcus cohnii subsp. urealyticum		3.00	0.17	XYL 92	99		
362750	**Good identification**						**362750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		92.18	0.75		99	99	
Staphylococcus warneri		7.56	0.63		1	8	
Staphylococcus xylosus		0.11	0.12		99	65	
362751	**Low discrimination**						**362751**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		67.26	0.63		99	99	
Staphylococcus xylosus		32.08	0.53		99	65	
Staphylococcus warneri		0.45	0.28		1	8	
362752	**Low discrimination**						**362752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		88.29	0.63		99	99	
Staphylococcus warneri		9.65	0.53		1	8	
Staphylococcus xylosus		1.21	0.21		99	65	
362753	**Low discrimination**						**362753**
Staphylococcus xylosus		82.25	0.62	VPT 14			
Staphylococcus saprophyticus subsp. bovis		15.15	0.50	XYL 92			
Staphylococcus cohnii subsp. urealyticum		1.91	0.38				
362761	**Good identification**						**362761**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		100.0	0.50	VPT 14			
362763	**Very good identification**						**362763**
Staphylococcus xylosus		100.0	0.59	VPT 14			
362770	**Low discrimination**						**362770**
					NOV	PYR	
Staphylococcus xylosus		55.89	0.33	VPT 14	99	65	
Staphylococcus saprophyticus subsp. bovis		40.76	0.34	MNS 99	99	99	
Staphylococcus warneri		3.34	0.21		1	8	
362771	**Very good identification**						**362771**
					PYR		
Staphylococcus xylosus		99.78	0.74	VPT 14	65		
Staphylococcus saprophyticus subsp. bovis		0.18	0.22		99		
Staphylococcus equorum		0.03	0.04		5		
362772	**Acceptable identification**						**362772**
					NOV	PYR	
Staphylococcus xylosus		93.37	0.43	VPT 14	99	65	
Staphylococcus saprophyticus subsp. bovis		5.98	0.22	MNS 99	99	99	
Staphylococcus warneri		0.65	0.12		1	8	
362773	**Very good identification**						**362773**
					PYR		
Staphylococcus xylosus		99.96	0.84	VPT 14	65		
Staphylococcus equorum		0.03	0.12		5		
Staphylococcus saprophyticus subsp. bovis		0.02	0.09		99		
363053	**Good identification**						**363053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	TRE 99			
363371	**Good identification**						**363371**
Staphylococcus xylosus		96.99	0.34	VPT 14			
Staphylococcus equorum		3.01	0.04	TRE 96			
363373	**Good identification**						**363373**
Staphylococcus xylosus		97.27	0.43	VPT 14			
Staphylococcus equorum		2.73	0.12	TRE 96			
363403	**Good identification**						**363403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	MAN 91			
363411	**Good identification**						**363411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.32	LAC 91			
363413	**Good identification**						**363413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.52				
363443	**Good identification**						**363443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	MAN 91			
363451	**Good identification**						**363451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	LAC 91			
363452	**Good identification**						**363452**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	MNS 99			
363453	**Very good identification**						**363453**
Staphylococcus cohnii subsp. urealyticum		99.81	0.68				
Staphylococcus xylosus		0.19	0.03				
363513	**Good identification**						**363513**
Staphylococcus cohnii subsp. urealyticum		98.12	0.45				
Staphylococcus xylosus		1.89	0.00				
363543	**Good identification**						**363543**
Staphylococcus cohnii subsp. urealyticum		97.18	0.39	MAN 91			
Staphylococcus xylosus		2.83	-.01				
363551	**Low discrimination**						**363551**
Staphylococcus cohnii subsp. urealyticum		81.67	0.39	LAC 91			
Staphylococcus xylosus		13.07	0.14				
Staphylococcus saprophyticus subsp. bovis		5.26	0.09				
363553	**Very good identification**						**363553**
Staphylococcus cohnii subsp. urealyticum		95.69	0.60				
Staphylococcus xylosus		4.31	0.23				
363571	**Good identification**						**363571**
Staphylococcus xylosus		97.60	0.36	VPT 14			
Staphylococcus equorum		2.40	0.04	SUC 95			
363573	**Good identification**						**363573**
Staphylococcus xylosus		96.00	0.45	VPT 14			
Staphylococcus equorum		2.13	0.12	SUC 95			
Staphylococcus cohnii subsp. urealyticum		1.87	0.19				
363633	**Low discrimination**						**363633**
Staphylococcus xylosus		86.79	0.28	VPT 14			
Staphylococcus equorum		13.21	0.12	GAL 91			
363640	**Good identification**						**363640**
Staphylococcus warneri		100.0	0.35	ESL 1			
363650	**Low discrimination**						**363650**
					NOV	PYR	
Staphylococcus warneri		41.43	0.32	ESL 1	1	8	
Staphylococcus saprophyticus subsp. bovis		39.89	0.22		99	99	
Staphylococcus pasteuri		18.68	0.10		1	1	
363653	**Low discrimination**						**363653**
					NOV		
Staphylococcus xylosus		53.08	0.29	VPT 14	99		
Staphylococcus cohnii subsp. urealyticum		45.25	0.37	GAL 91	99		
Staphylococcus simulans		1.67	0.02	XYL 92	1		
363671	**Low discrimination**						**363671**
Staphylococcus xylosus		88.64	0.41	VPT 14			
Staphylococcus equorum		11.36	0.24	GAL 91			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
363673	**Low discrimination**						**363673**
Staphylococcus xylosus		89.62	0.51	VPT 14			
Staphylococcus equorum		10.38	0.33	GAL 91			
363713	**Acceptable identification**						**363713**
Staphylococcus xylosus		92.19	0.26	VPT 14			
Staphylococcus cohnii subsp. urealyticum		7.81	0.14	XYL 92			
363731	**Low discrimination**						**363731**
Staphylococcus xylosus		85.58	0.39	VPT 14			
Staphylococcus equorum		14.42	0.24	MLT 93			
363733	**Low discrimination**						**363733**
Staphylococcus xylosus		86.79	0.48	VPT 14			
Staphylococcus equorum		13.21	0.33	MLT 93			
363750	**Low discrimination**						**363750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		89.95	0.34	ESL 1	99	99	
Staphylococcus warneri		7.37	0.21		1	8	
Staphylococcus xylosus		2.68	-.01		99	65	
363751	**Acceptable identification**						**363751**
					PYR		
Staphylococcus xylosus		90.11	0.40	VPT 14	65		
Staphylococcus saprophyticus subsp. bovis		7.63	0.22	XYL 92	99		
Staphylococcus equorum		1.34	0.04		5		
363753	**Acceptable identification**						**363753**
Staphylococcus xylosus		94.61	0.50	VPT 14			
Staphylococcus cohnii subsp. urealyticum		3.42	0.29	XYL 92			
Staphylococcus equorum		1.27	0.12				
363761	**Good identification**						**363761**
Staphylococcus xylosus		98.01	0.37	VPT 14			
Staphylococcus equorum		1.99	0.04	MAN 94			
363763	**Good identification**						**363763**
Staphylococcus xylosus		97.87	0.47	VPT 14			
Staphylococcus equorum		1.79	0.12	MAN 94			
Staphylococcus gallinarum		0.34	-.05				
363771	**Low discrimination**						**363771**
Staphylococcus xylosus		88.44	0.62	VPT 14			
Staphylococcus equorum		11.34	0.45				
Staphylococcus gallinarum		0.23	0.07				
363772	**Low discrimination**						**363772**
Staphylococcus xylosus		89.62	0.30	VPT 14			
Staphylococcus equorum		10.38	0.12	MNS 99			
363773	**Low discrimination**						**363773**
Staphylococcus xylosus		89.44	0.71	VPT 14			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus equorum		10.36	0.54			
Staphylococcus gallinarum		0.20	0.15			
364013	**Good identification**					**364013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	TRE 99		
364043	**Good identification**					**364043**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	TRE 99		
364050	**Good identification**					**364050**
Staphylococcus kloosii		100.0	0.27	VPT 10		
364051	**Good identification**					**364051**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	TRE 99		
364052	**Low discrimination**					**364052**
Staphylococcus kloosii		58.10	0.28	VPT 10		
Staphylococcus cohnii subsp. urealyticum		41.90	0.13	TRE 99		
364053	**Very good identification**					**364053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.54	TRE 99		
364113	**Good identification**					**364113**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	TRE 99		
364150	**Good identification**					**364150**
Staphylococcus kloosii		100.0	0.33	VPT 10		
364152	**Low discrimination**					**364152**
Staphylococcus kloosii		86.26	0.34	VPT 10		
Staphylococcus cohnii subsp. urealyticum		13.74	0.06	TRE 99		
364153	**Good identification**					**364153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	TRE 99		
364401	**Good identification**					**364401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	MAN 91		
364403	**Very good identification**					**364403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	MAN 91		
364410	**Low discrimination**					**364410**
Staphylococcus kloosii		79.42	0.42	VPT 10		
Staphylococcus cohnii subsp. urealyticum		20.58	0.18	MLT 95		
364411	**Very good identification**					**364411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	LAC 91		
364412	**Low discrimination**					**364412**
Staphylococcus cohnii subsp. urealyticum		70.75	0.39	MNS 99		
Staphylococcus kloosii		29.26	0.43			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
364413	**Very good identification**					**364413**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.80			
364430	**Good identification**					**364430**
	Staphylococcus kloosii	100.0	0.30	VPT 10		
364432	**Good identification**					**364432**
	Staphylococcus kloosii	100.0	0.30	VPT 10		
364433	**Good identification**					**364433**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.39	XYL 1		
364440	**Low discrimination**					**364440**
					NOV	
	Staphylococcus kloosii	41.15	0.27	VPT 10	99	
	Staphylococcus cohnii subsp. urealyticum	31.13	0.13	MAN 99	99	
	Staphylococcus warneri	27.72	0.16		1	
364441	**Very good identification**					**364441**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.54	MAN 91		
364442	**Low discrimination**					**364442**
	Staphylococcus cohnii subsp. urealyticum	87.60	0.34	MAN 91		
	Staphylococcus kloosii	12.40	0.28	MNS 99		
364443	**Very good identification**					**364443**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.75	MAN 91		
364450	**Good identification**					**364450**
					NOV	
	Staphylococcus kloosii	92.14	0.68	VPT 10	99	
	Staphylococcus cohnii subsp. urealyticum	7.12	0.34		99	
	Staphylococcus warneri	0.45	0.13		1	
364451	**Very good identification**					**364451**
	Staphylococcus cohnii subsp. urealyticum	99.87	0.75	LAC 91		
	Staphylococcus kloosii	0.13	0.27			
364452	**Low discrimination**					**364452**
	Staphylococcus kloosii	57.56	0.69	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	41.51	0.54			
	Staphylococcus saprophyticus subsp. saprophyticus	0.93	0.03			
364453	**Excellent identification**					**364453**
	Staphylococcus cohnii subsp. urealyticum	99.99	0.95			
	Staphylococcus kloosii	0.01	0.28			
364463	**Good identification**					**364463**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.34	MAN 91		
364470	**Very good identification**					**364470**
	Staphylococcus kloosii	100.0	0.56	VPT 10		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
364471	**Good identification**					**364471**
	Staphylococcus cohnii subsp. urealyticum	95.30	0.34	XYL 1		
	Staphylococcus xylosus	4.70	-.03	LAC 91		
364472	**Very good identification**					**364472**
	Staphylococcus kloosii	97.17	0.57	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	2.83	0.13			
364473	**Very good identification**					**364473**
	Staphylococcus cohnii subsp. urealyticum	98.63	0.54	XYL 1		
	Staphylococcus xylosus	1.37	0.07			
364501	**Good identification**					**364501**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.31	MAN 91		
364503	**Good identification**					**364503**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.52	MAN 91		
364510	**Acceptable identification**					**364510**
	Staphylococcus kloosii	94.59	0.48	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	5.41	0.10	MLT 95		
364511	**Good identification**					**364511**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.52	LAC 91		
364512	**Low discrimination**					**364512**
	Staphylococcus kloosii	65.19	0.49	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	34.81	0.31	MLT 95		
364513	**Very good identification**					**364513**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.72			
364530	**Good identification**					**364530**
	Staphylococcus kloosii	100.0	0.35	VPT 10		
364532	**Good identification**					**364532**
	Staphylococcus kloosii	100.0	0.36	VPT 10		
364533	**Low discrimination**					**364533**
	Staphylococcus cohnii subsp. urealyticum	87.75	0.31	XYL 1		
	Staphylococcus xylosus	12.25	0.04			
364540	**Low discrimination**					**364540**
	Staphylococcus kloosii	85.69	0.33	VPT 10		
	Staphylococcus cohnii subsp. urealyticum	14.31	0.05	MAN 99		
364541	**Good identification**					**364541**
	Staphylococcus cohnii subsp. urealyticum	100.0	0.46	MAN 91		
364543	**Very good identification**					**364543**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.67	MAN 91		
364550	**Very good identification**					**364550**
Staphylococcus kloosii		97.84	0.74	VPT 10		
Staphylococcus cohnii subsp. urealyticum		1.67	0.26			
Staphylococcus saprophyticus subsp. bovis		0.49	0.09			
364551	**Very good identification**					**364551**
Staphylococcus cohnii subsp. urealyticum		99.31	0.67	LAC 91		
Staphylococcus kloosii		0.59	0.33			
Staphylococcus xylosus		0.10	-.04			
364552	**Low discrimination**					**364552**
Staphylococcus kloosii		86.26	0.75	VPT 10		
Staphylococcus cohnii subsp. urealyticum		13.74	0.47			
364553	**Very good identification**					**364553**
Staphylococcus cohnii subsp. urealyticum		99.91	0.88			
Staphylococcus kloosii		0.06	0.34			
Staphylococcus xylosus		0.03	0.06			
364570	**Very good identification**					**364570**
Staphylococcus kloosii		100.0	0.62	VPT 10		
364572	**Very good identification**					**364572**
Staphylococcus kloosii		99.36	0.62	VPT 10		
Staphylococcus cohnii subsp. urealyticum		0.64	0.06			
364573	**Low discrimination**					**364573**
Staphylococcus cohnii subsp. urealyticum		74.46	0.47	XYL 1		
Staphylococcus xylosus		24.36	0.27			
Staphylococcus kloosii		1.18	0.21			
364603	**Acceptable identification**					**364603**
					NOV	
Staphylococcus cohnii subsp. urealyticum		93.11	0.28	SUC 3	99	
Staphylococcus simulans		6.89	-.01	MAN 91	1	
364611	**Good identification**					**364611**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	SUC 3		
364613	**Good identification**					**364613**
					NOV	
Staphylococcus cohnii subsp. urealyticum		97.16	0.49	SUC 3	99	
Staphylococcus simulans		2.84	0.12		1	
364640	**Good identification**					**364640**
Staphylococcus warneri		100.0	0.42	PHS 1		
364642	**Low discrimination**					**364642**
					NOV	
Staphylococcus warneri		85.43	0.32	PHS 1	1	
Staphylococcus saprophyticus subsp. saprophyticus		14.57	-.04		99	
364643	**Good identification**					**364643**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	SUC 3			
364650	**Low discrimination**						**364650**
					NOV		
Staphylococcus kloosii		41.09	0.48	VPT 10	99		
Staphylococcus warneri		38.90	0.39	SUC 9	1		
Staphylococcus saprophyticus subsp. saprophyticus		12.93	0.09		99		
364651	**Good identification**						**364651**
Staphylococcus cohnii subsp. urealyticum		97.41	0.44	SUC 3			
Staphylococcus xylosus		2.59	0.02	LAC 91			
364653	**Very good identification**						**364653**
					NOV		
Staphylococcus cohnii subsp. urealyticum		99.03	0.64	SUC 3	99		
Staphylococcus xylosus		0.74	0.11		99		
Staphylococcus simulans		0.23	0.04		1		
364670	**Good identification**						**364670**
Staphylococcus kloosii		100.0	0.35	VPT 10			
364672	**Good identification**						**364672**
Staphylococcus kloosii		100.0	0.36	VPT 10			
364673	**Low discrimination**						**364673**
Staphylococcus xylosus		89.50	0.33	VPT 14			
Staphylococcus cohnii subsp. urealyticum		10.50	0.23	NIT 89			
364710	**Good identification**						**364710**
Staphylococcus kloosii		100.0	0.27	VPT 10			
364712	**Good identification**						**364712**
Staphylococcus kloosii		100.0	0.28	VPT 10			
364713	**Acceptable identification**						**364713**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		90.81	0.41	SUC 3	1	99	
Staphylococcus xylosus		6.84	0.09		1	99	
Staphylococcus intermedius		2.35	0.01		99	1	
364733	**Good identification**						**364733**
Staphylococcus xylosus		100.0	0.31	VPT 14			
364740	**Good identification**						**364740**
Staphylococcus warneri		100.0	0.32	PHS 1			
364743	**Low discrimination**						**364743**
Staphylococcus cohnii subsp. urealyticum		89.77	0.36	SUC 3			
Staphylococcus xylosus		10.23	0.07	MAN 91			
364750	**Low discrimination**						**364750**
					NOV	PYR	
Staphylococcus kloosii		73.78	0.53	VPT 10	99	52	
Staphylococcus saprophyticus subsp. bovis		14.85	0.22	SUC 9	99	99	
Staphylococcus warneri		11.36	0.29		1	8	
364751	**Low discrimination**						**364751**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		58.05	0.36	SUC 3			
Staphylococcus xylosus		36.42	0.23	LAC 91			
Staphylococcus saprophyticus subsp. bovis		5.53	0.09				
364752	**Low discrimination**						**364752**
Staphylococcus kloosii		82.14	0.54	VPT 10			
Staphylococcus saprophyticus subsp. saprophyticus		6.06	0.02	SUC 9			
Staphylococcus cohnii subsp. urealyticum		4.09	0.16				
364753	**Low discrimination**						**364753**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		84.50	0.57	SUC 3	1	99	
Staphylococcus xylosus		14.92	0.32		1	99	
Staphylococcus intermedius		0.58	0.05		99	1	
364763	**Good identification**						**364763**
Staphylococcus xylosus		100.0	0.29	VPT 14			
364770	**Low discrimination**						**364770**
Staphylococcus kloosii		86.66	0.41	VPT 10			
Staphylococcus xylosus		13.34	0.03	SUC 9			
364771	**Good identification**						**364771**
Staphylococcus xylosus		100.0	0.44	VPT 14			
364772	**Low discrimination**						**364772**
Staphylococcus kloosii		62.77	0.42	VPT 10			
Staphylococcus xylosus		25.39	0.13	SUC 9			
Staphylococcus arlettae		11.85	0.15				
364773	**Very good identification**						**364773**
Staphylococcus xylosus		98.89	0.54	VPT 14			
Staphylococcus arlettae		0.62	0.18	NIT 89			
Staphylococcus cohnii subsp. urealyticum		0.49	0.16				
365013	**Good identification**						**365013**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	TRE 99			
365050	**Good identification**						**365050**
Staphylococcus kloosii		100.0	0.27	VPT 10			
365052	**Low discrimination**						**365052**
Staphylococcus kloosii		78.10	0.28	VPT 10			
Staphylococcus cohnii subsp. urealyticum		21.91	0.05	TRE 99			
365053	**Good identification**						**365053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.46	TRE 99			
365150	**Good identification**						**365150**
Staphylococcus kloosii		100.0	0.33	VPT 10			
365152	**Good identification**						**365152**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.34	VPT 10		
365153	**Good identification**					**365153**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	TRE 99		
365401	**Good identification**					**365401**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	MAN 91		
365403	**Good identification**					**365403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	MAN 91		
365410	**Acceptable identification**					**365410**
Staphylococcus kloosii		90.85	0.42	VPT 10		
Staphylococcus cohnii subsp. urealyticum		9.15	0.10	MLT 95		
365411	**Good identification**					**365411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	LAC 91		
365412	**Low discrimination**					**365412**
Staphylococcus kloosii		51.54	0.43	VPT 10		
Staphylococcus cohnii subsp. urealyticum		48.47	0.30	MLT 95		
365413	**Very good identification**					**365413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.71			
365430	**Good identification**					**365430**
Staphylococcus kloosii		100.0	0.30	VPT 10		
365432	**Good identification**					**365432**
Staphylococcus kloosii		100.0	0.30	VPT 10		
365433	**Good identification**					**365433**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	XYL 1		
365440	**Low discrimination**					**365440**
Staphylococcus kloosii		77.27	0.27	VPT 10		
Staphylococcus cohnii subsp. urealyticum		22.74	0.04	MAN 99		
365441	**Good identification**					**365441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.45	MAN 91		
365443	**Very good identification**					**365443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.66	MAN 91		
365450	**Very good identification**					**365450**
Staphylococcus kloosii		97.08	0.68	VPT 10		
Staphylococcus cohnii subsp. urealyticum		2.92	0.25			
365451	**Very good identification**					**365451**
Staphylococcus cohnii subsp. urealyticum		99.66	0.66	LAC 91		
Staphylococcus kloosii		0.34	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
365452	**Low discrimination**					**365452**
Staphylococcus kloosii		78.10	0.69	VPT 10		
Staphylococcus cohnii subsp. urealyticum		21.90	0.46			
365453	**Very good identification**					**365453**
Staphylococcus cohnii subsp. urealyticum		99.96	0.87			
Staphylococcus kloosii		0.04	0.28			
365470	**Very good identification**					**365470**
Staphylococcus kloosii		100.0	0.56	VPT 10		
365472	**Very good identification**					**365472**
Staphylococcus kloosii		98.88	0.57	VPT 10		
Staphylococcus cohnii subsp. urealyticum		1.12	0.05			
365473	**Good identification**					**365473**
Staphylococcus cohnii subsp. urealyticum		100.0	0.46	XYL 1		
365503	**Good identification**					**365503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91		
365510	**Good identification**					**365510**
Staphylococcus kloosii		100.0	0.48	VPT 10		
365511	**Good identification**					**365511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	LAC 91		
365512	**Low discrimination**					**365512**
Staphylococcus kloosii		82.81	0.49	VPT 10		
Staphylococcus cohnii subsp. urealyticum		17.19	0.23	MLT 95		
365513	**Very good identification**					**365513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.64			
365530	**Good identification**					**365530**
Staphylococcus kloosii		100.0	0.35	VPT 10		
365532	**Good identification**					**365532**
Staphylococcus kloosii		100.0	0.36	VPT 10		
365540	**Good identification**					**365540**
Staphylococcus kloosii		100.0	0.33	VPT 10		
365541	**Good identification**					**365541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	MAN 91		
365542	**Low discrimination**					**365542**
Staphylococcus kloosii		62.25	0.34	VPT 10		
Staphylococcus cohnii subsp. urealyticum		37.75	0.18	MAN 99		
365543	**Very good identification**					**365543**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	MAN 91		
365550	**Very good identification**					**365550**
Staphylococcus kloosii		99.34	0.74	VPT 10		
Staphylococcus cohnii subsp. urealyticum		0.66	0.18			
365551	**Very good identification**					**365551**
Staphylococcus cohnii subsp. urealyticum		98.49	0.59	LAC 91		
Staphylococcus kloosii		1.52	0.33			
365552	**Good identification**					**365552**
Staphylococcus kloosii		94.17	0.75	VPT 10		
Staphylococcus cohnii subsp. urealyticum		5.83	0.38			
365553	**Very good identification**					**365553**
Staphylococcus cohnii subsp. urealyticum		99.84	0.79			
Staphylococcus kloosii		0.16	0.34			
365570	**Very good identification**					**365570**
Staphylococcus kloosii		100.0	0.62	VPT 10		
365572	**Very good identification**					**365572**
Staphylococcus kloosii		100.0	0.62	VPT 10		
365573	**Low discrimination**					**365573**
Staphylococcus cohnii subsp. urealyticum		79.93	0.38	XYL 1		
Staphylococcus xylosus		16.81	0.15			
Staphylococcus kloosii		3.26	0.21			
365613	**Good identification**					**365613**
Staphylococcus cohnii subsp. urealyticum		100.0	0.40	SUC 3		
365643	**Good identification**					**365643**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	SUC 3		
365650	**Good identification**					**365650**
Staphylococcus kloosii		100.0	0.48	VPT 10		
365651	**Good identification**					**365651**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	SUC 3		
365652	**Acceptable identification**					**365652**
Staphylococcus kloosii		91.94	0.48	VPT 10		
Staphylococcus cohnii subsp. urealyticum		8.06	0.15	SUC 9		
365653	**Very good identification**					**365653**
Staphylococcus cohnii subsp. urealyticum		99.52	0.56	SUC 3		
Staphylococcus xylosus		0.48	-.01			
365670	**Good identification**					**365670**
Staphylococcus kloosii		100.0	0.35	VPT 10		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
365672	**Good identification**					**365672**
Staphylococcus kloosii		100.0	0.36	VPT 10		
365710	**Good identification**					**365710**
Staphylococcus kloosii		100.0	0.27	VPT 10		
365712	**Good identification**					**365712**
Staphylococcus kloosii		100.0	0.28	VPT 10		
365713	**Good identification**					**365713**
Staphylococcus cohnii subsp. urealyticum		95.38	0.33	SUC 3		
Staphylococcus xylosus		4.62	-.04			
365743	**Good identification**					**365743**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	SUC 3		
365750	**Very good identification**					**365750**
Staphylococcus kloosii		100.0	0.53	VPT 10		
365751	**Low discrimination**					**365751**
Staphylococcus cohnii subsp. urealyticum		71.26	0.28	SUC 3		
Staphylococcus xylosus		28.74	0.10	LAC 91		
365752	**Very good identification**					**365752**
Staphylococcus kloosii		98.10	0.54	VPT 10		
Staphylococcus cohnii subsp. urealyticum		1.90	0.07	SUC 9		
365753	**Low discrimination**					**365753**
Staphylococcus cohnii subsp. urealyticum		89.81	0.48	SUC 3		
Staphylococcus xylosus		10.19	0.20			
365770	**Good identification**					**365770**
Staphylococcus kloosii		100.0	0.41	VPT 10		
365771	**Low discrimination**					**365771**
Staphylococcus xylosus		64.38	0.32	VPT 14		
Staphylococcus equorum		30.88	0.27	NIT 89		
Staphylococcus gallinarum		4.74	0.07			
365772	**Low discrimination**					**365772**
Staphylococcus kloosii		77.53	0.42	VPT 10		
Staphylococcus arlettae		14.63	0.15	SUC 9		
Staphylococcus xylosus		7.84	0.00			
365773	**Low discrimination**					**365773**
Staphylococcus xylosus		65.38	0.41	VPT 14		
Staphylococcus equorum		28.33	0.35	NIT 89		
Staphylococcus gallinarum		4.14	0.15			
366053	**Good identification**					**366053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	TRE 99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
366153	**Acceptable identification**					**366153**
Staphylococcus cohnii subsp. urealyticum		94.84	0.31	TRE 99		
Staphylococcus xylosus		5.16	-.04			
366243	**Good identification**					**366243**
Staphylococcus epidermidis		100.0	0.30	GLR 1		
366340	**Good identification**					**366340**
Staphylococcus epidermidis		100.0	0.29	GLR 1		
366341	**Good identification**					**366341**
Staphylococcus epidermidis		100.0	0.35	GLR 1		
366342	**Good identification**					**366342**
Staphylococcus epidermidis		100.0	0.40	GLR 1		
366343	**Good identification**					**366343**
					NOV	
Staphylococcus epidermidis		99.22	0.45	GLR 1	1	
Staphylococcus xylosus		0.78	-.02		99	
366371	**Good identification**					**366371**
Staphylococcus xylosus		100.0	0.35	VPT 14		
366373	**Good identification**					**366373**
Staphylococcus xylosus		100.0	0.44	VPT 14		
366403	**Good identification**					**366403**
					NOV	
Staphylococcus cohnii subsp. urealyticum		98.72	0.44	MAN 91	99	
Staphylococcus simulans		1.28	-.01		1	
366411	**Good identification**					**366411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.44	LAC 91		
366413	**Very good identification**					**366413**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.49	0.64		99	
Staphylococcus simulans		0.51	0.12		1	
366441	**Good identification**					**366441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	MAN 91		
366443	**Very good identification**					**366443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.59	MAN 91		
366450	**Acceptable identification**					**366450**
Staphylococcus saprophyticus subsp. bovis		90.97	0.38			
Staphylococcus cohnii subsp. urealyticum		5.19	0.18			
Staphylococcus kloosii		3.84	0.27			
366451	**Very good identification**					**366451**
Staphylococcus cohnii subsp. urealyticum		95.76	0.59	LAC 91		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus saprophyticus subsp. bovis		4.24	0.26			
366452	**Low discrimination**					**366452**
Staphylococcus cohnii subsp. urealyticum		66.10	0.39	MNS 99		
Staphylococcus saprophyticus subsp. bovis		28.66	0.26			
Staphylococcus kloosii		5.25	0.28			
366453	**Very good identification**					**366453**
Staphylococcus cohnii subsp. urealyticum		99.79	0.80			
Staphylococcus saprophyticus subsp. bovis		0.11	0.13			
Staphylococcus xylosus		0.06	0.04			
366473	**Low discrimination**					**366473**
Staphylococcus cohnii subsp. urealyticum		61.13	0.39	XYL 1		
Staphylococcus xylosus		38.87	0.26			
366503	**Good identification**					**366503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	MAN 91		
366511	**Good identification**					**366511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	LAC 91		
366513	**Very good identification**					**366513**
Staphylococcus cohnii subsp. urealyticum		99.44	0.57			
Staphylococcus xylosus		0.56	0.01			
366541	**Good identification**					**366541**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	MAN 91		
366543	**Good identification**					**366543**
Staphylococcus cohnii subsp. urealyticum		99.16	0.52	MAN 91		
Staphylococcus xylosus		0.84	0.00			
366550	**Good identification**					**366550**
Staphylococcus saprophyticus subsp. bovis		97.41	0.50			
Staphylococcus kloosii		2.00	0.33			
Staphylococcus cohnii subsp. urealyticum		0.60	0.10			
366551	**Low discrimination**					**366551**
Staphylococcus cohnii subsp. urealyticum		68.49	0.52	LAC 91		
Staphylococcus saprophyticus subsp. bovis		28.30	0.38			
Staphylococcus xylosus		3.21	0.15			
366552	**Low discrimination**					**366552**
Staphylococcus saprophyticus subsp. bovis		74.85	0.38			
Staphylococcus cohnii subsp. urealyticum		18.50	0.31			
Staphylococcus kloosii		6.65	0.34			
366553	**Very good identification**					**366553**
Staphylococcus cohnii subsp. urealyticum		97.71	0.72			
Staphylococcus xylosus		1.29	0.24			
Staphylococcus saprophyticus subsp. bovis		1.00	0.26			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
366571	**Good identification**						**366571**
Staphylococcus xylosus		98.16	0.37	VPT 14			
Staphylococcus cohnii subsp. urealyticum		1.84	0.10	SUC 95			
366573	**Low discrimination**						**366573**
					COA	NOV	
Staphylococcus xylosus		88.01	0.46	VPT 14	1	99	
Staphylococcus lutrae		6.12	0.18	SUC 95	99	1	
Staphylococcus cohnii subsp. urealyticum		5.87	0.31		1	99	
366603	**Acceptable identification**						**366603**
					NOV		
Staphylococcus simulans		90.96	0.28	VPT 4	1		
Staphylococcus cohnii subsp. urealyticum		9.04	0.13	ARG 92	99		
366612	**Good identification**						**366612**
Staphylococcus simulans		100.0	0.29	VPT 4			
366613	**Low discrimination**						**366613**
					NOV		
Staphylococcus simulans		77.64	0.40	VPT 4	1		
Staphylococcus cohnii subsp. urealyticum		19.50	0.33	ARG 92	99		
Staphylococcus xylosus		2.85	0.07		99		
366633	**Good identification**						**366633**
					NOV		
Staphylococcus xylosus		97.67	0.29	VPT 14	99		
Staphylococcus simulans		2.33	-.01	GAL 91	1		
366640	**Low discrimination**						**366640**
					NOV	PYR	
Staphylococcus warneri		85.16	0.35	PHS 1	1	8	
Staphylococcus saprophyticus subsp. bovis		14.85	0.09		99	99	
366643	**Low discrimination**						**366643**
					NOV		
Staphylococcus cohnii subsp. urealyticum		49.52	0.28	SUC 3	99		
Staphylococcus simulans		39.51	0.20	MAN 91	1		
Staphylococcus xylosus		10.97	0.05		99		
366650	**Acceptable identification**						**366650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		94.26	0.50		99	99	
Staphylococcus warneri		3.96	0.32		1	8	
Staphylococcus pasteuri		1.78	0.10		1	1	
366651	**Low discrimination**						**366651**
Staphylococcus saprophyticus subsp. bovis		72.07	0.38				
Staphylococcus xylosus		15.34	0.21				
Staphylococcus cohnii subsp. urealyticum		12.59	0.28				
366652	**Low discrimination**						**366652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		78.67	0.38		99	99	
Staphylococcus simulans		13.10	0.22		1	70	
Staphylococcus warneri		4.40	0.22		1	8	
366653	**Low discrimination**						**366653**
					NOV		
Staphylococcus cohnii subsp. urealyticum		55.55	0.49	SUC 3	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		19.06	0.30		99		
Staphylococcus simulans		17.53	0.33		1		
366663	**Good identification**						**366663**
Staphylococcus xylosus		100.0	0.27	VPT 14			
366671	**Good identification**						**366671**
Staphylococcus xylosus		100.0	0.42	VPT 14			
366673	**Good identification**						**366673**
Staphylococcus xylosus		99.75	0.52	VPT 14			
Staphylococcus cohnii subsp. urealyticum		0.26	0.08	GAL 91			
366713	**Low discrimination**						**366713**
					COA	NOV	
Staphylococcus xylosus		46.72	0.28	VPT 14	1	99	
Staphylococcus intermedius		26.32	0.24	XYL 92	99	1	
Staphylococcus cohnii subsp. urealyticum		13.53	0.26	MLT 93	1	99	
366731	**Good identification**						**366731**
Staphylococcus xylosus		100.0	0.40	VPT 14			
366733	**Good identification**						**366733**
Staphylococcus xylosus		100.0	0.49	VPT 14			
366750	**Very good identification**						**366750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		99.54	0.63		99	99	
Staphylococcus warneri		0.33	0.21		1	8	
Staphylococcus xylosus		0.14	0.00		99	65	
366751	**Low discrimination**						**366751**
Staphylococcus saprophyticus subsp. bovis		63.59	0.50				
Staphylococcus xylosus		34.21	0.41				
Staphylococcus cohnii subsp. urealyticum		1.19	0.20				
366752	**Good identification**						**366752**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		97.70	0.50		99	99	
Staphylococcus xylosus		1.51	0.10		99	65	
Staphylococcus warneri		0.43	0.12		1	8	
366753	**Low discrimination**						**366753**
Staphylococcus xylosus		73.03	0.51	VPT 14			
Staphylococcus saprophyticus subsp. bovis		11.92	0.38	XYL 92			
Staphylococcus cohnii subsp. urealyticum		9.02	0.41				
366761	**Good identification**						**366761**
Staphylococcus xylosus		100.0	0.39	VPT 14			
366763	**Good identification**						**366763**
					COA	NOV	
Staphylococcus xylosus		96.84	0.48	VPT 14	1	99	
Staphylococcus lutrae		2.86	0.12	MAN 94	99	1	
Staphylococcus gallinarum		0.29	-.05		1	99	
366771	**Very good identification**						**366771**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		99.44	0.63	VPT 14			
Staphylococcus gallinarum		0.23	0.07				
Staphylococcus equorum		0.17	0.08				
366772	**Acceptable identification**						**366772**
Staphylococcus xylosus		94.63	0.31	VPT 14			
Staphylococcus saprophyticus subsp. bovis		5.37	0.09	MNS 99			
366773	**Very good identification**						**366773**
					COA	NOV	
Staphylococcus xylosus		99.29	0.73	VPT 14	1	99	
Staphylococcus lutrae		0.36	0.18		99	1	
Staphylococcus gallinarum		0.19	0.15		1	99	
367053	**Good identification**						**367053**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	TRE 99			
367373	**Low discrimination**						**367373**
Staphylococcus xylosus		74.78	0.32	VPT 14			
Staphylococcus gallinarum		14.06	0.15	TRE 96			
Staphylococcus equorum		11.17	0.16				
367403	**Good identification**						**367403**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	MAN 91			
367411	**Good identification**						**367411**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	LAC 91			
367413	**Very good identification**						**367413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.56				
367441	**Good identification**						**367441**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	MAN 91			
367443	**Good identification**						**367443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	MAN 91			
367450	**Low discrimination**						**367450**
Staphylococcus kloosii		65.57	0.27	VPT 10			
Staphylococcus cohnii subsp. urealyticum		34.43	0.10	NIT 1			
367451	**Good identification**						**367451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	LAC 91			
367452	**Low discrimination**						**367452**
Staphylococcus cohnii subsp. urealyticum		83.05	0.30	MNS 99			
Staphylococcus kloosii		16.95	0.28				
367453	**Very good identification**						**367453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.71				
367473	**Low discrimination**						**367473**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		70.98	0.30	XYL 1		
Staphylococcus xylosus		29.02	0.13			
367503	**Good identification**					**367503**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	MAN 91		
367511	**Good identification**					**367511**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	LAC 91		
367513	**Good identification**					**367513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.48			
367543	**Good identification**					**367543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	MAN 91		
367550	**Low discrimination**					**367550**
Staphylococcus kloosii		66.99	0.33	VPT 10		
Staphylococcus saprophyticus subsp. bovis		33.01	0.09	NIT 1		
367551	**Good identification**					**367551**
Staphylococcus cohnii subsp. urealyticum		97.08	0.43	LAC 91		
Staphylococcus xylosus		2.92	0.03			
367553	**Very good identification**					**367553**
Staphylococcus cohnii subsp. urealyticum		99.16	0.64			
Staphylococcus xylosus		0.84	0.12			
367573	**Low discrimination**					**367573**
Staphylococcus xylosus		72.95	0.34	VPT 14		
Staphylococcus gallinarum		10.86	0.15	SUC 95		
Staphylococcus equorum		8.62	0.16			
367653	**Low discrimination**					**367653**
Staphylococcus cohnii subsp. urealyticum		81.93	0.40	SUC 3		
Staphylococcus xylosus		18.07	0.18			
367671	**Low discrimination**					**367671**
Staphylococcus xylosus		56.38	0.30	VPT 14		
Staphylococcus equorum		38.43	0.28	GAL 91		
Staphylococcus gallinarum		5.19	0.07			
367673	**Low discrimination**					**367673**
Staphylococcus xylosus		58.99	0.39	VPT 14		
Staphylococcus equorum		36.34	0.36	GAL 91		
Staphylococcus gallinarum		4.67	0.15			
367731	**Low discrimination**					**367731**
Staphylococcus xylosus		49.58	0.28	VPT 14		
Staphylococcus equorum		44.42	0.28	MLT 93		
Staphylococcus gallinarum		6.00	0.07			
367733	**Low discrimination**					**367733**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		52.26	0.37	VPT 14		
Staphylococcus equorum		42.30	0.36	MLT 93		
Staphylococcus gallinarum		5.44	0.15			
367751	**Low discrimination**					**367751**
Staphylococcus xylosus		76.16	0.29	VPT 14		
Staphylococcus gallinarum		7.98	0.07	XYL 92		
Staphylococcus equorum		6.03	0.08			
367753	**Low discrimination**					**367753**
Staphylococcus xylosus		73.87	0.38	VPT 14		
Staphylococcus cohnii subsp. urealyticum		14.19	0.33	XYL 92		
Staphylococcus gallinarum		6.66	0.15			
367761	**Low discrimination**					**367761**
Staphylococcus gallinarum		55.77	0.28	VPT 1		
Staphylococcus xylosus		39.92	0.26	MAN 91		
Staphylococcus equorum		4.31	0.08			
367763	**Low discrimination**					**367763**
Staphylococcus gallinarum		52.27	0.36	VPT 1		
Staphylococcus xylosus		43.49	0.36	MAN 91		
Staphylococcus equorum		4.24	0.16			
367771	**Low discrimination**					**367771**
Staphylococcus xylosus		38.71	0.51	VPT 14		
Staphylococcus gallinarum		34.90	0.48			
Staphylococcus equorum		26.39	0.49			
367773	**Low discrimination**					**367773**
Staphylococcus xylosus		41.82	0.60	VPT 14		
Staphylococcus gallinarum		32.43	0.56			
Staphylococcus equorum		25.76	0.57			
370413	**Good identification**					**370413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1		
370443	**Good identification**					**370443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
370451	**Good identification**					**370451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
370453	**Good identification**					**370453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	ORN 1		
370513	**Good identification**					**370513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ORN 1		
370553	**Good identification**					**370553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ORN 1		
370640	**Good identification**					**370640**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		100.0	0.42	ORN 1			
370642	**Low discrimination**						**370642**
					NOV		
Staphylococcus warneri		85.43	0.32	ORN 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		14.57	-.04		99		
370650	**Low discrimination**						**370650**
					NOV		
Staphylococcus warneri		70.85	0.39	ORN 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		23.56	0.09		99		
Staphylococcus pasteuri		5.59	0.02		1		
370740	**Good identification**						**370740**
Staphylococcus warneri		100.0	0.32	ORN 1			
370750	**Good identification**						**370750**
Staphylococcus warneri		100.0	0.29	ORN 1			
371413	**Good identification**						**371413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1			
371453	**Good identification**						**371453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	ORN 1			
371553	**Good identification**						**371553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1			
372453	**Good identification**						**372453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ORN 1			
372553	**Good identification**						**372553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ORN 1			
372640	**Good identification**						**372640**
Staphylococcus warneri		100.0	0.35	ORN 1			
372641	**Good identification**						**372641**
Staphylococcus lugdunensis		100.0	0.33	GLR 2			
372643	**Good identification**						**372643**
Staphylococcus lugdunensis		100.0	0.53	GLR 2			
372650	**Low discrimination**						**372650**
					NOV	PYR	
Staphylococcus warneri		41.43	0.32	ORN 1	1	8	
Staphylococcus saprophyticus subsp. bovis		39.89	0.22		99	99	
Staphylococcus pasteuri		18.68	0.10		1	1	
372741	**Good identification**						**372741**
Staphylococcus lugdunensis		100.0	0.33	GLR 2			
372743	**Good identification**						**372743**
					NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lugdunensis		99.83	0.53	GLR 2	1		
Staphylococcus xylosus		0.17	-.04		99		
372750	**Acceptable identification**						**372750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		92.42	0.34	ORN 1	99	99	
Staphylococcus warneri		7.58	0.21		1	8	
372771	**Good identification**						**372771**
Staphylococcus xylosus		100.0	0.33	VPT 14			
372773	**Good identification**						**372773**
Staphylococcus xylosus		100.0	0.43	VPT 14			
373453	**Good identification**						**373453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ORN 1			
373773	**Low discrimination**						**373773**
Staphylococcus xylosus		89.62	0.30	VPT 14			
Staphylococcus equorum		10.38	0.12	ORN 1			
374413	**Good identification**						**374413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ORN 1			
374443	**Good identification**						**374443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	ORN 1			
374450	**Good identification**						**374450**
Staphylococcus kloosii		100.0	0.27	VPT 10			
374451	**Good identification**						**374451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	ORN 1			
374452	**Low discrimination**						**374452**
Staphylococcus kloosii		58.10	0.28	VPT 10			
Staphylococcus cohnii subsp. urealyticum		41.90	0.13	ORN 1			
374453	**Very good identification**						**374453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.54	ORN 1			
374513	**Good identification**						**374513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1			
374550	**Good identification**						**374550**
Staphylococcus kloosii		100.0	0.33	VPT 10			
374552	**Low discrimination**						**374552**
Staphylococcus kloosii		86.26	0.34	VPT 10			
Staphylococcus cohnii subsp. urealyticum		13.74	0.06	ORN 1			
374553	**Good identification**						**374553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
375413	**Good identification**					**375413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
375450	**Good identification**					**375450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
375452	**Low discrimination**					**375452**
Staphylococcus kloosii		78.10	0.28	VPT 10		
Staphylococcus cohnii subsp. urealyticum		21.91	0.05	ORN 1		
375453	**Good identification**					**375453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.46	ORN 1		
375550	**Good identification**					**375550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
375552	**Good identification**					**375552**
Staphylococcus kloosii		100.0	0.34	VPT 10		
375553	**Good identification**					**375553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	ORN 1		
376343	**Good identification**					**376343**
Staphylococcus epidermidis		100.0	0.27	ORN 12		
376453	**Good identification**					**376453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ORN 1		
376553	**Good identification**					**376553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ORN 1		
376773	**Good identification**					**376773**
Staphylococcus xylosus		100.0	0.31	VPT 14		
377453	**Good identification**					**377453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ORN 1		
400000	**Low discrimination**					**400000**
					OXI	
Staphylococcus carnosus subsp. utilis		71.94	0.70		1	
Micrococcus sp.		18.97	0.59		89	
Dermacoccus nishinomiyaensis		6.74	0.59		93	
400001	**Low discrimination**					**400001**
					OXI	
Staphylococcus capitis subsp. capitis		58.42	0.49	NIT 89	1	
Staphylococcus carnosus subsp. utilis		21.74	0.29	MAN 93	1	
Micrococcus sp.		11.58	0.24		89	
400002	**Low discrimination**					**400002**
					OXI	
Staphylococcus carnosus subsp. utilis		62.06	0.29	LAC 1	1	
Micrococcus sp.		16.37	0.18		89	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		12.67	0.31		1			
400010	**Low discrimination**							**400010**
					OXI	NOV		
Macrococcus bovicus		46.72	0.48	ARG 1	99	99		
Staphylococcus carnosus subsp. utilis		26.12	0.29		1	1		
Staphylococcus capitis subsp. capitis		9.42	0.31		1	1		
400011	**Very good identification**							**400011**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		99.88	0.72	NIT 89	1	1	1	
Macrococcus bovicus		0.05	0.07		99	99	99	
Staphylococcus cohnii subsp. cohnii		0.03	0.08		1	99	12	
400013	**Good identification**							**400013**
Staphylococcus capitis subsp. capitis		100.0	0.37	NIT 89				
400020	**Low discrimination**							**400020**
					OXI			
Staphylococcus carnosus subsp. utilis		51.78	0.29	XYL 1	1			
Kocuria rosea		26.32	0.32		35			
Micrococcus sp.		13.66	0.18		89			
400031	**Good identification**							**400031**
Staphylococcus capitis subsp. capitis		100.0	0.31	NIT 89				
400040	**Low discrimination**							**400040**
					OXI			
Staphylococcus auricularis		55.77	0.59	NIT 90	1			
Micrococcus sp.		25.56	0.36	TRE 92	89			
Staphylococcus carnosus subsp. utilis		13.01	0.29		1			
400041	**Low discrimination**							**400041**
					OXI			
Staphylococcus auricularis		82.43	0.39	NIT 90	1			
Micrococcus sp.		6.94	0.01	TRE 92	89			
Staphylococcus capitis subsp. capitis		4.70	0.08	MNS 10	1			
400042	**Acceptable identification**							**400042**
					OXI			
Staphylococcus auricularis		92.57	0.39	NIT 90	1			
Micrococcus sp.		3.86	-.05	TRE 92	89			
Staphylococcus carnosus subsp. utilis		1.96	-.12	LAC 10	1			
400050	**Low discrimination**							**400050**
					OXI	NOV		
Macrococcus bovicus		89.15	0.55	ARG 1	99	99		
Macrococcus equipersicus		8.68	0.36		99	99		
Staphylococcus auricularis		0.93	0.17		1	1		
400051	**Low discrimination**							**400051**
					OXI	NOV	YEP	
Staphylococcus capitis subsp. capitis		83.16	0.31	NIT 89	1	1	1	
Macrococcus bovicus		9.72	0.14	MLT 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		7.12	0.18		1	99	12	
400100	**Low discrimination**							**400100**
					OXI			
Dermacoccus nishinomiyaensis		51.01	0.44	ARG 1	93			
Staphylococcus carnosus subsp. utilis		28.88	0.29		1			
Micrococcus sp.		7.62	0.18		89			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
400110	**Low discrimination**						**400110**
					YEP		
Macrococcus bovicus		89.25	0.40	ARG 1	99		
Macrococcus equipersicus		7.50	0.20		99		
Dermacoccus nishinomiyaensis		2.08	0.03		1		
400111	**Good identification**						**400111**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		97.90	0.31	NIT 89	1	1	
Macrococcus bovicus		2.10	-.01	GAL 1	99	99	
400140	**Low discrimination**						**400140**
					OXI		
Staphylococcus auricularis		80.01	0.39	NIT 90	1		
Kocuria varians		7.39	0.19	GAL 10	13		
Micrococcus sp.		3.33	-.05	TRE 92	89		
400150	**Acceptable identification**						**400150**
Macrococcus bovicus		91.13	0.48	ARG 1			
Macrococcus equipersicus		8.88	0.28				
400200	**Low discrimination**						**400200**
					OXI		
Micrococcus sp.		40.82	0.38	ARG 1	89		
Staphylococcus auricularis		26.82	0.50	SUC 9	1		
Staphylococcus carnosus subsp. utilis		15.81	0.29		1		
400201	**Very good identification**						**400201**
					OXI		
Staphylococcus capitis subsp. capitis		95.08	0.55	NIT 89	1		
Staphylococcus auricularis		3.59	0.31	MAN 93	1		
Micrococcus sp.		1.00	0.03		89		
400202	**Low discrimination**						**400202**
					OXI		
Staphylococcus auricularis		83.90	0.31	NIT 90	1		
Micrococcus sp.		11.61	-.03	TRE 92	89		
Staphylococcus carnosus subsp. utilis		4.50	-.12	LAC 10	1		
400210	**Low discrimination**						**400210**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		48.10	0.37	NIT 89	1	1	
Macrococcus bovicus		32.12	0.35	MNS 99	99	99	
Staphylococcus vitulinus		12.02	0.21		99	99	
400211	**Very good identification**						**400211**
Staphylococcus capitis subsp. capitis		100.0	0.78	NIT 89			
400213	**Good identification**						**400213**
Staphylococcus capitis subsp. capitis		99.31	0.43	NIT 89			
Staphylococcus capitis subsp. ureolyticus		0.69	-.11	LAC 2			
400230	**Good identification**						**400230**
					NOV	OXI	
Staphylococcus vitulinus		97.20	0.29	ARG 1	99	99	
Staphylococcus capitis subsp. capitis		1.45	-.05	NIT 91	1	1	
Kocuria rosea		1.35	-.08		1	35	
400231	**Good identification**						**400231**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus capitis subsp. capitis		100.0	0.37	NIT 89			
400240	**Low discrimination**						**400240**
					COA	OXI	
Staphylococcus aureus subsp. anaerobius		82.43	0.59	PHS 99	99	1	
Staphylococcus auricularis		15.34	0.58		1	1	
Micrococcus sp.		0.75	0.15		1	89	
400241	**Low discrimination**						**400241**
					OXI	COA	
Staphylococcus auricularis		57.86	0.38	NIT 90	1	1	
Staphylococcus aureus subsp. anaerobius		28.26	0.18	TRE 92	1	99	
Kocuria kristinae		7.25	0.13	MNS 10	99	1	
400242	**Low discrimination**						**400242**
					COA	PYR	
Staphylococcus auricularis		61.92	0.38	NIT 90	1	40	
Staphylococcus aureus subsp. anaerobius		30.25	0.18	TRE 92	99	1	
Staphylococcus haemolyticus		3.43	0.13	LAC 10	1	98	
400250	**Low discrimination**						**400250**
					OXI	COA	
Macrococcus bovicus		74.54	0.43	ARG 1	99	1	
Staphylococcus aureus subsp. anaerobius		15.35	0.18		1	99	
Macrococcus equipersicus		2.87	0.15		99	1	
400251	**Acceptable identification**						**400251**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		94.20	0.37	NIT 89	1	1	
Staphylococcus capitis subsp. ureolyticus		4.32	0.00	MLT 1	1	1	
Macrococcus bovicus		1.48	0.02		99	99	
400300	**Low discrimination**						**400300**
					OXI		
Staphylococcus auricularis		49.39	0.31	NIT 90	1		
Dermacoccus nishinomiyaensis		29.54	0.20	GAL 10	93		
Kocuria varians		7.17	0.15	TRE 92	13		
400310	**Low discrimination**						**400310**
					OXI	NOV	
Macrococcus bovicus		88.90	0.28	ARG 1	99	99	
Macrococcus carouselicus		5.00	-.02		99	99	
Staphylococcus capitis subsp. capitis		3.14	-.05		1	1	
400311	**Good identification**						**400311**
Staphylococcus capitis subsp. capitis		100.0	0.37	NIT 89			
400340	**Low discrimination**						**400340**
					COA	PYR	
Staphylococcus auricularis		61.40	0.38	NIT 90	1	40	
Staphylococcus aureus subsp. anaerobius		30.00	0.18	GAL 10	99	1	
Staphylococcus haemolyticus		6.82	0.20	TRE 92	1	98	
400350	**Low discrimination**						**400350**
					OXI	PYR	
Macrococcus bovicus		89.02	0.35	ARG 1	99	1	
Staphylococcus haemolyticus		5.71	0.15		1	98	
Macrococcus equipersicus		3.43	0.08		99	1	
400400	**Very good identification**						**400400**
					NOV		
Staphylococcus carnosus subsp. utilis		95.84	0.85		1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		3.90	0.73		1			
Staphylococcus cohnii subsp. cohnii		0.13	0.46		99			
400401	**Low discrimination**							**400401**
					NOV			
Staphylococcus carnosus subsp. utilis		62.42	0.44	MNS 1	1			
Staphylococcus auricularis		27.92	0.53		1			
Staphylococcus cohnii subsp. cohnii		9.01	0.47		99			
400402	**Low discrimination**							**400402**
Staphylococcus carnosus subsp. utilis		68.96	0.44	LAC 1				
Staphylococcus auricularis		30.84	0.53					
Staphylococcus piscifermentans		0.11	0.12					
400403	**Low discrimination**							**400403**
					NOV			
Staphylococcus auricularis		79.29	0.34	NIT 90	1			
Staphylococcus carnosus subsp. utilis		16.12	0.03	MNS 10	1			
Staphylococcus cohnii subsp. cohnii		2.33	0.05	LAC 10	99			
400410	**Low discrimination**							**400410**
					OXI	NOV	YEP	
Staphylococcus carnosus subsp. utilis		55.92	0.44	MAN 1	1	1	10	
Macrococcus bovicus		28.58	0.51		99	99	99	
Staphylococcus cohnii subsp. cohnii		9.49	0.48		1	99	12	
400411	**Low discrimination**							**400411**
					NOV			
Staphylococcus cohnii subsp. cohnii		53.68	0.49	ARG 1	99			
Staphylococcus capitis subsp. capitis		40.54	0.37		1			
Staphylococcus carnosus subsp. utilis		2.95	0.03		1			
400413	**Low discrimination**							**400413**
					NOV	PYR		
Aerococcus viridans		47.68	0.27	ARG 1	1	95		
Staphylococcus capitis subsp. capitis		31.61	0.03	ESL 90	1	1		
Staphylococcus cohnii subsp. cohnii		20.71	0.08		99	1		
400420	**Good identification**							**400420**
					NOV			
Staphylococcus carnosus subsp. utilis		95.97	0.44	XYL 1	1			
Staphylococcus auricularis		3.90	0.32		1			
Staphylococcus cohnii subsp. cohnii		0.13	0.05		99			
400433	**Good identification**							**400433**
Aerococcus viridans		100.0	0.27	ARG 1				
400440	**Low discrimination**							**400440**
					NOV			
Staphylococcus auricularis		86.21	0.80	NIT 90	1			
Staphylococcus carnosus subsp. utilis		9.18	0.44		1			
Staphylococcus cohnii subsp. cohnii		3.87	0.56		99			
400441	**Low discrimination**							**400441**
					NOV			
Staphylococcus auricularis		68.98	0.61	NIT 90	1			
Staphylococcus cohnii subsp. cohnii		30.22	0.57	MNS 10	99			
Staphylococcus carnosus subsp. utilis		0.67	0.03		1			
400442	**Very good identification**							**400442**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		97.72	0.61	NIT 90	1			
Staphylococcus carnosus subsp. utilis		0.95	0.03	LAC 10	1			
Staphylococcus cohnii subsp. cohnii		0.40	0.15		99			
400443	**Acceptable identification**							**400443**
					NOV	PYR		
Staphylococcus auricularis		91.83	0.41	NIT 90	1	40		
Aerococcus viridans		4.51	0.30	MNS 10	1	95		
Staphylococcus cohnii subsp. cohnii		3.66	0.16	LAC 10	99	1		
400450	**Low discrimination**							**400450**
					OXI	NOV	YEP	
Macrococcus bovicus		62.19	0.59	ARG 1	99	99	99	
Staphylococcus cohnii subsp. cohnii		28.01	0.58		1	99	12	
Staphylococcus auricularis		4.95	0.39		1	1	1	
400451	**Very good identification**							**400451**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		95.74	0.59	ARG 1	1	99	12	
Macrococcus bovicus		1.98	0.18		99	99	99	
Staphylococcus auricularis		1.74	0.20		1	1	1	
400453	**Low discrimination**							**400453**
					NOV	PYR		
Aerococcus viridans		74.41	0.39	ARG 1	1	95		
Staphylococcus cohnii subsp. cohnii		25.59	0.18	ESL 90	99	1		
400460	**Low discrimination**							**400460**
					NOV			
Staphylococcus auricularis		86.85	0.39	NIT 90	1			
Staphylococcus carnosus subsp. utilis		9.25	0.03	XYL 1	1			
Staphylococcus cohnii subsp. cohnii		3.90	0.15		99			
400463	**Good identification**							**400463**
Aerococcus viridans		100.0	0.30	ARG 1				
400472	**Good identification**							**400472**
Aerococcus viridans		100.0	0.32	ARG 1				
400473	**Good identification**							**400473**
Aerococcus viridans		100.0	0.39	ARG 1				
400500	**Low discrimination**							**400500**
					OXI	NOV		
Staphylococcus carnosus subsp. utilis		66.80	0.44	GAL 1	1	1		
Staphylococcus auricularis		29.88	0.53		1	1		
Macrococcus carouselicus		1.66	0.20		99	99		
400501	**Low discrimination**							**400501**
					NOV			
Staphylococcus auricularis		65.53	0.34	NIT 90	1			
Staphylococcus cohnii subsp. cohnii		21.15	0.27	GAL 10	99			
Staphylococcus carnosus subsp. utilis		13.32	0.03	MNS 10	1			
400502	**Low discrimination**							**400502**
Staphylococcus auricularis		79.92	0.34	NIT 90				
Staphylococcus carnosus subsp. utilis		16.24	0.03	GAL 10				
Staphylococcus piscifermentans		3.84	0.16	LAC 10				
400510	**Low discrimination**							**400510**

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
					OXI	YEP			
Macrococcus bovicus		78.05	0.44	ARG 1	99	99			
Staphylococcus cohnii subsp. cohnii		6.72	0.28		1	12			
Macrococcus carouselicus		5.65	0.16		99	67			
400511	**Low discrimination**								**400511**
					OXI	NOV	YEP		
Staphylococcus cohnii subsp. cohnii		84.96	0.29	ARG 1	1	99	12		
Macrococcus bovicus		9.21	0.03	GAL 10	99	99	99		
Staphylococcus capitis subsp. capitis		5.83	-.04		1	1	1		
400513	**Good identification**								**400513**
Aerococcus viridans		100.0	0.27	ARG 1					
400533	**Good identification**								**400533**
Aerococcus viridans		100.0	0.27	ARG 1					
400540	**Good identification**								**400540**
					NOV				
Staphylococcus auricularis		93.78	0.61	NIT 90	1				
Staphylococcus cohnii subsp. cohnii		4.21	0.36	GAL 10	99				
Staphylococcus carnosus subsp. utilis		0.91	0.03		1				
400541	**Low discrimination**								**400541**
					NOV				
Staphylococcus auricularis		69.54	0.41	NIT 90	1				
Staphylococcus cohnii subsp. cohnii		30.46	0.37	GAL 10	99				
400542	**Good identification**								**400542**
Staphylococcus auricularis		96.00	0.41	NIT 90					
Aerococcus viridans		2.02	0.22	GAL 10					
Staphylococcus piscifermentans		1.98	0.16	LAC 10					
400550	**Low discrimination**								**400550**
					OXI	YEP			
Macrococcus bovicus		82.76	0.51	ARG 1	99	99			
Staphylococcus cohnii subsp. cohnii		9.67	0.39		1	12			
Macrococcus equipersicus		5.37	0.28		99	99			
400551	**Acceptable identification**								**400551**
					OXI	NOV	PYR	YEP	
Staphylococcus cohnii subsp. cohnii		90.15	0.39	ARG 1	1	99	1	12	
Macrococcus bovicus		7.20	0.10	GAL 10	99	99	1	99	
Aerococcus viridans		2.65	0.20		1	1	95	1	
400552	**Low discrimination**								**400552**
					OXI	NOV	YEP		
Aerococcus viridans		52.80	0.32	ARG 1	1	1	1		
Macrococcus bovicus		37.21	0.10	ESL 90	99	99	99		
Staphylococcus kloosii		9.99	0.18		1	99	99		
400553	**Good identification**								**400553**
Aerococcus viridans		100.0	0.39	ARG 1					
400563	**Good identification**								**400563**
Aerococcus viridans		100.0	0.30	ARG 1					
400572	**Good identification**								**400572**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.32	ARG 1			
400573	**Good identification**						**400573**
Aerococcus viridans		100.0	0.39	ARG 1			
400600	**Low discrimination**						**400600**
					OXI	NOV	
Staphylococcus auricularis		75.78	0.72	NIT 90	1	1	
Staphylococcus carnosus subsp. utilis		20.40	0.44		1	1	
Macrococcus carouselicus		2.40	0.34		99	99	
400601	**Low discrimination**						**400601**
					OXI	PYR	
Staphylococcus auricularis		88.60	0.52	NIT 90	1	40	
Kocuria kristinae		4.76	0.20	MNS 10	99	93	
Staphylococcus capitis subsp. capitis		4.16	0.20		1	1	
400602	**Good identification**						**400602**
Staphylococcus auricularis		95.73	0.52	NIT 90			
Staphylococcus carnosus subsp. utilis		2.34	0.03	LAC 10			
Staphylococcus petrasii subsp. jettensis		0.78	-.05				
400603	**Low discrimination**						**400603**
					OXI	PYR	
Staphylococcus auricularis		79.94	0.33	NIT 90	1	40	
Kocuria kristinae		9.65	0.07	MNS 10	99	93	
Staphylococcus hyicus		5.22	0.00	LAC 10	1	1	
400610	**Low discrimination**						**400610**
					OXI	NOV	
Macrococcus bovicus		45.44	0.39	ARG 1	99	99	
Macrococcus carouselicus		26.72	0.30		99	99	
Staphylococcus auricularis		13.35	0.31		1	1	
400611	**Good identification**						**400611**
					NOV		
Staphylococcus capitis subsp. capitis		97.24	0.43	NIT 89	1		
Staphylococcus auricularis		1.57	0.11	TRE 2	1		
Staphylococcus cohnii subsp. cohnii		0.70	0.08		99		
400620	**Low discrimination**						**400620**
Staphylococcus auricularis		78.80	0.31	NIT 90			
Staphylococcus carnosus subsp. utilis		21.21	0.03	XYL 1			
400633	**Good identification**						**400633**
Aerococcus viridans		100.0	0.27	ARG 1			
400640	**Low discrimination**						**400640**
Staphylococcus auricularis		60.92	0.80	NIT 90			
Staphylococcus petrasii subsp. pragensis		23.52	0.54				
Staphylococcus petrasii subsp. jettensis		13.33	0.51				
400641	**Low discrimination**						**400641**
					OXI		
Staphylococcus auricularis		78.40	0.60	NIT 90	1		
Kocuria kristinae		16.23	0.39	MNS 10	99		
Staphylococcus petrasii subsp. pragensis		2.75	0.13		1		
400642	**Low discrimination**						**400642**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		66.57	0.60	NIT 90			
Staphylococcus petrasii subsp. jettensis		23.14	0.36	LAC 10			
Staphylococcus haemolyticus		4.26	0.37				
400643	**Low discrimination**						**400643**
					OXI		
Staphylococcus auricularis		63.76	0.40	NIT 90	1		
Kocuria kristinae		29.70	0.27	MNS 10	99		
Aerococcus viridans		3.39	0.30	LAC 10	1		
400650	**Low discrimination**						**400650**
					OXI	NOV	
Macrococcus bovicus		47.80	0.47	ARG 1	99	99	
Staphylococcus warneri		16.95	0.49		1	1	
Staphylococcus auricularis		14.04	0.39		1	1	
400652	**Low discrimination**						**400652**
					NOV	PYR	
Staphylococcus haemolyticus		34.48	0.32	VPT 86	2	98	
Staphylococcus warneri		33.54	0.39	NIT 99	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		9.67	0.07		99	7	
400653	**Low discrimination**						**400653**
					OXI		
Aerococcus viridans		77.82	0.39	ARG 1	1		
Staphylococcus simulans		12.72	0.06	ESL 90	1		
Kocuria kristinae		9.46	-.02		99		
400660	**Low discrimination**						**400660**
Staphylococcus auricularis		61.66	0.39	NIT 90			
Staphylococcus petrasii subsp. pragensis		23.81	0.13	XYL 1			
Staphylococcus petrasii subsp. jettensis		13.49	0.10				
400663	**Good identification**						**400663**
Aerococcus viridans		100.0	0.30	ARG 1			
400672	**Low discrimination**						**400672**
					NOV	YEP	
Aerococcus viridans		64.70	0.32	ARG 1	1	1	
Staphylococcus arlettae		35.30	0.15	ESL 90	99	99	
400673	**Low discrimination**						**400673**
					NOV	YEP	
Aerococcus viridans		76.34	0.39	ARG 1	1	1	
Staphylococcus arlettae		23.66	0.18	ESL 90	99	99	
400700	**Low discrimination**						**400700**
					OXI	NOV	
Staphylococcus auricularis		86.13	0.52	NIT 90	1	1	
Macrococcus carouselicus		10.50	0.27	GAL 10	99	99	
Staphylococcus carnosus subsp. utilis		2.11	0.03		1	1	
400701	**Low discrimination**						**400701**
					OXI	PYR	
Staphylococcus auricularis		81.55	0.33	NIT 90	1	40	
Kocuria kristinae		13.13	0.10	GAL 10	99	93	
Staphylococcus hyicus		5.33	0.00	MNS 10	1	1	
400702	**Acceptable identification**						**400702**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		92.75	0.33	NIT 90				
Staphylococcus piscifermentans		7.25	0.20	GAL 10				
400710	**Low discrimination**							**400710**
					OXI	NOV		
Macrococcus bovicus		58.20	0.31	ARG 1	99	99		
Macrococcus carouselicus		34.21	0.23		99	99		
Staphylococcus auricularis		4.43	0.11		1	1		
400713	**Good identification**							**400713**
Aerococcus viridans		100.0	0.27	ARG 1				
400733	**Good identification**							**400733**
Aerococcus viridans		100.0	0.27	ARG 1				
400740	**Low discrimination**							**400740**
					PYR			
Staphylococcus auricularis		80.97	0.60	NIT 90	40			
Staphylococcus haemolyticus		10.40	0.43	GAL 10	98			
Staphylococcus warneri		3.88	0.41		8			
400741	**Low discrimination**							**400741**
					OXI			
Staphylococcus auricularis		59.91	0.40	NIT 90	1			
Kocuria kristinae		37.21	0.30	GAL 10	99			
Staphylococcus haemolyticus		2.14	0.12	MNS 10	1			
400742	**Low discrimination**							**400742**
					PYR			
Staphylococcus auricularis		53.15	0.40	NIT 90	40			
Staphylococcus haemolyticus		34.55	0.38	GAL 10	98			
Staphylococcus warneri		7.63	0.31	LAC 10	8			
400750	**Low discrimination**							**400750**
					OXI	PYR	NOV	
Macrococcus bovicus		51.77	0.39	ARG 1	99	1	99	
Staphylococcus haemolyticus		29.43	0.38		1	98	2	
Staphylococcus warneri		13.53	0.38		1	8	1	
400752	**Low discrimination**							**400752**
					PYR			
Staphylococcus haemolyticus		65.75	0.33	VPT 86	98			
Staphylococcus warneri		17.91	0.28	NIT 99	8			
Aerococcus viridans		11.79	0.32		95			
400753	**Acceptable identification**							**400753**
Aerococcus viridans		93.12	0.39	ARG 1				
Staphylococcus haemolyticus		6.89	0.02	ESL 90				
400763	**Good identification**							**400763**
Aerococcus viridans		100.0	0.30	ARG 1				
400772	**Low discrimination**							**400772**
					NOV	YEP		
Staphylococcus arlettae		75.56	0.31	ARG 1	99	99		
Aerococcus viridans		24.44	0.32	GLR 92	1	1		
400773	**Low discrimination**							**400773**
					NOV	YEP		
Staphylococcus arlettae		63.72	0.34	ARG 1	99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		36.28	0.39	GLR 92	1	1	
401000	**Low discrimination**						**401000**
					NOV	OXI	
Staphylococcus carnosus subsp. utilis		58.53	0.29	ESL 1	1	1	
Macrococcus carouselicus		17.78	0.28		99	99	
Micrococcus sp.		15.43	0.18		1	89	
401010	**Low discrimination**						**401010**
Macrococcus equipersicus		44.21	0.34	URE 86			
Macrococcus bovicus		27.51	0.28	ARG 1			
Macrococcus carouselicus		26.88	0.24				
401011	**Good identification**						**401011**
Staphylococcus capitis subsp. capitis		100.0	0.31	ESL 1			
401050	**Low discrimination**						**401050**
Macrococcus equipersicus		65.07	0.43	URE 86			
Macrococcus bovicus		34.93	0.36	ARG 1			
401053	**Good identification**						**401053**
Aerococcus viridans		100.0	0.33	ARG 1			
401073	**Good identification**						**401073**
Aerococcus viridans		100.0	0.33	ARG 1			
401150	**Low discrimination**						**401150**
Macrococcus equipersicus		65.07	0.35	URE 86			
Macrococcus bovicus		34.93	0.28	ARG 1			
401153	**Good identification**						**401153**
Aerococcus viridans		100.0	0.33	ARG 1			
401173	**Good identification**						**401173**
Aerococcus viridans		100.0	0.33	ARG 1			
401200	**Acceptable identification**						**401200**
					NOV	OXI	
Macrococcus carouselicus		92.04	0.34	ARG 1	99	99	
Micrococcus sp.		3.89	-.03	TRE 94	1	89	
Staphylococcus auricularis		2.56	0.09		1	1	
401210	**Low discrimination**						**401210**
Staphylococcus vitulinus		48.40	0.29	ARG 1			
Macrococcus carouselicus		42.18	0.30	NIT 91			
Macrococcus bovicus		5.32	0.16				
401211	**Good identification**						**401211**
Staphylococcus capitis subsp. capitis		100.0	0.37	ESL 1			
401230	**Good identification**						**401230**
Staphylococcus vitulinus		100.0	0.38	ARG 1			
401253	**Good identification**						**401253**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.33	ARG 1			
401273	**Good identification**						**401273**
Aerococcus viridans		100.0	0.33	ARG 1			
401300	**Good identification**						**401300**
Macrococcus carouselicus		100.0	0.27	ARG 1			
401353	**Good identification**						**401353**
Aerococcus viridans		100.0	0.33	ARG 1			
401373	**Good identification**						**401373**
Aerococcus viridans		100.0	0.33	ARG 1			
401400	**Low discrimination**						**401400**
					OXI	NOV	
Staphylococcus carnosus subsp. utilis		50.99	0.44	ESL 1	1	1	
Macrococcus carouselicus		46.22	0.53		99	99	
Staphylococcus auricularis		2.07	0.32		1	1	
401402	**Low discrimination**						**401402**
					OXI	NOV	
Staphylococcus piscifermentans		32.88	0.32	URE 90	1	1	
Staphylococcus carnosus subsp. utilis		23.16	0.03	NIT 99	1	1	
Macrococcus carouselicus		21.00	0.12		99	99	
401403	**Good identification**						**401403**
Aerococcus viridans		100.0	0.37	ARG 1			
401410	**Low discrimination**						**401410**
Macrococcus carouselicus		81.07	0.49	ARG 1			
Macrococcus equipersicus		8.51	0.34				
Macrococcus bovicus		7.94	0.32				
401412	**Low discrimination**						**401412**
					OXI	NOV	
Aerococcus viridans		57.87	0.39	ARG 1	1	1	
Staphylococcus piscifermentans		21.58	0.25		1	1	
Macrococcus carouselicus		20.56	0.08		99	99	
401413	**Good identification**						**401413**
Aerococcus viridans		100.0	0.47	ARG 1			
401422	**Good identification**						**401422**
Aerococcus viridans		100.0	0.29	ARG 1			
401423	**Good identification**						**401423**
Aerococcus viridans		100.0	0.37	ARG 1			
401431	**Good identification**						**401431**
Aerococcus viridans		100.0	0.27	ARG 1			
401432	**Good identification**						**401432**
Aerococcus viridans		100.0	0.39	ARG 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests				Code
401433	**Good identification**								**401433**
Aerococcus viridans		100.0	0.47	ARG 1					
401440	**Low discrimination**								**401440**
Staphylococcus auricularis		67.58	0.39	ESL 1					
Staphylococcus carnosus subsp. utilis		7.20	0.03	NIT 90					
Staphylococcus piscifermentans		6.81	0.28						
401442	**Low discrimination**								**401442**
Aerococcus viridans		46.90	0.42	ARG 1					
Staphylococcus piscifermentans		30.60	0.32						
Staphylococcus auricularis		22.50	0.20						
401443	**Good identification**								**401443**
Aerococcus viridans		100.0	0.49	ARG 1					
401450	**Low discrimination**								**401450**
					OXI	NOV			
Macrococcus equipersicus		51.46	0.43	URE 86	99	99			
Macrococcus bovicus		41.44	0.39	ARG 1	99	99			
Aerococcus viridans		2.29	0.32		1	1			
401451	**Low discrimination**								**401451**
					OXI	NOV	PYR	YEP	
Aerococcus viridans		65.82	0.39	ARG 1	1	1	95	1	
Staphylococcus cohnii subsp. cohnii		22.63	0.18	LAC 90	1	99	1	12	
Macrococcus equipersicus		6.40	0.02		99	99	1	99	
401452	**Low discrimination**								**401452**
					OXI	NOV			
Aerococcus viridans		87.82	0.51	ARG 1	1	1			
Staphylococcus piscifermentans		8.19	0.25		1	1			
Macrococcus equipersicus		2.21	0.02		99	99			
401453	**Very good identification**								**401453**
Aerococcus viridans		100.0	0.59	ARG 1					
401461	**Good identification**								**401461**
Aerococcus viridans		100.0	0.30	ARG 1					
401462	**Good identification**								**401462**
Aerococcus viridans		100.0	0.42	ARG 1					
401463	**Good identification**								**401463**
Aerococcus viridans		100.0	0.49	ARG 1					
401470	**Low discrimination**								**401470**
					OXI	NOV			
Aerococcus viridans		70.94	0.32	ARG 1	1	1			
Macrococcus equipersicus		16.09	0.02	LAC 90	99	99			
Macrococcus bovicus		12.96	-.02		99	99			
401471	**Good identification**								**401471**
Aerococcus viridans		100.0	0.39	ARG 1					
401472	**Good identification**								**401472**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.51	ARG 1			
401473	**Very good identification**						**401473**
Aerococcus viridans		100.0	0.59	ARG 1			
401500	**Acceptable identification**						**401500**
					OXI	NOV	
Macrococcus carouselicus		92.76	0.45	ARG 1	99	99	
Staphylococcus piscifermentans		3.42	0.32		1	1	
Staphylococcus carnosus subsp. utilis		2.41	0.03		1	1	
401502	**Low discrimination**						**401502**
					OXI	NOV	
Staphylococcus piscifermentans		69.55	0.36	URE 90	1	1	
Aerococcus viridans		17.76	0.29	NIT 99	1	1	
Macrococcus carouselicus		12.69	0.04		99	99	
401503	**Good identification**						**401503**
Aerococcus viridans		100.0	0.37	ARG 1			
401510	**Low discrimination**						**401510**
Macrococcus carouselicus		80.97	0.41	ARG 1			
Macrococcus equipersicus		8.50	0.26				
Macrococcus bovicus		7.93	0.24				
401511	**Low discrimination**						**401511**
					OXI	NOV	
Aerococcus viridans		63.01	0.27	ARG 1	1	1	
Macrococcus carouselicus		37.00	0.00	LAC 90	99	99	
401512	**Low discrimination**						**401512**
					OXI	NOV	
Aerococcus viridans		58.43	0.39	ARG 1	1	1	
Staphylococcus piscifermentans		32.68	0.28		1	1	
Macrococcus carouselicus		8.89	0.00		99	99	
401513	**Good identification**						**401513**
Aerococcus viridans		100.0	0.47	ARG 1			
401522	**Good identification**						**401522**
Aerococcus viridans		100.0	0.29	ARG 1			
401523	**Good identification**						**401523**
Aerococcus viridans		100.0	0.37	ARG 1			
401531	**Good identification**						**401531**
Aerococcus viridans		100.0	0.27	ARG 1			
401532	**Good identification**						**401532**
Aerococcus viridans		100.0	0.39	ARG 1			
401533	**Good identification**						**401533**
Aerococcus viridans		100.0	0.47	ARG 1			
401540	**Low discrimination**						**401540**
					OXI	NOV	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		41.02	0.32	URE 90	1	1	
Staphylococcus auricularis		30.16	0.20	NIT 99	1	1	
Macrococcus carouselicus		11.23	0.04		99	99	
401541	**Good identification**						**401541**
Aerococcus viridans		100.0	0.30	ARG 1			
401542	**Low discrimination**						**401542**
Aerococcus viridans		50.54	0.42	ARG 1			
Staphylococcus piscifermentans		49.46	0.36				
401543	**Good identification**						**401543**
Aerococcus viridans		100.0	0.49	ARG 1			
401550	**Low discrimination**						**401550**
					OXI	NOV	
Macrococcus equipersicus		48.78	0.35	URE 86	99	99	
Macrococcus bovicus		39.28	0.32	ARG 1	99	99	
Aerococcus viridans		5.07	0.32		1	1	
401551	**Good identification**						**401551**
Aerococcus viridans		100.0	0.39	ARG 1			
401552	**Low discrimination**						**401552**
					NOV	YEP	
Aerococcus viridans		86.15	0.51	ARG 1	1	1	
Staphylococcus piscifermentans		12.05	0.28		1	1	
Staphylococcus kloosii		1.81	0.18		99	99	
401553	**Very good identification**						**401553**
Aerococcus viridans		100.0	0.59	ARG 1			
401561	**Good identification**						**401561**
Aerococcus viridans		100.0	0.30	ARG 1			
401562	**Good identification**						**401562**
Aerococcus viridans		100.0	0.42	ARG 1			
401563	**Good identification**						**401563**
Aerococcus viridans		100.0	0.49	ARG 1			
401570	**Good identification**						**401570**
Aerococcus viridans		100.0	0.32	ARG 1			
401571	**Good identification**						**401571**
Aerococcus viridans		100.0	0.39	ARG 1			
401572	**Good identification**						**401572**
Aerococcus viridans		100.0	0.51	ARG 1			
401573	**Very good identification**						**401573**
Aerococcus viridans		100.0	0.59	ARG 1			
401600	**Very good identification**						**401600**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Micrococcus carouselicus		96.74	0.59	ARG 1	99	99	
Staphylococcus auricularis		1.97	0.31		1	1	
Staphylococcus piscifermentans		0.75	0.32		1	1	
401602	**Low discrimination**						**401602**
					OXI	NOV	
Staphylococcus piscifermentans		43.22	0.36	URE 90	1	1	
Micrococcus carouselicus		37.36	0.18	NIT 99	99	99	
Aerococcus viridans		11.04	0.29		1	1	
401603	**Acceptable identification**						**401603**
					OXI		
Aerococcus viridans		90.25	0.37	ARG 1	1		
Kocuria kristinae		9.75	-.05		99		
401610	**Good identification**						**401610**
Micrococcus carouselicus		93.88	0.55	ARG 1			
Staphylococcus vitulinus		3.87	0.24				
Micrococcus bovicus		1.13	0.19				
401612	**Low discrimination**						**401612**
					OXI	NOV	
Aerococcus viridans		43.85	0.39	ARG 1	1	1	
Micrococcus carouselicus		31.62	0.14		99	99	
Staphylococcus piscifermentans		24.53	0.28		1	1	
401613	**Good identification**						**401613**
Aerococcus viridans		100.0	0.47	ARG 1			
401622	**Good identification**						**401622**
Aerococcus viridans		100.0	0.29	ARG 1			
401623	**Good identification**						**401623**
Aerococcus viridans		100.0	0.37	ARG 1			
401630	**Acceptable identification**						**401630**
					OXI	NOV	
Staphylococcus vitulinus		90.53	0.33	ARG 1	99	99	
Micrococcus carouselicus		8.21	0.14	NIT 91	99	99	
Aerococcus viridans		1.27	0.19		1	1	
401631	**Low discrimination**						**401631**
					OXI	NOV	
Aerococcus viridans		76.34	0.27	ARG 1	1	1	
Staphylococcus vitulinus		23.66	-.08	LAC 90	99	99	
401632	**Acceptable identification**						**401632**
					OXI	NOV	
Aerococcus viridans		92.56	0.39	ARG 1	1	1	
Staphylococcus vitulinus		7.44	-.08		99	99	
401633	**Good identification**						**401633**
Aerococcus viridans		100.0	0.47	ARG 1			
401640	**Low discrimination**						**401640**
Staphylococcus auricularis		49.37	0.39	ESL 1			
Staphylococcus petrasii subsp. pragensis		19.06	0.13	NIT 90			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus petrasii subsp. jettensis		10.80	0.10					
401641	**Low discrimination**							**401641**
					OXI			
Kocuria kristinae		76.37	0.27	VPT 90	99			
Staphylococcus auricularis		14.91	0.19	ARG 1	1			
Aerococcus viridans		8.73	0.30		1			
401642	**Low discrimination**							**401642**
Aerococcus viridans		38.83	0.42	ARG 1				
Staphylococcus piscifermentans		38.00	0.36					
Staphylococcus auricularis		17.19	0.19					
401643	**Low discrimination**							**401643**
					OXI			
Aerococcus viridans		80.45	0.49	ARG 1	1			
Kocuria kristinae		19.56	0.15		99			
401650	**Low discrimination**							**401650**
Macrococcus bovicus		44.31	0.27	ARG 1				
Macrococcus equipersicus		21.77	0.22	ESL 10				
Macrococcus carouselicus		15.90	0.14					
401651	**Low discrimination**							**401651**
					OXI	NOV	PYR	
Aerococcus viridans		68.84	0.39	ARG 1	1	1	95	
Staphylococcus sciuri		22.79	0.13	LAC 90	99	99	1	
Kocuria kristinae		8.37	-.02		99	1	93	
401652	**Low discrimination**							**401652**
Aerococcus viridans		87.73	0.51	ARG 1				
Staphylococcus piscifermentans		12.27	0.28					
401653	**Very good identification**							**401653**
Aerococcus viridans		100.0	0.59	ARG 1				
401661	**Good identification**							**401661**
Aerococcus viridans		100.0	0.30	ARG 1				
401662	**Good identification**							**401662**
Aerococcus viridans		100.0	0.42	ARG 1				
401663	**Good identification**							**401663**
Aerococcus viridans		100.0	0.49	ARG 1				
401670	**Low discrimination**							**401670**
					OXI	NOV		
Aerococcus viridans		84.69	0.32	ARG 1	1	1		
Staphylococcus vitulinus		15.31	-.08	LAC 90	99	99		
401671	**Low discrimination**							**401671**
					OXI	NOV		
Aerococcus viridans		80.02	0.39	ARG 1	1	1		
Staphylococcus sciuri		19.98	0.11	LAC 90	99	99		
401672	**Acceptable identification**							**401672**
					NOV	YEP		
Aerococcus viridans		94.28	0.51	ARG 1	1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus arlettae		5.72	0.15		99	99	
401673	**Very good identification**						**401673**
					NOV	YEP	
Aerococcus viridans		96.67	0.59	ARG 1	1	1	
Staphylococcus arlettae		3.33	0.18		99	99	
401700	**Good identification**						**401700**
					OXI	NOV	
Macrococcus carouselicus		96.85	0.51	ARG 1	99	99	
Staphylococcus piscifermentans		2.64	0.36		1	1	
Staphylococcus auricularis		0.51	0.11		1	1	
401702	**Low discrimination**						**401702**
					OXI	NOV	
Staphylococcus piscifermentans		70.56	0.39	URE 90	1	1	
Macrococcus carouselicus		17.43	0.10	NIT 99	99	99	
Aerococcus viridans		12.02	0.29		1	1	
401703	**Good identification**						**401703**
Aerococcus viridans		100.0	0.37	ARG 1			
401710	**Good identification**						**401710**
					OXI	NOV	
Macrococcus carouselicus		96.24	0.47	ARG 1	99	99	
Staphylococcus piscifermentans		1.76	0.28		1	1	
Macrococcus bovicus		1.16	0.12		99	99	
401712	**Low discrimination**						**401712**
					OXI	NOV	
Aerococcus viridans		46.55	0.39	ARG 1	1	1	
Staphylococcus piscifermentans		39.06	0.32		1	1	
Macrococcus carouselicus		14.39	0.06		99	99	
401713	**Acceptable identification**						**401713**
					OXI	NOV	
Aerococcus viridans		91.12	0.47	ARG 1	1	1	
Staphylococcus lentus		8.88	0.12		99	99	
401722	**Good identification**						**401722**
Aerococcus viridans		100.0	0.29	ARG 1			
401723	**Good identification**						**401723**
Aerococcus viridans		100.0	0.37	ARG 1			
401731	**Low discrimination**						**401731**
					OXI	NOV	
Aerococcus viridans		65.99	0.27	ARG 1	1	1	
Staphylococcus sciuri		34.01	0.05	LAC 90	99	99	
401732	**Good identification**						**401732**
Aerococcus viridans		100.0	0.39	ARG 1			
401733	**Acceptable identification**						**401733**
					OXI	NOV	
Aerococcus viridans		91.12	0.47	ARG 1	1	1	
Staphylococcus lentus		8.88	0.12		99	99	
401740	**Low discrimination**						**401740**
					OXI	NOV	
Staphylococcus piscifermentans		50.14	0.36	URE 90	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		22.68	0.19	NIT 99	1	1		
Macrococcus carouselicus		18.57	0.10		99	99		
401742	**Low discrimination**							**401742**
Staphylococcus piscifermentans		59.48	0.39	URE 90				
Aerococcus viridans		40.52	0.42	NIT 99				
401743	**Acceptable identification**							**401743**
					OXI			
Aerococcus viridans		92.51	0.49	ARG 1	1			
Kocuria kristinae		7.50	0.05		99			
401751	**Low discrimination**							**401751**
					OXI	NOV		
Aerococcus viridans		56.56	0.39	ARG 1	1	1		
Staphylococcus sciuri		36.34	0.19	LAC 90	99	99		
Staphylococcus lentus		7.10	0.07		99	99		
401752	**Low discrimination**							**401752**
Aerococcus viridans		82.66	0.51	ARG 1				
Staphylococcus piscifermentans		17.34	0.32					
401753	**Good identification**							**401753**
					OXI	NOV		
Aerococcus viridans		94.25	0.59	ARG 1	1	1		
Staphylococcus lentus		4.66	0.18		99	99		
Staphylococcus sciuri		1.10	0.03		99	99		
401761	**Good identification**							**401761**
Aerococcus viridans		100.0	0.30	ARG 1				
401762	**Good identification**							**401762**
Aerococcus viridans		100.0	0.42	ARG 1				
401763	**Good identification**							**401763**
Aerococcus viridans		100.0	0.49	ARG 1				
401770	**Good identification**							**401770**
Aerococcus viridans		100.0	0.32	ARG 1				
401771	**Low discrimination**							**401771**
					OXI	NOV		
Aerococcus viridans		62.10	0.39	ARG 1	1	1		
Staphylococcus sciuri		30.11	0.17	LAC 90	99	99		
Staphylococcus lentus		7.79	0.07		99	99		
401772	**Low discrimination**							**401772**
					NOV	YEP		
Aerococcus viridans		74.43	0.51	ARG 1	1	1		
Staphylococcus arlettae		25.57	0.31		99	99		
401773	**Low discrimination**							**401773**
					OXI	NOV	YEP	
Aerococcus viridans		79.78	0.59	ARG 1	1	1	1	
Staphylococcus arlettae		15.57	0.34		1	99	99	
Staphylococcus lentus		3.94	0.18		99	99	1	
402000	**Good identification**							**402000**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. utilis		93.75	0.85				
Staphylococcus auricularis		2.98	0.71				
Kocuria rosea		1.93	0.59				
402001	**Low discrimination**						**402001**
Staphylococcus capitis subsp. capitis		73.51	0.68	MAN 93			
Staphylococcus carnosus subsp. utilis		17.75	0.44				
Staphylococcus auricularis		6.21	0.51				
402002	**Low discrimination**						**402002**
Staphylococcus carnosus subsp. utilis		70.43	0.44	LAC 1			
Staphylococcus auricularis		24.65	0.51				
Kocuria varians		2.47	0.32				
402003	**Low discrimination**						**402003**
Staphylococcus capitis subsp. capitis		52.24	0.33	MAN 93			
Staphylococcus auricularis		24.04	0.31	LAC 2			
Staphylococcus carnosus subsp. carnosus		15.60	0.23				
402010	**Low discrimination**						**402010**
Staphylococcus carnosus subsp. utilis		53.52	0.44	MAN 1			
Staphylococcus capitis subsp. capitis		29.74	0.50				
Kocuria rosea		12.09	0.39				
402011	**Very good identification**						**402011**
Staphylococcus capitis subsp. capitis		99.91	0.91				
Staphylococcus carnosus subsp. carnosus		0.05	0.23				
Kocuria rosea		0.02	0.13				
402013	**Good identification**						**402013**
Staphylococcus capitis subsp. capitis		91.65	0.56	LAC 2			
Staphylococcus carnosus subsp. carnosus		8.24	0.36				
Staphylococcus capitis subsp. ureolyticus		0.12	-.13				
402020	**Low discrimination**						**402020**
Staphylococcus carnosus subsp. utilis		51.03	0.44	XYL 1			
Kocuria rosea		44.47	0.51				
Kocuria varians		2.23	0.34				
402030	**Low discrimination**						**402030**
					NOV	OXI	
Kocuria rosea		79.79	0.32	ARG 1	1	35	
Staphylococcus carnosus subsp. utilis		8.33	0.03	MAN 10	1	1	
Staphylococcus vitulinus		7.26	0.09		99	99	
402031	**Good identification**						**402031**
Staphylococcus capitis subsp. capitis		99.09	0.50	XYL 1			
Kocuria rosea		0.91	0.05				
402040	**Low discrimination**						**402040**
Staphylococcus auricularis		87.13	0.78	TRE 92			
Staphylococcus carnosus subsp. utilis		11.85	0.44				
Kocuria varians		0.32	0.29				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
402041	**Good identification**						**402041**
Staphylococcus auricularis		93.77	0.59	TRE 92			
Staphylococcus capitis subsp. capitis		4.80	0.27	MNS 10			
Staphylococcus carnosus subsp. utilis		1.16	0.03				
402042	**Very good identification**						**402042**
Staphylococcus auricularis		97.14	0.59	TRE 92			
Staphylococcus carnosus subsp. utilis		1.20	0.03	LAC 10			
Kocuria varians		0.92	0.18				
402043	**Good identification**						**402043**
Staphylococcus auricularis		98.47	0.39	TRE 92			
Staphylococcus capitis subsp. ureolyticus		1.54	-.13	MNS 10			
402050	**Low discrimination**						**402050**
					OXI	NOV	
Macrococcus bovicus		47.64	0.36	ARG 1	99	99	
Staphylococcus auricularis		40.04	0.37	NIT 10	1	1	
Staphylococcus carnosus subsp. utilis		5.45	0.03		1	1	
402051	**Good identification**						**402051**
Staphylococcus capitis subsp. capitis		97.74	0.50	MLT 1			
Staphylococcus auricularis		1.45	0.17				
Staphylococcus capitis subsp. ureolyticus		0.81	-.02				
402060	**Low discrimination**						**402060**
Staphylococcus auricularis		72.27	0.37	TRE 92			
Staphylococcus carnosus subsp. utilis		9.83	0.03	XYL 1			
Kocuria varians		9.33	0.20				
402100	**Low discrimination**						**402100**
					OXI		
Staphylococcus carnosus subsp. utilis		57.13	0.44	GAL 1	1		
Staphylococcus auricularis		20.00	0.51		1		
Dermacoccus nishinomiyaensis		10.81	0.39		93		
402101	**Low discrimination**						**402101**
Staphylococcus capitis subsp. capitis		36.85	0.27	GAL 1			
Staphylococcus auricularis		34.26	0.31	MAN 93			
Kocuria varians		14.28	0.25				
402102	**Low discrimination**						**402102**
Kocuria varians		49.67	0.35	ARG 1			
Staphylococcus auricularis		36.73	0.31				
Staphylococcus carnosus subsp. utilis		9.54	0.03				
402111	**Good identification**						**402111**
Staphylococcus capitis subsp. capitis		98.88	0.50	GAL 1			
Staphylococcus carnosus subsp. carnosus		1.12	0.11				
402120	**Low discrimination**						**402120**
Kocuria varians		72.90	0.37	ARG 1			
Staphylococcus carnosus subsp. utilis		11.24	0.03				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Kocuria rosea		9.79	0.10				
402140	**Good identification**						**402140**
Staphylococcus auricularis		92.75	0.59	GAL 10			
Kocuria varians		4.65	0.33	TRE 92			
Staphylococcus carnosus subsp. utilis		1.15	0.03				
402141	**Good identification**						**402141**
Staphylococcus auricularis		96.23	0.39	GAL 10			
Kocuria varians		3.77	0.11	TRE 92			
402142	**Low discrimination**						**402142**
					PYR		
Staphylococcus auricularis		82.56	0.39	GAL 10	40		
Kocuria varians		10.50	0.22	TRE 92	10		
Staphylococcus haemolyticus		4.76	0.14	LAC 10	98		
402150	**Low discrimination**						**402150**
					OXI	NOV	
Macrococcus bovicus		73.40	0.28	ARG 1	99	99	
Staphylococcus auricularis		16.00	0.17	NIT 10	1	1	
Staphylococcus haemolyticus		10.60	0.15		1	2	
402200	**Low discrimination**						**402200**
Staphylococcus auricularis		67.15	0.70	TRE 92			
Staphylococcus carnosus subsp. utilis		23.09	0.44				
Kocuria rosea		5.22	0.39				
402201	**Very good identification**						**402201**
Staphylococcus capitis subsp. capitis		95.66	0.73	MAN 93			
Staphylococcus auricularis		4.02	0.50				
Kocuria rosea		0.15	0.13				
402202	**Acceptable identification**						**402202**
Staphylococcus auricularis		92.61	0.50	TRE 92			
Staphylococcus carnosus subsp. utilis		2.90	0.03	LAC 10			
Kocuria varians		1.37	0.14				
402203	**Low discrimination**						**402203**
Staphylococcus capitis subsp. capitis		70.69	0.39	MAN 93			
Staphylococcus auricularis		16.17	0.31	LAC 2			
Staphylococcus hyicus		11.62	0.19				
402210	**Low discrimination**						**402210**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		72.01	0.55	MNS 99	1	1	
Staphylococcus vitulinus		22.49	0.41		99	99	
Staphylococcus auricularis		2.05	0.29		1	1	
402211	**Excellent identification**						**402211**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		99.99	0.96		1	1	
Staphylococcus auricularis		0.00	0.09		1	1	
Staphylococcus vitulinus		0.00	0.00		99	99	
402213	**Very good identification**						**402213**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus capitis subsp. capitis		98.04	0.62	LAC 2			
Staphylococcus capitis subsp. ureolyticus		1.60	0.16				
Staphylococcus simulans		0.37	0.16				
402220	**Low discrimination**						**402220**
Kocuria rosea		64.09	0.32	ARG 1			
Staphylococcus auricularis		19.45	0.29	SUC 10			
Staphylococcus carnosus subsp. utilis		6.69	0.03				
402221	**Acceptable identification**						**402221**
Staphylococcus capitis subsp. capitis		90.29	0.32	MAN 93			
Kocuria rosea		5.92	0.05	XYL 1			
Staphylococcus auricularis		3.79	0.09				
402230	**Good identification**						**402230**
					NOV	OXI	
Staphylococcus vitulinus		97.63	0.50	ARG 1	99	99	
Kocuria rosea		1.21	0.12		1	35	
Staphylococcus capitis subsp. capitis		1.17	0.14		1	1	
402231	**Very good identification**						**402231**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		99.15	0.55	XYL 1	1	1	
Staphylococcus vitulinus		0.85	0.09		99	99	
402240	**Low discrimination**						**402240**
Staphylococcus auricularis		87.70	0.78	TRE 92			
Staphylococcus haemolyticus		10.56	0.60				
Staphylococcus petrasii subsp. jettensis		0.58	0.18				
402241	**Low discrimination**						**402241**
					PYR		
Staphylococcus auricularis		87.18	0.58	TRE 92	40		
Staphylococcus capitis subsp. capitis		8.98	0.32	MNS 10	1		
Staphylococcus haemolyticus		2.92	0.29		98		
402242	**Low discrimination**						**402242**
					PYR		
Staphylococcus auricularis		60.17	0.58	TRE 92	40		
Staphylococcus haemolyticus		36.69	0.54	LAC 10	98		
Staphylococcus hominis subsp. hominis		1.87	0.29		1		
402243	**Low discrimination**						**402243**
					PYR		
Staphylococcus auricularis		58.18	0.38	TRE 92	40		
Staphylococcus capitis subsp. ureolyticus		23.58	0.16	MNS 10	1		
Staphylococcus haemolyticus		9.87	0.23	LAC 10	98		
402250	**Low discrimination**						**402250**
					OXI	NOV	
Staphylococcus haemolyticus		81.40	0.55	VPT 86	1	2	
Staphylococcus auricularis		11.62	0.36	TRE 93	1	1	
Macrococcus bovicus		3.75	0.23		99	99	
402251	**Low discrimination**						**402251**
					PYR		
Staphylococcus capitis subsp. capitis		88.54	0.55	MLT 1	1		
Staphylococcus capitis subsp. ureolyticus		9.53	0.26		1		
Staphylococcus haemolyticus		1.27	0.24		98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
402252	**Acceptable identification**					**402252**
					PYR	
Staphylococcus haemolyticus		93.31	0.50	VPT 86	98	
Staphylococcus capitis subsp. ureolyticus		2.84	0.03	TRE 93	1	
Staphylococcus auricularis		2.63	0.17		40	
402253	**Acceptable identification**					**402253**
					COA	
Staphylococcus capitis subsp. ureolyticus		92.07	0.44	URE 99	1	
Staphylococcus delphini		4.20	0.18		99	
Staphylococcus capitis subsp. capitis		2.38	0.21		1	
402260	**Low discrimination**					**402260**
					PYR	
Staphylococcus auricularis		87.76	0.36	TRE 92	40	
Staphylococcus haemolyticus		10.57	0.19	XYL 1	98	
Kocuria varians		1.67	0.02		10	
402300	**Low discrimination**					**402300**
					OXI	
Staphylococcus auricularis		83.99	0.50	GAL 10	1	
Kocuria varians		6.61	0.29	TRE 92	13	
Dermacoccus nishinomiyaensis		3.14	0.14		93	
402301	**Low discrimination**					**402301**
Staphylococcus capitis subsp. capitis		54.87	0.32	GAL 1		
Staphylococcus auricularis		25.36	0.31	MAN 93		
Staphylococcus hyicus		18.22	0.19			
402302	**Low discrimination**					**402302**
					PYR	
Staphylococcus auricularis		65.41	0.31	GAL 10	40	
Kocuria varians		13.07	0.17	TRE 92	10	
Staphylococcus haemolyticus		9.54	0.14	LAC 10	98	
402303	**Good identification**					**402303**
Staphylococcus hyicus		97.75	0.41	PHS 92		
Staphylococcus auricularis		1.32	0.11	TRE 92		
Staphylococcus capitis subsp. capitis		0.52	-.03			
402311	**Very good identification**					**402311**
Staphylococcus capitis subsp. capitis		100.0	0.55	GAL 1		
402340	**Low discrimination**					**402340**
					PYR	
Staphylococcus haemolyticus		53.65	0.61	VPT 86	98	
Staphylococcus auricularis		43.89	0.58	TRE 93	40	
Staphylococcus hominis subsp. hominis		1.26	0.29		1	
402341	**Low discrimination**					**402341**
					PYR	
Staphylococcus auricularis		63.94	0.38	GAL 10	40	
Staphylococcus haemolyticus		21.76	0.30	TRE 92	98	
Staphylococcus epidermidis		14.30	0.24	MNS 10	1	
402342	**Low discrimination**					**402342**
					PYR	
Staphylococcus haemolyticus		77.84	0.56	VPT 86	98	
Staphylococcus auricularis		12.58	0.38	TRE 93	40	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		4.80	0.29		1		
402343	**Low discrimination**						**402343**
					PYR		
Staphylococcus epidermidis		66.72	0.34	VPT 96	1		
Staphylococcus haemolyticus		18.03	0.24	URE 92	98		
Staphylococcus auricularis		10.47	0.19		40		
402350	**Very good identification**						**402350**
					OXI	NOV	
Staphylococcus haemolyticus		96.38	0.56	VPT 86	1	2	
Macrococcus bovicus		1.69	0.16	TRE 93	99	99	
Staphylococcus auricularis		1.36	0.17		1	1	
402352	**Good identification**						**402352**
					PYR		
Staphylococcus haemolyticus		98.28	0.51	VPT 86	98		
Staphylococcus hominis subsp. hominis		1.22	0.13	TRE 93	1		
Staphylococcus piscifermentans		0.50	0.12		15		
402400	**Good identification**						**402400**
Staphylococcus carnosus subsp. utilis		93.46	1.00				
Staphylococcus auricularis		6.52	0.92				
Staphylococcus piscifermentans		0.02	0.50				
402401	**Low discrimination**						**402401**
Staphylococcus carnosus subsp. utilis		55.95	0.59	MNS 1			
Staphylococcus auricularis		42.90	0.73				
Staphylococcus capitis subsp. capitis		0.90	0.33				
402402	**Low discrimination**						**402402**
Staphylococcus carnosus subsp. utilis		55.64	0.59	LAC 1			
Staphylococcus auricularis		42.66	0.73				
Staphylococcus piscifermentans		1.67	0.54				
402403	**Low discrimination**						**402403**
Staphylococcus auricularis		86.18	0.53	MNS 10			
Staphylococcus carnosus subsp. utilis		10.22	0.18	LAC 10			
Staphylococcus carnosus subsp. carnosus		2.08	0.16				
402410	**Good identification**						**402410**
					OXI	NOV	
Staphylococcus carnosus subsp. utilis		91.33	0.59	MAN 1	1	1	
Staphylococcus auricularis		6.37	0.51		1	1	
Macrococcus bovicus		0.99	0.32		99	99	
402411	**Good identification**						**402411**
Staphylococcus capitis subsp. capitis		90.47	0.56	TRE 2			
Staphylococcus carnosus subsp. utilis		4.27	0.18				
Staphylococcus auricularis		3.28	0.32				
402412	**Low discrimination**						**402412**
Staphylococcus piscifermentans		41.43	0.46	URE 90			
Staphylococcus carnosus subsp. utilis		32.50	0.18	ESL 90			
Staphylococcus auricularis		24.92	0.32				
402413	**Low discrimination**						**402413**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		56.25	0.28	BGA 90			
Staphylococcus capitis subsp. capitis		29.79	0.21	PHS 99			
Staphylococcus simulans		8.09	0.14				
402420	**Good identification**						**402420**
Staphylococcus carnosus subsp. utilis		93.28	0.59	XYL 1			
Staphylococcus auricularis		6.50	0.51				
Kocuria rosea		0.16	0.10				
402440	**Good identification**						**402440**
Staphylococcus auricularis		93.84	1.00				
Staphylococcus carnosus subsp. utilis		5.83	0.59				
Staphylococcus piscifermentans		0.12	0.50				
402441	**Very good identification**						**402441**
					NOV		
Staphylococcus auricularis		99.04	0.80	MNS 10	1		
Staphylococcus carnosus subsp. utilis		0.56	0.18		1		
Staphylococcus cohnii subsp. cohnii		0.36	0.34		99		
402442	**Very good identification**						**402442**
Staphylococcus auricularis		97.10	0.80	LAC 10			
Staphylococcus piscifermentans		1.63	0.54				
Staphylococcus haemolyticus		0.64	0.37				
402443	**Very good identification**						**402443**
Staphylococcus auricularis		99.67	0.61	MNS 10			
Staphylococcus haemolyticus		0.18	0.05	LAC 10			
Staphylococcus piscifermentans		0.15	0.12				
402450	**Low discrimination**						**402450**
					OXI	NOV	
Staphylococcus auricularis		71.65	0.59	MAN 1	1	1	
Macrococcus bovicus		11.12	0.39		99	99	
Staphylococcus haemolyticus		5.41	0.37		1	2	
402451	**Low discrimination**						**402451**
					NOV		
Staphylococcus auricularis		61.87	0.39	MAN 1	1		
Staphylococcus cohnii subsp. cohnii		28.56	0.36	MNS 10	99		
Staphylococcus capitis subsp. capitis		7.40	0.15		1		
402452	**Low discrimination**						**402452**
Staphylococcus auricularis		47.43	0.39	MAN 1			
Staphylococcus piscifermentans		33.79	0.46	LAC 10			
Staphylococcus haemolyticus		18.11	0.32				
402460	**Good identification**						**402460**
Staphylococcus auricularis		94.15	0.59	XYL 1			
Staphylococcus carnosus subsp. utilis		5.85	0.18				
402461	**Good identification**						**402461**
Staphylococcus auricularis		100.0	0.39	XYL 1			
402462	**Good identification**						**402462**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		98.35	0.39	XYL 1			
Staphylococcus piscifermentans		1.65	0.12	LAC 10			
402500	**Low discrimination**						**402500**
Staphylococcus carnosus subsp. utilis		55.54	0.59	GAL 1			
Staphylococcus auricularis		42.59	0.73				
Staphylococcus piscifermentans		1.67	0.54				
402501	**Low discrimination**						**402501**
Staphylococcus auricularis		86.63	0.53	GAL 10			
Staphylococcus carnosus subsp. utilis		10.27	0.18	MNS 10			
Staphylococcus schleiferi subsp. schleiferi		1.74	0.14				
402502	**Low discrimination**						**402502**
Staphylococcus auricularis		60.37	0.53	GAL 10			
Staphylococcus piscifermentans		31.94	0.57	LAC 10			
Staphylococcus carnosus subsp. utilis		7.16	0.18				
402503	**Low discrimination**						**402503**
Staphylococcus auricularis		55.69	0.34	GAL 10			
Staphylococcus hyicus		38.61	0.22	MNS 10			
Staphylococcus carnosus subsp. carnosus		3.03	0.03	LAC 10			
402510	**Low discrimination**						**402510**
Staphylococcus piscifermentans		36.14	0.46	URE 90			
Staphylococcus carnosus subsp. utilis		28.35	0.18	ESL 90			
Staphylococcus auricularis		21.74	0.32				
402512	**Good identification**						**402512**
Staphylococcus piscifermentans		95.73	0.50	URE 90			
Staphylococcus auricularis		4.27	0.12	ESL 90			
402540	**Very good identification**						**402540**
Staphylococcus auricularis		96.32	0.80	GAL 10			
Staphylococcus piscifermentans		1.62	0.54				
Staphylococcus haemolyticus		1.27	0.43				
402541	**Very good identification**						**402541**
					NOV	PYR	
Staphylococcus auricularis		99.12	0.61	GAL 10	1	40	
Staphylococcus cohnii subsp. cohnii		0.36	0.14	MNS 10	99	1	
Staphylococcus haemolyticus		0.36	0.12		2	98	
402542	**Low discrimination**						**402542**
Staphylococcus auricularis		75.80	0.61	GAL 10			
Staphylococcus piscifermentans		17.18	0.57	LAC 10			
Staphylococcus haemolyticus		5.05	0.38				
402543	**Good identification**						**402543**
Staphylococcus auricularis		96.23	0.41	GAL 10			
Staphylococcus piscifermentans		1.98	0.16	MNS 10			
Staphylococcus haemolyticus		1.79	0.07	LAC 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
402550	**Low discrimination**						**402550**
Staphylococcus auricularis		31.83	0.39	GAL 10			
Staphylococcus haemolyticus		24.37	0.38	MAN 1			
Staphylococcus piscifermentans		22.68	0.46				
402552	**Low discrimination**						**402552**
Staphylococcus piscifermentans		66.36	0.50	URE 90			
Staphylococcus haemolyticus		26.74	0.33	ESL 90			
Staphylococcus auricularis		6.90	0.20				
402560	**Good identification**						**402560**
Staphylococcus auricularis		97.09	0.39	GAL 10			
Staphylococcus piscifermentans		1.63	0.12	XYL 1			
Staphylococcus haemolyticus		1.28	0.02				
402600	**Low discrimination**						**402600**
Staphylococcus auricularis		85.69	0.92				
Staphylococcus carnosus subsp. utilis		13.45	0.59				
Staphylococcus piscifermentans		0.40	0.54				
402601	**Good identification**						**402601**
Staphylococcus auricularis		93.88	0.72	MNS 10			
Staphylococcus capitis subsp. capitis		3.97	0.39				
Staphylococcus carnosus subsp. utilis		1.34	0.18				
402602	**Good identification**						**402602**
Staphylococcus auricularis		90.87	0.72	LAC 10			
Staphylococcus piscifermentans		5.79	0.57				
Staphylococcus haemolyticus		1.51	0.37				
402603	**Low discrimination**						**402603**
Staphylococcus auricularis		54.62	0.52	MNS 10			
Staphylococcus hyicus		39.24	0.41	LAC 10			
Staphylococcus simulans		4.94	0.30				
402610	**Low discrimination**						**402610**
					OXI	NOV	
Staphylococcus auricularis		45.25	0.51	MAN 1	1	1	
Staphylococcus vitulinus		24.28	0.36		99	99	
Staphylococcus piscifermentans		9.05	0.46		1	1	
402611	**Very good identification**						**402611**
					PYR		
Staphylococcus capitis subsp. capitis		97.99	0.62	TRE 2	1		
Staphylococcus auricularis		1.76	0.31		40		
Staphylococcus haemolyticus		0.09	0.06		98		
402612	**Low discrimination**						**402612**
Staphylococcus piscifermentans		45.76	0.50	URE 90			
Staphylococcus simulans		20.12	0.31	ESL 90			
Staphylococcus auricularis		16.94	0.31				
402613	**Low discrimination**						**402613**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		74.48	0.42	URE 98			
Staphylococcus capitis subsp. capitis		21.24	0.27	BGA 90			
Staphylococcus auricularis		2.08	0.11				
402620	**Low discrimination**						**402620**
Staphylococcus auricularis		72.75	0.51	XYL 1			
Staphylococcus muscae		14.42	0.18				
Staphylococcus carnosus subsp. utilis		11.42	0.18				
402621	**Good identification**						**402621**
Staphylococcus auricularis		95.95	0.31	XYL 1			
Staphylococcus capitis subsp. capitis		4.05	-.03	MNS 10			
402622	**Acceptable identification**						**402622**
Staphylococcus auricularis		94.01	0.31	XYL 1			
Staphylococcus piscifermentans		5.99	0.16	LAC 10			
402630	**Good identification**						**402630**
					OXI	NOV	
Staphylococcus vitulinus		99.31	0.45	ARG 1	99	99	
Staphylococcus auricularis		0.69	0.10		1	1	
402640	**Low discrimination**						**402640**
Staphylococcus auricularis		82.99	0.99				
Staphylococcus haemolyticus		11.55	0.83				
Staphylococcus petrasii subsp. jettensis		4.71	0.59				
402641	**Very good identification**						**402641**
Staphylococcus auricularis		95.35	0.80	MNS 10			
Staphylococcus haemolyticus		3.69	0.52				
Staphylococcus petrasii subsp. jettensis		0.49	0.18				
402642	**Low discrimination**						**402642**
Staphylococcus auricularis		54.53	0.80	LAC 10			
Staphylococcus haemolyticus		38.41	0.78				
Staphylococcus petrasii subsp. jettensis		4.91	0.43				
402643	**Low discrimination**						**402643**
Staphylococcus auricularis		80.36	0.60	MNS 10			
Staphylococcus haemolyticus		15.76	0.47	LAC 10			
Staphylococcus simulans		1.40	0.22				
402650	**Low discrimination**						**402650**
Staphylococcus haemolyticus		86.46	0.78	VPT 86			
Staphylococcus auricularis		10.69	0.58				
Staphylococcus piscifermentans		0.92	0.46				
402651	**Low discrimination**						**402651**
					PYR		
Staphylococcus haemolyticus		63.49	0.47	VPT 86	98		
Staphylococcus auricularis		28.19	0.39	MNS 3	40		
Staphylococcus capitis subsp. capitis		6.78	0.21		1		
402652	**Good identification**						**402652**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		93.65	0.73	VPT 86			
Staphylococcus piscifermentans		2.65	0.50				
Staphylococcus auricularis		2.29	0.39				
402653	**Low discrimination**						**402653**
Staphylococcus haemolyticus		56.61	0.42	VPT 86			
Staphylococcus simulans		34.22	0.35	MNS 3			
Staphylococcus auricularis		4.97	0.19				
402660	**Low discrimination**						**402660**
Staphylococcus auricularis		83.15	0.58	XYL 1			
Staphylococcus haemolyticus		11.57	0.42				
Staphylococcus petrasii subsp. jettensis		4.72	0.18				
402661	**Good identification**						**402661**
Staphylococcus auricularis		96.27	0.39	XYL 1			
Staphylococcus haemolyticus		3.73	0.11	MNS 10			
402662	**Low discrimination**						**402662**
Staphylococcus auricularis		54.89	0.39	XYL 1			
Staphylococcus haemolyticus		38.67	0.37	LAC 10			
Staphylococcus petrasii subsp. jettensis		4.95	0.02				
402670	**Low discrimination**						**402670**
					OXI	NOV	
Staphylococcus haemolyticus		83.30	0.37	VPT 86	1	2	
Staphylococcus auricularis		10.30	0.17	XYL 1	1	1	
Staphylococcus vitulinus		6.40	0.04		99	99	
402672	**Good identification**						**402672**
Staphylococcus haemolyticus		100.0	0.32	VPT 86			
402700	**Low discrimination**						**402700**
Staphylococcus auricularis		89.76	0.72	GAL 10			
Staphylococcus piscifermentans		5.72	0.57				
Staphylococcus haemolyticus		2.99	0.43				
402701	**Low discrimination**						**402701**
Staphylococcus auricularis		56.89	0.52	GAL 10			
Staphylococcus hyicus		40.87	0.41	MNS 10			
Rothia mucilaginosa		0.95	0.05				
402702	**Low discrimination**						**402702**
Staphylococcus auricularis		47.36	0.52	GAL 10			
Staphylococcus piscifermentans		40.71	0.61	LAC 10			
Staphylococcus haemolyticus		7.98	0.38				
402703	**Very good identification**						**402703**
					YEP		
Staphylococcus hyicus		98.01	0.63	PHS 92	1		
Staphylococcus auricularis		1.32	0.33		1		
Staphylococcus chromogenes		0.42	0.18		95		
402710	**Low discrimination**						**402710**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		45.77	0.50	URE 90				
Staphylococcus haemolyticus		32.79	0.38	ESL 90				
Staphylococcus auricularis		16.94	0.31					
402712	**Low discrimination**							**402712**
Staphylococcus piscifermentans		75.67	0.53	URE 90				
Staphylococcus haemolyticus		20.33	0.33	ESL 90				
Staphylococcus auricularis		2.08	0.11					
402720	**Acceptable identification**							**402720**
Staphylococcus auricularis		91.16	0.31	GAL 10				
Staphylococcus piscifermentans		5.80	0.16	XYL 1				
Staphylococcus haemolyticus		3.03	0.02					
402740	**Low discrimination**							**402740**
Staphylococcus haemolyticus		57.47	0.84	VPT 86				
Staphylococcus auricularis		40.69	0.80					
Staphylococcus piscifermentans		1.11	0.57					
402741	**Low discrimination**							**402741**
Staphylococcus auricularis		69.64	0.60	GAL 10				
Staphylococcus haemolyticus		27.37	0.53	MNS 10				
Rothia mucilaginosa		1.16	0.13					
402742	**Low discrimination**							**402742**
Staphylococcus haemolyticus		82.80	0.79	VPT 86				
Staphylococcus auricularis		11.58	0.60					
Staphylococcus piscifermentans		4.27	0.61					
402743	**Low discrimination**							**402743**
					PYR			
Staphylococcus haemolyticus		55.39	0.48	VPT 86	98			
Staphylococcus auricularis		27.84	0.40	MNS 3	40			
Staphylococcus hyicus		8.96	0.22		1			
402750	**Very good identification**							**402750**
Staphylococcus haemolyticus		96.84	0.79	VPT 86				
Staphylococcus piscifermentans		1.37	0.50					
Staphylococcus auricularis		1.18	0.39					
402751	**Acceptable identification**							**402751**
					OXI	NOV		
Staphylococcus haemolyticus		91.63	0.48	VPT 86	1	2		
Staphylococcus auricularis		4.01	0.19	MNS 3	1	1		
Staphylococcus sciuri		2.24	0.13		99	99		
402752	**Very good identification**							**402752**
					PYR			
Staphylococcus haemolyticus		95.69	0.74	VPT 86	98			
Staphylococcus piscifermentans		3.60	0.53		15			
Staphylococcus hominis subsp. hominis		0.28	0.23		1			
402753	**Low discrimination**							**402753**
					OXI	COA	PYR	
Staphylococcus haemolyticus		79.36	0.43	VPT 86	1	1	98	
Staphylococcus aureus subsp. aureus		5.52	0.01	MNS 3	1	99	10	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lentus		4.82	0.18		99	1	6	
402760	**Low discrimination**							**402760**
Staphylococcus haemolyticus		57.89	0.43	VPT 86				
Staphylococcus auricularis		40.99	0.39	XYL 1				
Staphylococcus piscifermentans		1.12	0.16					
402762	**Low discrimination**							**402762**
Staphylococcus haemolyticus		83.94	0.38	VPT 86				
Staphylococcus auricularis		11.74	0.19	XYL 1				
Staphylococcus piscifermentans		4.33	0.20					
402770	**Good identification**							**402770**
Staphylococcus haemolyticus		100.0	0.38	VPT 86				
402772	**Good identification**							**402772**
Staphylococcus haemolyticus		96.38	0.33	VPT 86				
Staphylococcus piscifermentans		3.62	0.12	XYL 1				
403000	**Acceptable identification**							**403000**
Staphylococcus carnosus subsp. utilis		92.63	0.44	ESL 1				
Staphylococcus auricularis		2.95	0.30					
Kocuria rosea		1.90	0.18					
403001	**Low discrimination**							**403001**
Staphylococcus capitis subsp. capitis		75.42	0.27	ESL 1				
Staphylococcus carnosus subsp. utilis		18.21	0.03	MAN 93				
Staphylococcus auricularis		6.37	0.10					
403002	**Low discrimination**							**403002**
Staphylococcus piscifermentans		51.26	0.32	URE 90				
Staphylococcus carnosus subsp. utilis		36.11	0.03	TRE 99				
Staphylococcus auricularis		12.64	0.10					
403011	**Good identification**							**403011**
Staphylococcus capitis subsp. capitis		99.50	0.50	ESL 1				
Staphylococcus carnosus subsp. carnosus		0.50	0.04					
403040	**Low discrimination**							**403040**
Staphylococcus auricularis		79.07	0.37	ESL 1				
Staphylococcus carnosus subsp. utilis		10.76	0.03	TRE 92				
Staphylococcus piscifermentans		10.18	0.28					
403042	**Low discrimination**							**403042**
Staphylococcus piscifermentans		63.48	0.32	URE 90				
Staphylococcus auricularis		36.52	0.17	TRE 99				
403100	**Low discrimination**							**403100**
Staphylococcus piscifermentans		42.99	0.32	URE 90				
Staphylococcus carnosus subsp. utilis		30.28	0.03	TRE 99				
Staphylococcus auricularis		10.60	0.10					
403102	**Good identification**							**403102**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403112	**Good identification**						**403112**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
403140	**Low discrimination**						**403140**
Staphylococcus piscifermentans		63.48	0.32	URE 90			
Staphylococcus auricularis		36.52	0.17	TRE 99			
403142	**Good identification**						**403142**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403152	**Good identification**						**403152**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
403200	**Low discrimination**						**403200**
Staphylococcus auricularis		40.57	0.29	ESL 1			
Staphylococcus piscifermentans		19.81	0.32	TRE 92			
Staphylococcus carnosus subsp. utilis		13.95	0.03				
403201	**Good identification**						**403201**
Staphylococcus capitis subsp. capitis		95.97	0.32	ESL 1			
Staphylococcus auricularis		4.03	0.09	MAN 93			
403202	**Low discrimination**						**403202**
Staphylococcus piscifermentans		86.83	0.36	URE 90			
Staphylococcus auricularis		13.18	0.09	TRE 99			
403210	**Good identification**						**403210**
					OXI	NOV	
Staphylococcus vitulinus		97.61	0.50	ARG 1	99	99	
Staphylococcus capitis subsp. capitis		1.17	0.14		1	1	
Staphylococcus piscifermentans		0.69	0.25		1	1	
403211	**Very good identification**						**403211**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		98.94	0.55	ESL 1	1	1	
Staphylococcus vitulinus		0.84	0.09		99	99	
Staphylococcus sciuri		0.22	0.05		99	99	
403212	**Low discrimination**						**403212**
					OXI	NOV	
Staphylococcus piscifermentans		51.15	0.28	URE 90	1	1	
Staphylococcus vitulinus		48.85	0.09	TRE 99	99	99	
403230	**Very good identification**						**403230**
Staphylococcus vitulinus		100.0	0.59	ARG 1			
403240	**Low discrimination**						**403240**
Staphylococcus auricularis		75.21	0.36	ESL 1			
Staphylococcus piscifermentans		15.73	0.32	TRE 92			
Staphylococcus haemolyticus		9.06	0.19				
403242	**Low discrimination**						**403242**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		63.70	0.36	URE 90			
Staphylococcus auricularis		22.55	0.17	TRE 99			
Staphylococcus haemolyticus		13.75	0.13				
403252	**Low discrimination**						**403252**
Staphylococcus piscifermentans		77.17	0.28	URE 90			
Staphylococcus haemolyticus		22.83	0.09	TRE 99			
403300	**Low discrimination**						**403300**
					OXI	NOV	
Staphylococcus piscifermentans		76.85	0.36	URE 90	1	1	
Staphylococcus auricularis		11.66	0.09	TRE 99	1	1	
Macrococcus carouselicus		11.48	0.02		99	99	
403302	**Good identification**						**403302**
Staphylococcus piscifermentans		100.0	0.39	URE 90			
403310	**Low discrimination**						**403310**
					OXI	NOV	
Staphylococcus piscifermentans		45.92	0.28	URE 90	1	1	
Staphylococcus vitulinus		43.85	0.09	TRE 99	99	99	
Macrococcus carouselicus		10.24	-.02		99	99	
403312	**Good identification**						**403312**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
403340	**Low discrimination**						**403340**
Staphylococcus piscifermentans		55.96	0.36	URE 90			
Staphylococcus haemolyticus		24.22	0.20	TRE 99			
Staphylococcus auricularis		19.82	0.17				
403342	**Low discrimination**						**403342**
Staphylococcus piscifermentans		86.04	0.39	URE 90			
Staphylococcus haemolyticus		13.96	0.14	TRE 99			
403350	**Low discrimination**						**403350**
					OXI	NOV	
Staphylococcus piscifermentans		55.06	0.28	URE 90	1	1	
Staphylococcus haemolyticus		32.66	0.15	TRE 99	1	2	
Staphylococcus sciuri		12.29	0.04		99	99	
403352	**Low discrimination**						**403352**
Staphylococcus piscifermentans		81.81	0.32	URE 90			
Staphylococcus haemolyticus		18.20	0.10	TRE 99			
403400	**Low discrimination**						**403400**
Staphylococcus carnosus subsp. utilis		79.30	0.59	ESL 1			
Staphylococcus piscifermentans		14.30	0.70				
Staphylococcus auricularis		5.53	0.51				
403402	**Good identification**						**403402**
Staphylococcus piscifermentans		93.81	0.73	URE 90			
Staphylococcus carnosus subsp. utilis		3.50	0.18				
Staphylococcus auricularis		2.69	0.32				
403403	**Low discrimination**						**403403**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		71.49	0.32	URE 90			
Staphylococcus auricularis		22.52	0.12	MNS 1			
Staphylococcus carnosus subsp. carnosus		5.99	-.04				
403410	**Low discrimination**						**403410**
					OXI	NOV	
Staphylococcus piscifermentans		79.44	0.62	URE 90	1	1	
Staphylococcus carnosus subsp. utilis		10.39	0.18		1	1	
Macrococcus carouselicus		7.25	0.24		99	99	
403412	**Very good identification**						**403412**
Staphylococcus piscifermentans		100.0	0.66	URE 90			
403422	**Good identification**						**403422**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
403440	**Low discrimination**						**403440**
Staphylococcus piscifermentans		51.06	0.70	URE 90			
Staphylococcus auricularis		46.07	0.59				
Staphylococcus carnosus subsp. utilis		2.86	0.18				
403441	**Acceptable identification**						**403441**
Staphylococcus auricularis		90.85	0.39	ESL 1			
Staphylococcus piscifermentans		9.15	0.28	MNS 10			
403442	**Good identification**						**403442**
Staphylococcus piscifermentans		93.74	0.73	URE 90			
Staphylococcus auricularis		6.27	0.39				
403443	**Low discrimination**						**403443**
Staphylococcus piscifermentans		57.63	0.32	URE 90			
Staphylococcus auricularis		42.37	0.20	MNS 1			
403450	**Good identification**						**403450**
					OXI	NOV	
Staphylococcus piscifermentans		94.19	0.62	URE 90	1	1	
Macrococcus bovicus		3.42	0.20		99	99	
Staphylococcus auricularis		2.00	0.18		1	1	
403452	**Very good identification**						**403452**
Staphylococcus piscifermentans		100.0	0.66	URE 90			
403460	**Low discrimination**						**403460**
Staphylococcus piscifermentans		52.57	0.28	URE 90			
Staphylococcus auricularis		47.43	0.18	XYL 1			
403462	**Good identification**						**403462**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
403500	**Good identification**						**403500**
Staphylococcus piscifermentans		92.30	0.73	URE 90			
Staphylococcus carnosus subsp. utilis		3.45	0.18				
Staphylococcus auricularis		2.64	0.32				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
403501	**Low discrimination**						**403501**
Staphylococcus piscifermentans		76.04	0.32	URE 90			
Staphylococcus auricularis		23.96	0.12	MNS 1			
403502	**Very good identification**						**403502**
Staphylococcus piscifermentans		99.79	0.77	URE 90			
Staphylococcus auricularis		0.21	0.12				
403503	**Good identification**						**403503**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403510	**Very good identification**						**403510**
					OXI	NOV	
Staphylococcus piscifermentans		97.04	0.66	URE 90	1	1	
Macrococcus carouselicus		2.53	0.16		99	99	
Macrococcus bovicus		0.43	0.05		99	99	
403512	**Very good identification**						**403512**
Staphylococcus piscifermentans		100.0	0.69	URE 90			
403513	**Low discrimination**						**403513**
					OXI	NOV	
Staphylococcus piscifermentans		65.07	0.28	URE 90	1	1	
Staphylococcus lentus		26.44	0.12	MNS 1	99	99	
Staphylococcus carnosus subsp. carnosus		8.48	-.04		1	1	
403520	**Good identification**						**403520**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
403522	**Good identification**						**403522**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403532	**Good identification**						**403532**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
403540	**Good identification**						**403540**
Staphylococcus piscifermentans		93.66	0.73	URE 90			
Staphylococcus auricularis		6.26	0.39				
Staphylococcus haemolyticus		0.08	0.02				
403541	**Low discrimination**						**403541**
Staphylococcus piscifermentans		54.43	0.32	URE 90			
Staphylococcus auricularis		40.02	0.20	MNS 1			
Rothia mucilaginosa		5.56	-.09				
403542	**Very good identification**						**403542**
Staphylococcus piscifermentans		99.51	0.77	URE 90			
Staphylococcus auricularis		0.49	0.20				
403543	**Good identification**						**403543**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403550	**Very good identification**						**403550**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		98.97	0.66	URE 90	1	1	
Macrococcus bovicus		1.03	0.12		99	99	
403552	**Very good identification**						**403552**
Staphylococcus piscifermentans		100.0	0.69	URE 90			
403553	**Low discrimination**						**403553**
					OXI	NOV	
Staphylococcus piscifermentans		50.16	0.28	URE 90	1	1	
Staphylococcus lentus		41.38	0.18	MNS 1	99	99	
Aerococcus viridans		8.46	0.18		1	1	
403560	**Good identification**						**403560**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
403562	**Good identification**						**403562**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403572	**Good identification**						**403572**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
403600	**Low discrimination**						**403600**
					OXI	NOV	
Staphylococcus piscifermentans		73.45	0.73	URE 90	1	1	
Staphylococcus auricularis		17.48	0.51		1	1	
Macrococcus carouselicus		6.08	0.34		99	99	
403601	**Low discrimination**						**403601**
					OXI	NOV	
Staphylococcus auricularis		67.34	0.31	ESL 1	1	1	
Staphylococcus piscifermentans		25.73	0.32	MNS 10	1	1	
Staphylococcus sciuri		4.09	0.05		99	99	
403602	**Very good identification**						**403602**
Staphylococcus piscifermentans		98.27	0.77	URE 90			
Staphylococcus auricularis		1.73	0.31				
403603	**Low discrimination**						**403603**
Staphylococcus piscifermentans		75.01	0.36	URE 90			
Staphylococcus auricularis		14.54	0.11	MNS 1			
Staphylococcus hyicus		10.45	0.00				
403610	**Low discrimination**						**403610**
					OXI	NOV	
Staphylococcus piscifermentans		51.22	0.66	URE 90	1	1	
Staphylococcus vitulinus		41.27	0.45		99	99	
Macrococcus carouselicus		6.32	0.30		99	99	
403611	**Low discrimination**						**403611**
					OXI	NOV	
Staphylococcus sciuri		69.62	0.36	ARG 1	99	99	
Staphylococcus capitis subsp. capitis		19.90	0.21		1	1	
Staphylococcus piscifermentans		5.81	0.25		1	1	
403612	**Very good identification**						**403612**
					OXI	NOV	
Staphylococcus piscifermentans		99.30	0.69	URE 90	1	1	
Staphylococcus vitulinus		0.54	0.04		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		0.16	0.02		99	99	
403620	**Low discrimination**						**403620**
					OXI	NOV	
Staphylococcus piscifermentans		46.05	0.32	URE 90	1	1	
Staphylococcus vitulinus		42.99	0.13	XYL 1	99	99	
Staphylococcus auricularis		10.96	0.10		1	1	
403622	**Good identification**						**403622**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403630	**Very good identification**						**403630**
					OXI	NOV	
Staphylococcus vitulinus		99.03	0.54	ARG 1	99	99	
Staphylococcus sciuri		0.51	0.15		99	99	
Staphylococcus piscifermentans		0.46	0.25		1	1	
403631	**Low discrimination**						**403631**
Staphylococcus sciuri		80.59	0.34	ARG 1			
Staphylococcus vitulinus		19.41	0.13				
403640	**Low discrimination**						**403640**
Staphylococcus piscifermentans		60.00	0.73	URE 90			
Staphylococcus auricularis		33.32	0.58				
Staphylococcus haemolyticus		4.64	0.42				
403641	**Low discrimination**						**403641**
					OXI	NOV	
Staphylococcus auricularis		73.53	0.39	ESL 1	1	1	
Staphylococcus piscifermentans		12.04	0.32	MNS 10	1	1	
Staphylococcus sciuri		7.20	0.17		99	99	
403642	**Good identification**						**403642**
Staphylococcus piscifermentans		93.13	0.77	URE 90			
Staphylococcus auricularis		3.83	0.39				
Staphylococcus haemolyticus		2.70	0.37				
403643	**Low discrimination**						**403643**
Staphylococcus piscifermentans		62.27	0.36	URE 90			
Staphylococcus auricularis		28.17	0.19	MNS 1			
Staphylococcus haemolyticus		5.52	0.05				
403650	**Low discrimination**						**403650**
					OXI	NOV	
Staphylococcus piscifermentans		84.07	0.66	URE 90	1	1	
Staphylococcus haemolyticus		8.90	0.37		1	2	
Staphylococcus sciuri		4.74	0.30		99	99	
403651	**Good identification**						**403651**
					OXI	NOV	
Staphylococcus sciuri		96.42	0.48	ARG 1	99	99	
Staphylococcus piscifermentans		2.14	0.25		1	1	
Staphylococcus lentus		0.75	0.07		99	99	
403652	**Very good identification**						**403652**
					OXI	NOV	
Staphylococcus piscifermentans		95.62	0.69	URE 90	1	1	
Staphylococcus haemolyticus		3.80	0.32		1	2	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		0.58	0.13		99	99	
403653	**Low discrimination**						**403653**
					OXI	NOV	
Staphylococcus sciuri		71.06	0.32	ARG 1	99	99	
Staphylococcus piscifermentans		14.51	0.28	LAC 14	1	1	
Staphylococcus lentus		11.98	0.18		99	99	
403660	**Low discrimination**						**403660**
Staphylococcus piscifermentans		64.30	0.32	URE 90			
Staphylococcus auricularis		35.70	0.17	XYL 1			
403662	**Good identification**						**403662**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403670	**Low discrimination**						**403670**
					OXI	NOV	
Staphylococcus sciuri		56.96	0.27	ARG 1	99	99	
Staphylococcus vitulinus		29.50	0.13	MNS 89	99	99	
Staphylococcus piscifermentans		13.54	0.25		1	1	
403671	**Good identification**						**403671**
Staphylococcus sciuri		98.99	0.46	ARG 1			
Staphylococcus lentus		1.02	0.07				
403672	**Low discrimination**						**403672**
					OXI	NOV	
Staphylococcus piscifermentans		68.66	0.28	URE 90	1	1	
Staphylococcus sciuri		31.34	0.11	XYL 1	99	99	
403673	**Low discrimination**						**403673**
					OXI	NOV	
Staphylococcus sciuri		78.80	0.30	ARG 1	99	99	
Staphylococcus lentus		17.60	0.18	LAC 14	99	99	
Aerococcus viridans		3.60	0.18		1	1	
403700	**Very good identification**						**403700**
					OXI	NOV	
Staphylococcus piscifermentans		95.10	0.77	URE 90	1	1	
Macrococcus carouselicus		2.25	0.27		99	99	
Staphylococcus auricularis		1.68	0.31		1	1	
403702	**Very good identification**						**403702**
Staphylococcus piscifermentans		99.87	0.80	URE 90			
Staphylococcus auricularis		0.13	0.11				
403710	**Good identification**						**403710**
					OXI	NOV	
Staphylococcus piscifermentans		94.33	0.69	URE 90	1	1	
Macrococcus carouselicus		3.33	0.23		99	99	
Staphylococcus sciuri		1.83	0.24		99	99	
403711	**Low discrimination**						**403711**
					OXI	NOV	
Staphylococcus sciuri		55.69	0.42	ARG 1	99	99	
Staphylococcus lentus		40.72	0.41		99	99	
Staphylococcus piscifermentans		3.59	0.28		1	1	
403712	**Very good identification**						**403712**
					OXI	NOV	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		99.52	0.73	URE 90	1	1	
Staphylococcus lentus		0.27	0.12		99	99	
Staphylococcus sciuri		0.21	0.07		99	99	
403713	**Acceptable identification**						**403713**
					OXI	NOV	
Staphylococcus lentus		90.90	0.53	ARG 1	99	99	
Staphylococcus sciuri		5.71	0.26		99	99	
Staphylococcus piscifermentans		3.39	0.32		1	1	
403720	**Good identification**						**403720**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
403722	**Good identification**						**403722**
Staphylococcus piscifermentans		100.0	0.39	URE 90			
403731	**Low discrimination**						**403731**
Staphylococcus sciuri		50.78	0.40	ARG 1			
Staphylococcus lentus		49.22	0.41				
403732	**Low discrimination**						**403732**
					OXI	NOV	
Staphylococcus piscifermentans		70.03	0.32	URE 90	1	1	
Staphylococcus lentus		18.97	0.12	XYL 1	99	99	
Staphylococcus sciuri		11.00	0.05		99	99	
403733	**Good identification**						**403733**
Staphylococcus lentus		95.48	0.53	ARG 1			
Staphylococcus sciuri		4.52	0.24				
403740	**Low discrimination**						**403740**
Staphylococcus piscifermentans		89.06	0.77	URE 90			
Staphylococcus haemolyticus		5.17	0.43				
Staphylococcus auricularis		3.66	0.39				
403741	**Low discrimination**						**403741**
					OXI	NOV	
Rothia mucilaginosa		72.45	0.33	VPT 95	1	1	
Staphylococcus piscifermentans		10.75	0.36	ARG 1	1	1	
Staphylococcus sciuri		8.32	0.23		99	99	
403742	**Very good identification**						**403742**
Staphylococcus piscifermentans		97.58	0.80	URE 90			
Staphylococcus haemolyticus		2.13	0.38				
Staphylococcus auricularis		0.30	0.19				
403743	**Low discrimination**						**403743**
					OXI	NOV	
Staphylococcus piscifermentans		69.84	0.39	URE 90	1	1	
Staphylococcus lentus		16.46	0.18	MNS 1	99	99	
Staphylococcus sciuri		5.87	0.07		99	99	
403750	**Low discrimination**						**403750**
					OXI	NOV	
Staphylococcus piscifermentans		86.46	0.69	URE 90	1	1	
Staphylococcus haemolyticus		6.88	0.38		1	2	
Staphylococcus sciuri		6.30	0.36		99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
403751	**Low discrimination**						**403751**
					OXI	NOV	
Staphylococcus sciuri		70.49	0.54	ARG 1	99	99	
Staphylococcus lentus		27.82	0.48		99	99	
Staphylococcus piscifermentans		1.21	0.28		1	1	
403752	**Very good identification**						**403752**
					OXI	NOV	
Staphylococcus piscifermentans		95.85	0.73	URE 90	1	1	
Staphylococcus haemolyticus		2.86	0.33		1	2	
Staphylococcus sciuri		0.76	0.19		99	99	
403753	**Low discrimination**						**403753**
					OXI	NOV	
Staphylococcus lentus		87.84	0.59	ARG 1	99	99	
Staphylococcus sciuri		10.22	0.38		99	99	
Staphylococcus piscifermentans		1.61	0.32		1	1	
403760	**Low discrimination**						**403760**
					OXI	NOV	
Staphylococcus piscifermentans		88.48	0.36	URE 90	1	1	
Staphylococcus sciuri		6.38	0.02	XYL 1	99	99	
Staphylococcus haemolyticus		5.14	0.02		1	2	
403762	**Good identification**						**403762**
Staphylococcus piscifermentans		100.0	0.39	URE 90			
403770	**Low discrimination**						**403770**
					OXI	NOV	
Staphylococcus sciuri		81.54	0.33	ARG 1	99	99	
Staphylococcus piscifermentans		14.98	0.28	MNS 89	1	1	
Staphylococcus lentus		3.49	0.07		99	99	
403771	**Low discrimination**						**403771**
Staphylococcus sciuri		65.66	0.52	ARG 1			
Staphylococcus lentus		34.34	0.48				
403772	**Low discrimination**						**403772**
					OXI	NOV	
Staphylococcus piscifermentans		46.71	0.32	URE 90	1	1	
Staphylococcus sciuri		27.60	0.17	XYL 1	99	99	
Staphylococcus lentus		25.69	0.18		99	99	
403773	**Good identification**						**403773**
					OXI	NOV	
Staphylococcus lentus		91.76	0.59	ARG 1	99	99	
Staphylococcus sciuri		8.06	0.35		99	99	
Aerococcus viridans		0.19	0.18		1	1	
404000	**Low discrimination**						**404000**
					OXI		
Staphylococcus carnosus subsp. utilis		50.07	0.29	PHS 1	1		
Dermacoccus nishinomiyaensis		24.44	0.33		93		
Micrococcus sp.		13.20	0.18		89		
404001	**Low discrimination**						**404001**
Staphylococcus capitis subsp. capitis		60.56	0.30	NIT 89			
Staphylococcus schleiferi subsp. schleiferi		31.56	0.26	PHS 11			
Staphylococcus carnosus subsp. carnosus		6.04	0.11	MAN 93			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
404011	**Very good identification**						**404011**
Staphylococcus capitis subsp. capitis		97.09	0.54	NIT 89			
Staphylococcus carnosus subsp. carnosus		2.92	0.23	PHS 11			
404013	**Low discrimination**						**404013**
Staphylococcus carnosus subsp. carnosus		85.48	0.36	BGA 90			
Staphylococcus capitis subsp. capitis		14.52	0.19	NIT 99			
404040	**Very good identification**						**404040**
					COA	OXI	
Staphylococcus aureus subsp. anaerobius		99.66	0.59	SUC 99	99	1	
Staphylococcus auricularis		0.20	0.17		1	1	
Micrococcus sp.		0.09	-.05		1	89	
404200	**Very good identification**						**404200**
					COA	OXI	
Staphylococcus aureus subsp. anaerobius		99.60	0.59	MLT 99	99	1	
Dermacoccus nishinomiyaensis		0.15	0.08		1	93	
Micrococcus sp.		0.12	-.03		1	89	
404201	**Low discrimination**						**404201**
					COA		
Staphylococcus capitis subsp. capitis		74.17	0.36	NIT 89	1		
Staphylococcus aureus subsp. anaerobius		25.83	0.18	PHS 11	99		
404211	**Very good identification**						**404211**
Staphylococcus capitis subsp. capitis		100.0	0.59	NIT 89			
404240	**Excellent identification**						**404240**
					COA		
Staphylococcus aureus subsp. anaerobius		100.0	1.00		99		
Staphylococcus auricularis		0.00	0.17		1		
404241	**Very good identification**						**404241**
Staphylococcus aureus subsp. anaerobius		100.0	0.59	MNS 1			
404242	**Very good identification**						**404242**
Staphylococcus aureus subsp. anaerobius		100.0	0.59	LAC 1			
404250	**Very good identification**						**404250**
					OXI	COA	
Staphylococcus aureus subsp. anaerobius		99.95	0.59	MAN 1	1	99	
Macrococcus bovicus		0.05	0.02		99	1	
404260	**Very good identification**						**404260**
Staphylococcus aureus subsp. anaerobius		100.0	0.59	XYL 1			
404340	**Very good identification**						**404340**
					COA		
Staphylococcus aureus subsp. anaerobius		99.96	0.59	GAL 1	99		
Staphylococcus epidermidis		0.04	0.03		1		
404400	**Acceptable identification**						**404400**
					NOV		
Staphylococcus carnosus subsp. utilis		94.89	0.44	PHS 1	1		
Staphylococcus auricularis		3.86	0.32		1		
Staphylococcus cohnii subsp. cohnii		1.25	0.25		99		
404401	**Low discrimination**						**404401**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV			
Staphylococcus schleiferi subsp. schleiferi		75.05	0.36	VPT 99	1			
Staphylococcus cohnii subsp. cohnii		10.83	0.26	NIT 90	99			
Staphylococcus carnosus subsp. utilis		7.66	0.03		1			
404410	**Low discrimination**							**404410**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		47.29	0.27	ARG 1	1	99	12	
Staphylococcus carnosus subsp. utilis		28.48	0.03	PHS 9	1	1	10	
Macrococcus bovicus		14.55	0.10		99	99	99	
404411	**Low discrimination**							**404411**
					NOV			
Staphylococcus cohnii subsp. cohnii		39.39	0.28	ARG 1	99			
Staphylococcus capitis subsp. capitis		37.17	0.19	PHS 9	1			
Staphylococcus carnosus subsp. carnosus		23.44	0.16		1			
404413	**Good identification**							**404413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	BGA 90				
404440	**Low discrimination**							**404440**
					COA	NOV		
Staphylococcus auricularis		50.56	0.39	NIT 90	1	1		
Staphylococcus cohnii subsp. cohnii		22.24	0.35	PHS 1	1	99		
Staphylococcus aureus subsp. anaerobius		21.81	0.18		99	1		
404441	**Low discrimination**							**404441**
					NOV			
Staphylococcus cohnii subsp. cohnii		81.09	0.36	ARG 1	99			
Staphylococcus auricularis		18.91	0.20	PHS 9	1			
404450	**Low discrimination**							**404450**
					OXI	YEP		
Staphylococcus kloosii		50.02	0.52	ARG 1	1	99		
Staphylococcus cohnii subsp. cohnii		40.74	0.38		1	12		
Macrococcus bovicus		9.24	0.18		99	99		
404451	**Good identification**							**404451**
Staphylococcus cohnii subsp. cohnii		100.0	0.38	ARG 1				
404452	**Good identification**							**404452**
Staphylococcus kloosii		100.0	0.53	ARG 1				
404470	**Good identification**							**404470**
Staphylococcus kloosii		100.0	0.40	ARG 1				
404472	**Good identification**							**404472**
Staphylococcus kloosii		100.0	0.40	ARG 1				
404501	**Good identification**							**404501**
					NOV			
Staphylococcus schleiferi subsp. schleiferi		98.42	0.36	VPT 99	1			
Staphylococcus cohnii subsp. cohnii		1.58	0.06	NIT 90	99			
404510	**Low discrimination**							**404510**
					OXI	YEP		
Staphylococcus kloosii		62.05	0.32	ARG 1	1	99		
Macrococcus bovicus		20.60	0.03	MLT 95	99	99		
Staphylococcus cohnii subsp. cohnii		17.36	0.08		1	12		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
404512	**Good identification**							**404512**
Staphylococcus kloosii		100.0	0.32	ARG 1				
404550	**Good identification**							**404550**
					OXI	YEP		
Staphylococcus kloosii		91.96	0.58	ARG 1	1	99		
Staphylococcus cohnii subsp. cohnii		4.29	0.18		1	12		
Macrococcus bovicus		3.75	0.10		99	99		
404552	**Very good identification**							**404552**
					NOV	YEP		
Staphylococcus kloosii		96.10	0.59	ARG 1	99	99		
Staphylococcus caprae		3.90	0.02		1	1		
404570	**Good identification**							**404570**
Staphylococcus kloosii		100.0	0.46	ARG 1				
404572	**Good identification**							**404572**
Staphylococcus kloosii		100.0	0.46	ARG 1				
404640	**Very good identification**							**404640**
					COA	PYR		
Staphylococcus aureus subsp. anaerobius		96.61	0.59	TRE 1	99	1		
Staphylococcus auricularis		2.09	0.39		1	40		
Staphylococcus petrasii subsp. pragensis		0.81	0.13		1	99		
404652	**Good identification**							**404652**
Staphylococcus kloosii		100.0	0.32	ARG 1				
404703	**Good identification**							**404703**
					YEP	PYR	COA	
Staphylococcus hyicus		96.00	0.44	NIT 99	1	1	24	
Staphylococcus chromogenes		3.51	0.18		95	45	1	
Staphylococcus intermedius		0.50	0.02		1	90	99	
404750	**Acceptable identification**							**404750**
					OXI			
Staphylococcus kloosii		90.66	0.37	ARG 1	1			
Macrococcus bovicus		9.34	-.02	SUC 9	99			
404752	**Low discrimination**							**404752**
					NOV	YEP		
Staphylococcus kloosii		86.82	0.38	ARG 1	99	99		
Staphylococcus caprae		13.18	-.07	SUC 9	1	1		
404772	**Acceptable identification**							**404772**
Staphylococcus arlettae		90.92	0.35	ARG 1				
Staphylococcus kloosii		9.08	0.26	GLR 92				
404773	**Good identification**							**404773**
Staphylococcus arlettae		100.0	0.37	ARG 1				
405240	**Very good identification**							**405240**
Staphylococcus aureus subsp. anaerobius		100.0	0.59	ESL 1				
405402	**Good identification**							**405402**
Staphylococcus piscifermentans		100.0	0.28	URE 90				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
405442	**Good identification**							**405442**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
405450	**Acceptable identification**							**405450**
					OXI	NOV	YEP	
Staphylococcus kloosii		91.92	0.52	ARG 1	1	99	99	
Staphylococcus piscifermentans		3.85	0.17		1	1	1	
Macrococcus equipersicus		2.34	0.02		99	99	99	
405452	**Acceptable identification**							**405452**
					NOV	YEP		
Staphylococcus kloosii		94.52	0.53	ARG 1	99	99		
Staphylococcus piscifermentans		5.48	0.21		1	1		
405470	**Good identification**							**405470**
Staphylococcus kloosii		100.0	0.40	ARG 1				
405472	**Good identification**							**405472**
Staphylococcus kloosii		100.0	0.40	ARG 1				
405500	**Low discrimination**							**405500**
					OXI	NOV		
Staphylococcus piscifermentans		70.90	0.28	URE 90	1	1		
Macrococcus carouselicus		29.11	0.04	NIT 99	99	99		
405502	**Good identification**							**405502**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
405510	**Low discrimination**							**405510**
					OXI	NOV	YEP	
Staphylococcus kloosii		50.22	0.32	ARG 1	1	99	99	
Staphylococcus piscifermentans		30.88	0.21	MLT 95	1	1	1	
Macrococcus carouselicus		18.91	0.00		99	99	67	
405512	**Low discrimination**							**405512**
					NOV	YEP		
Staphylococcus kloosii		54.02	0.32	ARG 1	99	99		
Staphylococcus piscifermentans		45.98	0.25	MLT 95	1	1		
405540	**Low discrimination**							**405540**
					NOV	YEP		
Staphylococcus piscifermentans		88.20	0.28	URE 90	1	1		
Staphylococcus kloosii		11.80	0.17	NIT 99	99	99		
405542	**Acceptable identification**							**405542**
					NOV	YEP		
Staphylococcus piscifermentans		91.19	0.32	URE 90	1	1		
Staphylococcus kloosii		8.81	0.18	NIT 99	99	99		
405550	**Very good identification**							**405550**
					NOV	YEP		
Staphylococcus kloosii		96.87	0.58	ARG 1	99	99		
Staphylococcus piscifermentans		3.14	0.21		1	1		
405552	**Very good identification**							**405552**
					NOV	YEP		
Staphylococcus kloosii		95.71	0.59	ARG 1	99	99		
Staphylococcus piscifermentans		4.29	0.25		1	1		
405570	**Good identification**							**405570**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.46	ARG 1				
405572	**Good identification**							**405572**
Staphylococcus kloosii		100.0	0.46	ARG 1				
405602	**Good identification**							**405602**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
405642	**Good identification**							**405642**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
405650	**Low discrimination**							**405650**
					OXI	NOV	YEP	
Staphylococcus kloosii		54.95	0.31	ARG 1	1	99	99	
Staphylococcus piscifermentans		34.90	0.21	SUC 9	1	1	1	
Staphylococcus sciuri		10.14	-.01		99	99	57	
405652	**Low discrimination**							**405652**
					NOV	YEP		
Staphylococcus kloosii		53.21	0.32	ARG 1	99	99		
Staphylococcus piscifermentans		46.79	0.25	SUC 9	1	1		
405700	**Low discrimination**							**405700**
					OXI	NOV		
Staphylococcus piscifermentans		64.28	0.32	URE 90	1	1		
Macrococcus carouselicus		35.72	0.10	NIT 99	99	99		
405702	**Good identification**							**405702**
Staphylococcus piscifermentans		100.0	0.36	URE 90				
405712	**Good identification**							**405712**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
405740	**Good identification**							**405740**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
405742	**Good identification**							**405742**
Staphylococcus piscifermentans		100.0	0.36	URE 90				
405750	**Low discrimination**							**405750**
					OXI	NOV	YEP	
Staphylococcus kloosii		59.69	0.37	ARG 1	1	99	99	
Staphylococcus piscifermentans		29.29	0.25	SUC 9	1	1	1	
Staphylococcus sciuri		11.02	0.05		99	99	57	
405752	**Low discrimination**							**405752**
					NOV	YEP		
Staphylococcus kloosii		59.54	0.38	ARG 1	99	99		
Staphylococcus piscifermentans		40.46	0.28	SUC 9	1	1		
405772	**Acceptable identification**							**405772**
Staphylococcus arlettae		90.92	0.35	ARG 1				
Staphylococcus kloosii		9.08	0.26	GLR 92				
405773	**Low discrimination**							**405773**
					OXI	YEP		
Staphylococcus arlettae		88.68	0.37	ARG 1	1	99		
Staphylococcus sciuri		4.52	0.05	GLR 92	99	57		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lentus		3.74	0.05		99	1	
406000	**Low discrimination**						**406000**
					OXI		
Staphylococcus carnosus subsp. utilis		89.05	0.44	PHS 1	1		
Dermacoccus nishinomiyaensis		4.66	0.28		93		
Staphylococcus auricularis		2.83	0.30		1		
406001	**Low discrimination**						**406001**
Staphylococcus carnosus subsp. carnosus		43.17	0.52	BGA 90			
Staphylococcus capitis subsp. capitis		35.38	0.49				
Staphylococcus schleiferi subsp. schleiferi		20.51	0.46				
406003	**Very good identification**						**406003**
Staphylococcus carnosus subsp. carnosus		99.35	0.64	BGA 90			
Staphylococcus capitis subsp. capitis		0.42	0.14				
Staphylococcus schleiferi subsp. schleiferi		0.12	0.04				
406010	**Low discrimination**						**406010**
Staphylococcus capitis subsp. capitis		64.61	0.31	PHS 11			
Staphylococcus carnosus subsp. carnosus		23.74	0.23	MNS 99			
Staphylococcus carnosus subsp. utilis		9.50	0.03				
406011	**Low discrimination**						**406011**
Staphylococcus capitis subsp. capitis		73.11	0.72	PHS 11			
Staphylococcus carnosus subsp. carnosus		26.86	0.64				
Staphylococcus schleiferi subsp. schleiferi		0.03	0.04				
406012	**Good identification**						**406012**
Staphylococcus carnosus subsp. carnosus		98.63	0.36	BGA 90			
Staphylococcus capitis subsp. capitis		1.37	-.04	MNS 99			
406013	**Very good identification**						**406013**
Staphylococcus carnosus subsp. carnosus		98.63	0.77	BGA 90			
Staphylococcus capitis subsp. capitis		1.37	0.37				
406031	**Low discrimination**						**406031**
Staphylococcus capitis subsp. capitis		73.13	0.31	PHS 11			
Staphylococcus carnosus subsp. carnosus		26.87	0.23	XYL 1			
406033	**Good identification**						**406033**
Staphylococcus carnosus subsp. carnosus		98.63	0.36	BGA 90			
Staphylococcus capitis subsp. capitis		1.37	-.04	XYL 1			
406040	**Low discrimination**						**406040**
					COA		
Staphylococcus auricularis		59.27	0.37	PHS 1	1		
Staphylococcus aureus subsp. anaerobius		32.67	0.18	TRE 92	99		
Staphylococcus carnosus subsp. utilis		8.06	0.03		1		
406051	**Low discrimination**						**406051**
Staphylococcus capitis subsp. capitis		73.13	0.31	PHS 11			
Staphylococcus carnosus subsp. carnosus		26.87	0.23	MLT 1			
406053	**Low discrimination**						**406053**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COA		
Staphylococcus carnosus subsp. carnosus		72.34	0.36	BGA 90	1		
Staphylococcus delphini		26.66	0.18	MLT 1	99		
Staphylococcus capitis subsp. capitis		1.01	-.04		1		
406101	**Low discrimination**						**406101**
Staphylococcus schleiferi subsp. schleiferi		64.36	0.46	VPT 99			
Staphylococcus carnosus subsp. carnosus		33.87	0.40				
Staphylococcus capitis subsp. capitis		1.12	0.08				
406103	**Acceptable identification**						**406103**
					COA		
Staphylococcus carnosus subsp. carnosus		93.50	0.52	BGA 90	1		
Staphylococcus hyicus		5.16	0.22		24		
Staphylococcus schleiferi subsp. coagulans		0.70	0.12		99		
406111	**Low discrimination**						**406111**
Staphylococcus carnosus subsp. carnosus		89.70	0.52	BGA 90			
Staphylococcus capitis subsp. capitis		9.87	0.31				
Staphylococcus schleiferi subsp. schleiferi		0.43	0.04				
406113	**Very good identification**						**406113**
					COA		
Staphylococcus carnosus subsp. carnosus		99.68	0.64	BGA 90	1		
Staphylococcus schleiferi subsp. coagulans		0.17	0.12		99		
Staphylococcus felis		0.09	-.04		1		
406201	**Very good identification**						**406201**
Staphylococcus capitis subsp. capitis		95.97	0.55	PHS 11			
Staphylococcus hyicus		2.60	0.19	MAN 93			
Staphylococcus carnosus subsp. carnosus		0.64	0.11				
406203	**Low discrimination**						**406203**
Staphylococcus hyicus		83.53	0.41	GAL 92			
Staphylococcus carnosus subsp. carnosus		7.11	0.23	TRE 92			
Staphylococcus capitis subsp. capitis		5.46	0.20				
406210	**Good identification**						**406210**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		97.51	0.37	PHS 11	1	1	
Staphylococcus vitulinus		2.49	0.00	MNS 99	99	99	
406211	**Very good identification**						**406211**
Staphylococcus capitis subsp. capitis		99.80	0.78	PHS 11			
Staphylococcus carnosus subsp. carnosus		0.20	0.23				
406213	**Low discrimination**						**406213**
					COA		
Staphylococcus capitis subsp. capitis		63.35	0.43	PHS 11	1		
Staphylococcus carnosus subsp. carnosus		24.81	0.36	LAC 2	1		
Staphylococcus delphini		9.14	0.18		99		
406231	**Good identification**						**406231**
Staphylococcus capitis subsp. capitis		100.0	0.37	PHS 11			
406240	**Very good identification**						**406240**
					COA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. anaerobius		97.90	0.59	NIT 1	99		
Staphylococcus auricularis		1.66	0.36		1		
Staphylococcus epidermidis		0.24	0.18		1		
406243	**Low discrimination**						**406243**
					COA	YEP	
Staphylococcus epidermidis		52.33	0.34	VPT 96	1	1	
Staphylococcus delphini		32.82	0.18	URE 92	99	1	
Staphylococcus chromogenes		12.21	0.17		1	95	
406251	**Low discrimination**						**406251**
					COA		
Staphylococcus capitis subsp. capitis		77.42	0.37	PHS 11	1		
Staphylococcus delphini		22.58	0.18	MLT 1	99		
406253	**Very good identification**						**406253**
					COA	YEP	
Staphylococcus delphini		99.60	0.59	URE 99	99	1	
Staphylococcus capitis subsp. ureolyticus		0.22	0.03		1	77	
Staphylococcus chromogenes		0.11	0.06		1	95	
406301	**Acceptable identification**						**406301**
					YEP		
Staphylococcus hyicus		91.37	0.41	TRE 92	1		
Staphylococcus chromogenes		4.26	0.17	LAC 92	95		
Staphylococcus capitis subsp. capitis		2.96	0.14		1		
406303	**Very good identification**						**406303**
					YEP		
Staphylococcus hyicus		96.12	0.63	TRE 92	1		
Staphylococcus chromogenes		3.51	0.37		95		
Staphylococcus carnosus subsp. carnosus		0.18	0.11		1		
406311	**Acceptable identification**						**406311**
					YEP		
Staphylococcus capitis subsp. capitis		90.62	0.37	PHS 11	1		
Staphylococcus carnosus subsp. carnosus		4.48	0.11	GAL 1	1		
Staphylococcus chromogenes		2.77	0.06		95		
406340	**Low discrimination**						**406340**
					COA	PYR	
Staphylococcus epidermidis		49.03	0.34	VPT 96	1	1	
Staphylococcus aureus subsp. anaerobius		35.09	0.18	URE 92	99	1	
Staphylococcus haemolyticus		7.98	0.20		1	98	
406341	**Low discrimination**						**406341**
					YEP		
Staphylococcus epidermidis		86.31	0.40	VPT 96	1		
Staphylococcus chromogenes		11.25	0.17	URE 92	95		
Staphylococcus hyicus		2.44	0.00		1		
406342	**Good identification**						**406342**
					PYR		
Staphylococcus epidermidis		95.32	0.44	VPT 96	1		
Staphylococcus haemolyticus		2.76	0.14	URE 92	98		
Staphylococcus piscifermentans		1.26	0.16		15		
406343	**Low discrimination**						**406343**
					COA	YEP	
Staphylococcus epidermidis		62.78	0.50	VPT 96	1	1	
Staphylococcus chromogenes		23.26	0.37	URE 92	1	95	
Staphylococcus delphini		6.95	0.18		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
406353	**Very good identification**						**406353**
					COA	YEP	
Staphylococcus delphini		98.12	0.59	URE 99	99	1	
Staphylococcus chromogenes		0.94	0.26		1	95	
Staphylococcus caprae		0.40	0.11		1	1	
406400	**Good identification**						**406400**
Staphylococcus carnosus subsp. utilis		92.14	0.59	PHS 1			
Staphylococcus auricularis		6.42	0.51				
Staphylococcus piscifermentans		1.23	0.46				
406401	**Low discrimination**						**406401**
Staphylococcus schleiferi subsp. schleiferi		70.37	0.55	VPT 99			
Staphylococcus carnosus subsp. carnosus		21.16	0.44				
Staphylococcus carnosus subsp. utilis		4.19	0.18				
406402	**Low discrimination**						**406402**
Staphylococcus piscifermentans		50.15	0.50	URE 90			
Staphylococcus carnosus subsp. utilis		25.29	0.18	ESL 90			
Staphylococcus auricularis		19.39	0.32				
406403	**Good identification**						**406403**
Staphylococcus carnosus subsp. carnosus		94.69	0.57	BGA 90			
Staphylococcus hyicus		3.05	0.22				
Staphylococcus felis		0.92	0.09				
406411	**Low discrimination**						**406411**
Staphylococcus carnosus subsp. carnosus		87.77	0.57	BGA 90			
Staphylococcus capitis subsp. capitis		11.38	0.37				
Staphylococcus schleiferi subsp. schleiferi		0.74	0.14				
406412	**Low discrimination**						**406412**
Staphylococcus piscifermentans		51.02	0.42	URE 90			
Staphylococcus carnosus subsp. carnosus		48.98	0.28	ESL 90			
406413	**Very good identification**						**406413**
Staphylococcus carnosus subsp. carnosus		98.28	0.69	BGA 90			
Staphylococcus felis		1.60	0.26				
Staphylococcus capitis subsp. capitis		0.07	0.03				
406433	**Good identification**						**406433**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	BGA 90			
406440	**Low discrimination**						**406440**
Staphylococcus auricularis		87.12	0.59	PHS 1			
Staphylococcus piscifermentans		7.15	0.46				
Staphylococcus carnosus subsp. utilis		5.41	0.18				
406441	**Low discrimination**						**406441**
					NOV		
Staphylococcus auricularis		86.27	0.39	PHS 1	1		
Staphylococcus schleiferi subsp. schleiferi		8.18	0.14	MNS 10	1		
Staphylococcus cohnii subsp. cohnii		3.10	0.13		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
406442	**Low discrimination**					**406442**
Staphylococcus piscifermentans		52.57	0.50	URE 90		
Staphylococcus auricularis		47.43	0.39	ESL 90		
406450	**Low discrimination**					**406450**
					NOV	
Staphylococcus piscifermentans		68.27	0.39	URE 90	1	
Staphylococcus auricularis		19.60	0.18	ESL 90	1	
Staphylococcus cohnii subsp. cohnii		9.09	0.14		99	
406452	**Good identification**					**406452**
Staphylococcus piscifermentans		100.0	0.42	URE 90		
406453	**Acceptable identification**					**406453**
Staphylococcus carnosus subsp. carnosus		92.10	0.28	BGA 90		
Staphylococcus caprae		7.90	-.02	MLT 1		
406500	**Low discrimination**					**406500**
Staphylococcus piscifermentans		50.59	0.50	URE 90		
Staphylococcus carnosus subsp. utilis		25.52	0.18	ESL 90		
Staphylococcus auricularis		19.56	0.32			
406501	**Low discrimination**					**406501**
Staphylococcus schleiferi subsp. schleiferi		89.23	0.55	VPT 99		
Staphylococcus carnosus subsp. carnosus		6.71	0.32			
Staphylococcus hyicus		3.45	0.22			
406502	**Good identification**					**406502**
Staphylococcus piscifermentans		93.36	0.54	URE 90		
Staphylococcus auricularis		2.67	0.12	ESL 90		
Staphylococcus hyicus		2.37	0.03			
406503	**Low discrimination**					**406503**
Staphylococcus hyicus		54.73	0.44	SUC 99		
Staphylococcus carnosus subsp. carnosus		36.98	0.44			
Staphylococcus felis		4.83	0.20			
406510	**Acceptable identification**					**406510**
Staphylococcus piscifermentans		94.34	0.42	URE 90		
Staphylococcus carnosus subsp. carnosus		5.66	0.03	ESL 90		
406511	**Good identification**					**406511**
Staphylococcus carnosus subsp. carnosus		95.45	0.44	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		3.21	0.14			
Staphylococcus felis		0.84	-.04			
406512	**Low discrimination**					**406512**
Staphylococcus piscifermentans		83.69	0.46	URE 90		
Staphylococcus carnosus subsp. carnosus		13.39	0.16	ESL 90		
Staphylococcus felis		2.93	-.04			
406513	**Low discrimination**					**406513**

Code Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus	79.54	0.57	BGA 90		
Staphylococcus felis	17.38	0.37			
Staphylococcus condimenti	2.21	0.14			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
406540	**Low discrimination**						**406540**
Staphylococcus piscifermentans		52.24	0.50	URE 90			
Staphylococcus auricularis		47.14	0.39	ESL 90			
Staphylococcus haemolyticus		0.62	0.02				
406542	**Good identification**						**406542**
Staphylococcus piscifermentans		92.29	0.54	URE 90			
Staphylococcus auricularis		6.17	0.20	ESL 90			
Staphylococcus caprae		1.54	-.05				
406550	**Acceptable identification**						**406550**
					NOV	YEP	
Staphylococcus piscifermentans		92.01	0.42	URE 90	1	1	
Staphylococcus caprae		5.38	-.05	ESL 90	1	1	
Staphylococcus kloosii		2.61	0.17		99	99	
406552	**Low discrimination**						**406552**
					NOV	YEP	
Staphylococcus piscifermentans		56.76	0.46	URE 90	1	1	
Staphylococcus caprae		42.07	0.22	ESL 90	1	1	
Staphylococcus kloosii		1.16	0.18		99	99	
406553	**Acceptable identification**						**406553**
					COA		
Staphylococcus caprae		93.40	0.39	VPT 99	1		
Staphylococcus carnosus subsp. carnosus		2.75	0.16	URE 90	1		
Staphylococcus aureus subsp. aureus		2.21	0.01		99		
406600	**Low discrimination**						**406600**
Staphylococcus auricularis		59.69	0.51	PHS 1			
Staphylococcus piscifermentans		18.58	0.50				
Staphylococcus muscae		11.83	0.18				
406601	**Low discrimination**						**406601**
Staphylococcus hyicus		77.46	0.41	GAL 92			
Staphylococcus auricularis		9.80	0.31	LAC 92			
Staphylococcus capitis subsp. capitis		5.07	0.20				
406602	**Low discrimination**						**406602**
Staphylococcus piscifermentans		67.11	0.54	URE 90			
Staphylococcus auricularis		15.97	0.31	ESL 90			
Staphylococcus hyicus		14.66	0.22				
406603	**Good identification**						**406603**
					YEP		
Staphylococcus hyicus		94.22	0.63	GAL 92	1		
Staphylococcus chromogenes		4.40	0.39		95		
Staphylococcus felis		0.59	0.16		1		
406610	**Low discrimination**						**406610**
Staphylococcus piscifermentans		85.17	0.42	URE 90			
Staphylococcus auricularis		6.45	0.10	ESL 90			
Staphylococcus capitis subsp. capitis		4.92	0.02				
406611	**Acceptable identification**						**406611**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus capitis subsp. capitis		92.30	0.43	PHS 11	1		
Staphylococcus carnosus subsp. carnosus		3.87	0.16	TRE 2	1		
Staphylococcus chromogenes		1.79	0.08		95		
406612	**Low discrimination**						**406612**
Staphylococcus piscifermentans		80.58	0.46	URE 90			
Staphylococcus simulans		17.72	0.21	ESL 90			
Staphylococcus felis		1.71	-.09				
406613	**Low discrimination**						**406613**
					YEP		
Staphylococcus felis		44.41	0.33	URE 99	1		
Staphylococcus simulans		15.57	0.32		1		
Staphylococcus chromogenes		14.05	0.28		95		
406620	**Very good identification**						**406620**
Staphylococcus muscae		99.95	0.59	ARG 1			
Staphylococcus auricularis		0.05	0.10				
406640	**Low discrimination**						**406640**
Staphylococcus auricularis		72.02	0.58	PHS 1			
Staphylococcus haemolyticus		10.02	0.42				
Staphylococcus piscifermentans		9.61	0.50				
406641	**Low discrimination**						**406641**
					YEP		
Staphylococcus auricularis		78.44	0.39	PHS 1	1		
Staphylococcus chromogenes		15.84	0.20	MNS 10	95		
Staphylococcus haemolyticus		3.04	0.11		20		
406642	**Low discrimination**						**406642**
Staphylococcus piscifermentans		49.22	0.54	URE 90			
Staphylococcus auricularis		27.33	0.39	ESL 90			
Staphylococcus haemolyticus		19.25	0.37				
406643	**Low discrimination**						**406643**
					YEP		
Staphylococcus chromogenes		74.92	0.39	URE 99	95		
Staphylococcus hyicus		16.22	0.22	GAL 90	1		
Staphylococcus auricularis		4.58	0.19		1		
406650	**Low discrimination**						**406650**
Staphylococcus haemolyticus		54.85	0.37	VPT 86			
Staphylococcus piscifermentans		38.37	0.42	PHS 1			
Staphylococcus auricularis		6.78	0.17				
406652	**Low discrimination**						**406652**
Staphylococcus piscifermentans		61.17	0.46	URE 90			
Staphylococcus haemolyticus		32.79	0.32	ESL 90			
Staphylococcus simulans		6.04	0.14				
406653	**Low discrimination**						**406653**
					COA	YEP	
Staphylococcus chromogenes		35.96	0.28	URE 99	1	95	
Staphylococcus delphini		29.80	0.18	GAL 90	99	1	
Staphylococcus simulans		17.91	0.25		1	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
406700	**Low discrimination**							**406700**
	Staphylococcus piscifermentans	67.82	0.54	URE 90				
	Staphylococcus auricularis	16.14	0.31	ESL 90				
	Staphylococcus hyicus	14.81	0.22					
406701	**Very good identification**							**406701**
					YEP			
	Staphylococcus hyicus	95.03	0.63	LAC 92	1			
	Staphylococcus chromogenes	4.44	0.39		95			
	Staphylococcus schleiferi subsp. schleiferi	0.25	0.14		1			
406702	**Low discrimination**							**406702**
					YEP			
	Staphylococcus hyicus	60.82	0.44	MNS 99	1			
	Staphylococcus piscifermentans	36.32	0.57		1			
	Staphylococcus chromogenes	2.22	0.18		95			
406703	**Very good identification**							**406703**
					YEP			
	Staphylococcus hyicus	96.23	0.85		1			
	Staphylococcus chromogenes	3.51	0.59		95			
	Staphylococcus felis	0.17	0.26		1			
406710	**Good identification**							**406710**
	Staphylococcus piscifermentans	100.0	0.46	URE 90				
406711	**Low discrimination**							**406711**
					YEP	PYR	COA	
	Staphylococcus chromogenes	43.22	0.28	URE 99	95	45	1	
	Staphylococcus hyicus	33.17	0.22	LAC 90	1	1	24	
	Staphylococcus intermedius	11.51	0.18		1	90	99	
406712	**Low discrimination**							**406712**
					YEP			
	Staphylococcus piscifermentans	87.74	0.50	URE 90	1			
	Staphylococcus felis	4.15	0.02	ESL 90	1			
	Staphylococcus chromogenes	3.53	0.07		95			
406713	**Low discrimination**							**406713**
					YEP			
	Staphylococcus felis	33.48	0.43	URE 99	1			
	Staphylococcus chromogenes	28.48	0.48		95			
	Staphylococcus hyicus	27.93	0.44		1			
406723	**Good identification**							**406723**
					YEP			
	Staphylococcus hyicus	96.48	0.44	XYL 1	1			
	Staphylococcus chromogenes	3.52	0.18		95			
406740	**Low discrimination**							**406740**
	Staphylococcus piscifermentans	42.32	0.54	URE 90				
	Staphylococcus haemolyticus	33.18	0.43	ESL 90				
	Staphylococcus auricularis	23.50	0.39					
406741	**Low discrimination**							**406741**
					YEP			
	Staphylococcus chromogenes	73.89	0.39	URE 99	95			
	Staphylococcus hyicus	16.00	0.22	LAC 90	1			
	Staphylococcus auricularis	4.52	0.19		1			

Code Taxon		ID score	T index	tests against	Differentiation tests				Code
406742	**Low discrimination**								**406742**
					YEP				
Staphylococcus piscifermentans		69.86	0.57	URE 90	1				
Staphylococcus haemolyticus		20.54	0.38	ESL 90	20				
Staphylococcus chromogenes		4.27	0.18		95				
406743	**Low discrimination**								**406743**
					YEP	PYR	COA		
Staphylococcus chromogenes		75.66	0.59	URE 99	95	45	1		
Staphylococcus hyicus		20.93	0.44		1	1	24		
Staphylococcus intermedius		2.15	0.29		1	90	99		
406750	**Low discrimination**								**406750**
					NOV				
Staphylococcus haemolyticus		51.19	0.38	VPT 86	2				
Staphylococcus piscifermentans		47.63	0.46	PHS 1	1				
Staphylococcus saprophyticus subsp. bovis		1.18	0.02		99				
406751	**Low discrimination**								**406751**
					OXI	COA	NOV	YEP	
Staphylococcus chromogenes		49.78	0.28	URE 99	1	1	1	95	
Staphylococcus intermedius		19.88	0.21	LAC 90	1	99	1	1	
Staphylococcus sciuri		10.26	0.18		99	1	99	57	
406752	**Low discrimination**								**406752**
Staphylococcus piscifermentans		58.42	0.50	URE 90					
Staphylococcus haemolyticus		23.54	0.33	ESL 90					
Staphylococcus caprae		10.68	0.13						
406753	**Low discrimination**								**406753**
					COA	YEP			
Staphylococcus aureus subsp. aureus		55.08	0.42	VPT 87	99	84			
Staphylococcus chromogenes		25.83	0.48	URE 99	1	95			
Staphylococcus caprae		8.69	0.30		1	1			
407001	**Low discrimination**								**407001**
Staphylococcus carnosus subsp. carnosus		89.47	0.32	BGA 90					
Staphylococcus capitis subsp. capitis		6.67	0.08	ESL 10					
Staphylococcus schleiferi subsp. schleiferi		3.86	0.04						
407002	**Low discrimination**								**407002**
Staphylococcus piscifermentans		77.32	0.28	URE 90					
Staphylococcus carnosus subsp. carnosus		22.68	0.04	TRE 99					
407003	**Good identification**								**407003**
Staphylococcus carnosus subsp. carnosus		100.0	0.45	BGA 90					
407011	**Low discrimination**								**407011**
Staphylococcus carnosus subsp. carnosus		80.16	0.45	BGA 90					
Staphylococcus capitis subsp. capitis		19.84	0.31	ESL 10					
407013	**Very good identification**								**407013**
Staphylococcus carnosus subsp. carnosus		99.87	0.57	BGA 90					
Staphylococcus capitis subsp. capitis		0.13	-.04	ESL 10					
407042	**Good identification**								**407042**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		100.0	0.28	URE 90			
407100	**Good identification**						**407100**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
407102	**Good identification**						**407102**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407103	**Acceptable identification**						**407103**
					COA		
Staphylococcus carnosus subsp. carnosus		93.89	0.32	BGA 90	1		
Staphylococcus schleiferi subsp. coagulans		6.11	0.12	ESL 10	99		
407111	**Good identification**						**407111**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	BGA 90			
407113	**Good identification**						**407113**
					COA		
Staphylococcus carnosus subsp. carnosus		98.52	0.45	BGA 90	1		
Staphylococcus schleiferi subsp. coagulans		1.48	0.11	ESL 10	99		
407140	**Good identification**						**407140**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
407142	**Good identification**						**407142**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407200	**Good identification**						**407200**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
407202	**Good identification**						**407202**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407211	**Good identification**						**407211**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		95.02	0.37	ESL 1	1	1	
Staphylococcus sciuri		2.89	0.10	PHS 11	99	99	
Staphylococcus carnosus subsp. carnosus		2.09	0.04		1	1	
407242	**Good identification**						**407242**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407300	**Good identification**						**407300**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407302	**Good identification**						**407302**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
407312	**Good identification**						**407312**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
407340	**Good identification**						**407340**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407342	**Acceptable identification**						**407342**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		92.16	0.36	URE 90		
Staphylococcus epidermidis		7.84	0.03	TRE 99		
407351	**Good identification**					**407351**
Staphylococcus sciuri		100.0	0.28	ARG 1		
407352	**Good identification**					**407352**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
407400	**Good identification**					**407400**
Staphylococcus piscifermentans		91.75	0.66	URE 90		
Staphylococcus carnosus subsp. utilis		7.71	0.18			
Staphylococcus auricularis		0.54	0.10			
407402	**Very good identification**					**407402**
Staphylococcus piscifermentans		99.87	0.70	URE 90		
Staphylococcus carnosus subsp. carnosus		0.13	-.04			
407403	**Acceptable identification**					**407403**
Staphylococcus carnosus subsp. carnosus		92.56	0.37	BGA 90		
Staphylococcus piscifermentans		7.44	0.28	ESL 10		
407410	**Very good identification**					**407410**
Staphylococcus piscifermentans		99.56	0.58	URE 90		
Staphylococcus carnosus subsp. carnosus		0.44	-.04			
407411	**Good identification**					**407411**
Staphylococcus carnosus subsp. carnosus		95.66	0.37	BGA 90		
Staphylococcus piscifermentans		2.20	0.17	ESL 10		
Staphylococcus capitis subsp. capitis		1.13	-.04			
407412	**Very good identification**					**407412**
Staphylococcus piscifermentans		98.83	0.62	URE 90		
Staphylococcus carnosus subsp. carnosus		1.17	0.08			
407413	**Good identification**					**407413**
Staphylococcus carnosus subsp. carnosus		99.15	0.50	BGA 90		
Staphylococcus piscifermentans		0.85	0.21	ESL 10		
407422	**Good identification**					**407422**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
407440	**Very good identification**					**407440**
Staphylococcus piscifermentans		98.65	0.66	URE 90		
Staphylococcus auricularis		1.35	0.18			
407442	**Very good identification**					**407442**
Staphylococcus piscifermentans		100.0	0.70	URE 90		
407443	**Low discrimination**					**407443**
Staphylococcus piscifermentans		88.83	0.28	URE 90		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		11.17	-.04	MNS 1		
407450	**Very good identification**					**407450**
Staphylococcus piscifermentans		100.0	0.58	URE 90		
407452	**Very good identification**					**407452**
Staphylococcus piscifermentans		100.0	0.62	URE 90		
407462	**Good identification**					**407462**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
407500	**Very good identification**					**407500**
Staphylococcus piscifermentans		100.0	0.70	URE 90		
407501	**Low discrimination**					**407501**
Staphylococcus piscifermentans		36.78	0.28	URE 90		
Staphylococcus schleiferi subsp. schleiferi		34.60	0.14	MNS 1		
Staphylococcus carnosus subsp. carnosus		28.61	0.12			
407502	**Very good identification**					**407502**
Staphylococcus piscifermentans		100.0	0.73	URE 90		
407510	**Very good identification**					**407510**
Staphylococcus piscifermentans		100.0	0.62	URE 90		
407512	**Very good identification**					**407512**
Staphylococcus piscifermentans		99.80	0.66	URE 90		
Staphylococcus carnosus subsp. carnosus		0.20	-.04			
407513	**Acceptable identification**					**407513**
Staphylococcus carnosus subsp. carnosus		93.33	0.37	BGA 90		
Staphylococcus piscifermentans		4.82	0.25	ESL 10		
Staphylococcus felis		1.85	-.04			
407520	**Good identification**					**407520**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
407522	**Good identification**					**407522**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
407540	**Very good identification**					**407540**
Staphylococcus piscifermentans		100.0	0.70	URE 90		
407541	**Good identification**					**407541**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
407542	**Very good identification**					**407542**
Staphylococcus piscifermentans		100.0	0.73	URE 90		
407543	**Good identification**					**407543**
Staphylococcus piscifermentans		100.0	0.32	URE 90		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
407550	**Very good identification**						**407550**
					NOV	YEP	
Staphylococcus piscifermentans		99.69	0.62	URE 90	1	1	
Staphylococcus kloosii		0.31	0.17		99	99	
407552	**Very good identification**						**407552**
					NOV	YEP	
Staphylococcus piscifermentans		99.77	0.66	URE 90	1	1	
Staphylococcus kloosii		0.23	0.18		99	99	
407560	**Good identification**						**407560**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
407562	**Good identification**						**407562**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407600	**Very good identification**						**407600**
Staphylococcus piscifermentans		99.64	0.70	URE 90			
Staphylococcus auricularis		0.36	0.10				
407601	**Low discrimination**						**407601**
					OXI	NOV	
Staphylococcus piscifermentans		58.16	0.28	URE 90	1	1	
Staphylococcus sciuri		23.61	0.10	MNS 1	99	99	
Staphylococcus hyicus		18.23	0.00		1	1	
407602	**Very good identification**						**407602**
Staphylococcus piscifermentans		100.0	0.73	URE 90			
407610	**Very good identification**						**407610**
					OXI	NOV	
Staphylococcus piscifermentans		95.20	0.62	URE 90	1	1	
Staphylococcus sciuri		3.64	0.23		99	99	
Staphylococcus vitulinus		1.16	0.04		99	99	
407611	**Acceptable identification**						**407611**
					OXI	NOV	
Staphylococcus sciuri		94.08	0.41	ARG 1	99	99	
Staphylococcus piscifermentans		3.07	0.21		1	1	
Staphylococcus capitis subsp. capitis		1.95	0.02		1	1	
407612	**Very good identification**						**407612**
					OXI	NOV	
Staphylococcus piscifermentans		99.59	0.66	URE 90	1	1	
Staphylococcus sciuri		0.41	0.06		99	99	
407622	**Good identification**						**407622**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
407631	**Good identification**						**407631**
Staphylococcus sciuri		100.0	0.39	ARG 1			
407640	**Very good identification**						**407640**
					OXI	NOV	
Staphylococcus piscifermentans		98.98	0.70	URE 90	1	1	
Staphylococcus auricularis		0.83	0.17		1	1	
Staphylococcus sciuri		0.19	0.03		99	99	
407642	**Very good identification**						**407642**

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		100.0	0.73	URE 90				
407643	**Low discrimination**							**407643**
					OXI	YEP	NOV	
Staphylococcus piscifermentans		78.26	0.32	URE 90	1	1	1	
Staphylococcus sciuri		12.97	0.06	MNS 1	99	57	99	
Staphylococcus chromogenes		8.77	-.02		1	95	1	
407650	**Low discrimination**							**407650**
					OXI	NOV		
Staphylococcus piscifermentans		87.42	0.62	URE 90	1	1		
Staphylococcus sciuri		12.58	0.34		99	99		
407651	**Very good identification**							**407651**
					OXI	NOV		
Staphylococcus sciuri		99.14	0.53	ARG 1	99	99		
Staphylococcus piscifermentans		0.86	0.21		1	1		
407652	**Very good identification**							**407652**
					OXI	NOV		
Staphylococcus piscifermentans		98.46	0.66	URE 90	1	1		
Staphylococcus sciuri		1.54	0.18		99	99		
407653	**Acceptable identification**							**407653**
					OXI	NOV		
Staphylococcus sciuri		90.52	0.37	ARG 1	99	99		
Staphylococcus piscifermentans		7.24	0.25	LAC 14	1	1		
Staphylococcus lentus		2.24	0.05		99	99		
407660	**Low discrimination**							**407660**
					OXI	NOV		
Staphylococcus piscifermentans		87.54	0.28	URE 90	1	1		
Staphylococcus sciuri		12.47	0.01	XYL 1	99	99		
407662	**Good identification**							**407662**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
407670	**Acceptable identification**							**407670**
					OXI	NOV		
Staphylococcus sciuri		91.49	0.32	ARG 1	99	99		
Staphylococcus piscifermentans		8.52	0.21	MNS 89	1	1		
407671	**Good identification**							**407671**
Staphylococcus sciuri		100.0	0.51	ARG 1				
407673	**Good identification**							**407673**
Staphylococcus sciuri		96.83	0.34	ARG 1				
Staphylococcus lentus		3.18	0.05	LAC 14				
407700	**Very good identification**							**407700**
Staphylococcus piscifermentans		100.0	0.73	URE 90				
407702	**Very good identification**							**407702**
Staphylococcus piscifermentans		99.81	0.77	URE 90				
Staphylococcus hyicus		0.19	0.03					
407703	**Acceptable identification**							**407703**
					YEP			
Staphylococcus hyicus		91.42	0.44	ESL 1	1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		4.96	0.36		1			
Staphylococcus chromogenes		3.34	0.18		95			
407710	**Very good identification**							**407710**
					OXI	NOV		
Staphylococcus piscifermentans		95.29	0.66	URE 90	1	1		
Staphylococcus sciuri		4.72	0.28		99	99		
407711	**Low discrimination**							**407711**
					OXI	NOV		
Staphylococcus sciuri		88.29	0.47	ARG 1	99	99		
Staphylococcus lentus		9.48	0.29		99	99		
Staphylococcus piscifermentans		2.23	0.25		1	1		
407712	**Very good identification**							**407712**
					OXI	NOV		
Staphylococcus piscifermentans		99.47	0.69	URE 90	1	1		
Staphylococcus sciuri		0.53	0.12		99	99		
407713	**Low discrimination**							**407713**
					OXI	NOV		
Staphylococcus lentus		59.56	0.40	ARG 1	99	99		
Staphylococcus sciuri		25.47	0.31		99	99		
Staphylococcus piscifermentans		5.92	0.28		1	1		
407720	**Good identification**							**407720**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
407722	**Good identification**							**407722**
Staphylococcus piscifermentans		100.0	0.36	URE 90				
407731	**Low discrimination**							**407731**
Staphylococcus sciuri		87.54	0.45	ARG 1				
Staphylococcus lentus		12.46	0.29					
407732	**Low discrimination**							**407732**
					OXI	NOV		
Staphylococcus piscifermentans		71.37	0.28	URE 90	1	1		
Staphylococcus sciuri		28.63	0.10	XYL 1	99	99		
407733	**Low discrimination**							**407733**
Staphylococcus lentus		75.61	0.40	ARG 1				
Staphylococcus sciuri		24.40	0.28					
407740	**Very good identification**							**407740**
					OXI	NOV		
Staphylococcus piscifermentans		99.67	0.73	URE 90	1	1		
Staphylococcus sciuri		0.25	0.09		99	99		
Staphylococcus haemolyticus		0.09	0.02		1	2		
407741	**Low discrimination**							**407741**
					OXI	YEP	NOV	
Staphylococcus sciuri		61.95	0.28	ARG 1	99	57	99	
Staphylococcus piscifermentans		31.34	0.32	MAN 97	1	1	1	
Staphylococcus chromogenes		3.51	-.02		1	95	1	
407742	**Very good identification**							**407742**
Staphylococcus piscifermentans		100.0	0.77	URE 90				
407743	**Low discrimination**							**407743**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					OXI	YEP	NOV	
Staphylococcus piscifermentans		46.26	0.36	URE 90	1	1	1	
Staphylococcus chromogenes		31.12	0.18	MNS 1	1	95	1	
Staphylococcus sciuri		9.92	0.12		99	57	99	
407750	**Low discrimination**							**407750**
					OXI	NOV		
Staphylococcus piscifermentans		84.31	0.66	URE 90	1	1		
Staphylococcus sciuri		15.69	0.40		99	99		
407751	**Good identification**							**407751**
					OXI	NOV		
Staphylococcus sciuri		93.93	0.59	ARG 1	99	99		
Staphylococcus lentus		5.44	0.35		99	99		
Staphylococcus piscifermentans		0.63	0.25		1	1		
407752	**Very good identification**							**407752**
					OXI	NOV		
Staphylococcus piscifermentans		97.82	0.69	URE 90	1	1		
Staphylococcus sciuri		1.98	0.24		99	99		
Staphylococcus lentus		0.20	0.05		99	99		
407753	**Low discrimination**							**407753**
					OXI	NOV		
Staphylococcus lentus		52.17	0.47	ARG 1	99	99		
Staphylococcus sciuri		41.35	0.43		99	99		
Staphylococcus piscifermentans		2.56	0.28		1	1		
407760	**Low discrimination**							**407760**
					OXI	NOV		
Staphylococcus piscifermentans		84.44	0.32	URE 90	1	1		
Staphylococcus sciuri		15.56	0.07	XYL 1	99	99		
407762	**Good identification**							**407762**
Staphylococcus piscifermentans		100.0	0.36	URE 90				
407770	**Acceptable identification**							**407770**
					OXI	NOV		
Staphylococcus sciuri		93.29	0.38	ARG 1	99	99		
Staphylococcus piscifermentans		6.71	0.25	MNS 89	1	1		
407771	**Low discrimination**							**407771**
					OXI			
Staphylococcus sciuri		89.22	0.56	ARG 1	99			
Staphylococcus lentus		6.85	0.35		99			
Staphylococcus gallinarum		3.93	0.23		1			
407773	**Low discrimination**							**407773**
					OXI			
Staphylococcus lentus		49.96	0.47	ARG 1	99			
Staphylococcus sciuri		29.87	0.40		99			
Staphylococcus gallinarum		19.79	0.31		1			
410000	**Low discrimination**							**410000**
					OXI			
Staphylococcus carnosus subsp. utilis		72.05	0.29	ORN 1	1			
Micrococcus sp.		19.00	0.18		89			
Dermacoccus nishinomiyaensis		6.75	0.18		93			
410011	**Good identification**							**410011**
Staphylococcus capitis subsp. capitis		100.0	0.31	ORN 1				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
410211	**Good identification**					**410211**
Staphylococcus capitis subsp. capitis		100.0	0.37	ORN 1		
410400	**Good identification**					**410400**
					NOV	
Staphylococcus carnosus subsp. utilis		95.97	0.44	ORN 1	1	
Staphylococcus auricularis		3.90	0.32		1	
Staphylococcus cohnii subsp. cohnii		0.13	0.05		99	
410440	**Low discrimination**					**410440**
					NOV	
Staphylococcus auricularis		86.85	0.39	ORN 1	1	
Staphylococcus carnosus subsp. utilis		9.25	0.03	NIT 90	1	
Staphylococcus cohnii subsp. cohnii		3.90	0.15		99	
410600	**Low discrimination**					**410600**
Staphylococcus auricularis		78.80	0.31	ORN 1		
Staphylococcus carnosus subsp. utilis		21.21	0.03	NIT 90		
410640	**Low discrimination**					**410640**
Staphylococcus auricularis		61.66	0.39	ORN 1		
Staphylococcus petrasii subsp. pragensis		23.81	0.13	NIT 90		
Staphylococcus petrasii subsp. jettensis		13.49	0.10			
412000	**Acceptable identification**					**412000**
Staphylococcus carnosus subsp. utilis		93.78	0.44	ORN 1		
Staphylococcus auricularis		2.98	0.30			
Kocuria rosea		1.93	0.18			
412001	**Low discrimination**					**412001**
Staphylococcus capitis subsp. capitis		75.42	0.27	ORN 1		
Staphylococcus carnosus subsp. utilis		18.21	0.03	MAN 93		
Staphylococcus auricularis		6.37	0.10			
412011	**Good identification**					**412011**
Staphylococcus capitis subsp. capitis		100.0	0.50	ORN 1		
412040	**Low discrimination**					**412040**
Staphylococcus auricularis		88.03	0.37	ORN 1		
Staphylococcus carnosus subsp. utilis		11.98	0.03	TRE 92		
412200	**Low discrimination**					**412200**
Staphylococcus auricularis		70.35	0.29	ORN 1		
Staphylococcus carnosus subsp. utilis		24.19	0.03	TRE 92		
Kocuria rosea		5.46	-.02			
412201	**Good identification**					**412201**
Staphylococcus capitis subsp. capitis		95.97	0.32	ORN 1		
Staphylococcus auricularis		4.03	0.09	MAN 93		
412211	**Very good identification**					**412211**
Staphylococcus capitis subsp. capitis		100.0	0.55	ORN 1		
412240	**Low discrimination**					**412240**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		89.25	0.36	ORN 1		
Staphylococcus haemolyticus		10.75	0.19	TRE 92		
412400	**Good identification**					**412400**
Staphylococcus carnosus subsp. utilis		93.48	0.59	ORN 1		
Staphylococcus auricularis		6.52	0.51			
412440	**Good identification**					**412440**
Staphylococcus auricularis		94.15	0.59	ORN 1		
Staphylococcus carnosus subsp. utilis		5.85	0.18			
412441	**Good identification**					**412441**
Staphylococcus auricularis		100.0	0.39	ORN 1		
412442	**Good identification**					**412442**
Staphylococcus auricularis		98.35	0.39	ORN 1		
Staphylococcus piscifermentans		1.65	0.12	LAC 10		
412540	**Good identification**					**412540**
Staphylococcus auricularis		97.09	0.39	ORN 1		
Staphylococcus piscifermentans		1.63	0.12	GAL 10		
Staphylococcus haemolyticus		1.28	0.02			
412600	**Low discrimination**					**412600**
Staphylococcus auricularis		86.08	0.51	ORN 1		
Staphylococcus carnosus subsp. utilis		13.51	0.18			
Staphylococcus piscifermentans		0.41	0.12			
412601	**Good identification**					**412601**
Staphylococcus auricularis		95.95	0.31	ORN 1		
Staphylococcus capitis subsp. capitis		4.05	-.03	MNS 10		
412602	**Acceptable identification**					**412602**
Staphylococcus auricularis		94.01	0.31	ORN 1		
Staphylococcus piscifermentans		5.99	0.16	LAC 10		
412640	**Low discrimination**					**412640**
Staphylococcus auricularis		83.15	0.58	ORN 1		
Staphylococcus haemolyticus		11.57	0.42			
Staphylococcus petrasii subsp. jettensis		4.72	0.18			
412641	**Acceptable identification**					**412641**
Staphylococcus auricularis		90.25	0.39	ORN 1		
Staphylococcus lugdunensis		6.26	0.09	MNS 10		
Staphylococcus haemolyticus		3.50	0.11			
412642	**Low discrimination**					**412642**
Staphylococcus auricularis		54.89	0.39	ORN 1		
Staphylococcus haemolyticus		38.67	0.37	LAC 10		
Staphylococcus petrasii subsp. jettensis		4.95	0.02			
412643	**Low discrimination**					**412643**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus lugdunensis		82.44	0.29	VPT 99		
Staphylococcus auricularis		14.68	0.19	ARG 5		
Staphylococcus haemolyticus		2.88	0.05			
412650	**Low discrimination**					**412650**
Staphylococcus haemolyticus		89.00	0.37	VPT 86		
Staphylococcus auricularis		11.00	0.17	ORN 1		
412652	**Good identification**					**412652**
Staphylococcus haemolyticus		100.0	0.32	VPT 86		
412700	**Acceptable identification**					**412700**
Staphylococcus auricularis		91.16	0.31	ORN 1		
Staphylococcus piscifermentans		5.80	0.16	GAL 10		
Staphylococcus haemolyticus		3.03	0.02			
412740	**Low discrimination**					**412740**
Staphylococcus haemolyticus		57.89	0.43	VPT 86		
Staphylococcus auricularis		40.99	0.39	ORN 1		
Staphylococcus piscifermentans		1.12	0.16			
412742	**Low discrimination**					**412742**
Staphylococcus haemolyticus		83.94	0.38	VPT 86		
Staphylococcus auricularis		11.74	0.19	ORN 1		
Staphylococcus piscifermentans		4.33	0.20			
412743	**Good identification**					**412743**
Staphylococcus lugdunensis		96.21	0.29	VPT 99		
Staphylococcus haemolyticus		3.79	0.07	ARG 5		
412750	**Good identification**					**412750**
Staphylococcus haemolyticus		100.0	0.38	VPT 86		
412752	**Good identification**					**412752**
Staphylococcus haemolyticus		96.38	0.33	VPT 86		
Staphylococcus piscifermentans		3.62	0.12	ORN 1		
413402	**Good identification**					**413402**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
413440	**Low discrimination**					**413440**
Staphylococcus piscifermentans		52.57	0.28	URE 90		
Staphylococcus auricularis		47.43	0.18	ORN 1		
413442	**Good identification**					**413442**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
413500	**Good identification**					**413500**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
413502	**Good identification**					**413502**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.36	URE 90		
413512	**Good identification**					**413512**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
413540	**Good identification**					**413540**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
413542	**Good identification**					**413542**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
413552	**Good identification**					**413552**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
413600	**Low discrimination**					**413600**
Staphylococcus piscifermentans		80.78	0.32	URE 90		
Staphylococcus auricularis		19.22	0.10	ORN 1		
413602	**Good identification**					**413602**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
413612	**Good identification**					**413612**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
413640	**Low discrimination**					**413640**
Staphylococcus piscifermentans		64.30	0.32	URE 90		
Staphylococcus auricularis		35.70	0.17	ORN 1		
413642	**Good identification**					**413642**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
413652	**Good identification**					**413652**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
413700	**Good identification**					**413700**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
413702	**Good identification**					**413702**
Staphylococcus piscifermentans		100.0	0.39	URE 90		
413710	**Good identification**					**413710**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
413712	**Good identification**					**413712**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
413740	**Acceptable identification**					**413740**
Staphylococcus piscifermentans		94.51	0.36	URE 90		
Staphylococcus haemolyticus		5.49	0.02	ORN 1		
413742	**Good identification**					**413742**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.39	URE 90		
413750	**Good identification**					**413750**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
413752	**Good identification**					**413752**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
414240	**Very good identification**					**414240**
Staphylococcus aureus subsp. anaerobius		100.0	0.59	ORN 1		
416011	**Low discrimination**					**416011**
Staphylococcus capitis subsp. capitis		73.13	0.31	ORN 1		
Staphylococcus carnosus subsp. carnosus		26.87	0.23	PHS 11		
416013	**Good identification**					**416013**
Staphylococcus carnosus subsp. carnosus		98.63	0.36	ORN 1		
Staphylococcus capitis subsp. capitis		1.37	-.04	BGA 90		
416211	**Good identification**					**416211**
Staphylococcus capitis subsp. capitis		100.0	0.37	ORN 1		
416343	**Good identification**					**416343**
					YEP	
Staphylococcus epidermidis		97.33	0.32	VPT 96	1	
Staphylococcus chromogenes		2.67	-.04	URE 92	95	
416413	**Good identification**					**416413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ORN 1		
416703	**Good identification**					**416703**
					YEP	
Staphylococcus hyicus		96.48	0.44	ORN 1	1	
Staphylococcus chromogenes		3.52	0.18		95	
417402	**Good identification**					**417402**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417442	**Good identification**					**417442**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417500	**Good identification**					**417500**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417502	**Good identification**					**417502**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
417540	**Good identification**					**417540**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417542	**Good identification**					**417542**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
417600	**Good identification**					**417600**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417602	**Good identification**					**417602**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
417640	**Good identification**					**417640**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417642	**Good identification**					**417642**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
417700	**Good identification**					**417700**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
417702	**Good identification**					**417702**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
417712	**Good identification**					**417712**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
417740	**Good identification**					**417740**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
417742	**Good identification**					**417742**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
417752	**Good identification**					**417752**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
420000	**Low discrimination**					**420000**
					OXI	
Staphylococcus carnosus subsp. utilis		58.38	0.29	BGA 1	1	
Staphylococcus auricularis		17.46	0.35		1	
Micrococcus sp.		15.39	0.18		89	
420011	**Good identification**					**420011**
Staphylococcus capitis subsp. capitis		96.77	0.31	BGA 1		
Staphylococcus carnosus subsp. carnosus		3.23	0.02	NIT 89		
420040	**Acceptable identification**					**420040**
					OXI	
Staphylococcus auricularis		94.62	0.42	BGA 14	1	
Micrococcus sp.		2.69	-.05	NIT 90	89	
Staphylococcus carnosus subsp. utilis		1.37	-.12	TRE 92	1	
420200	**Low discrimination**					**420200**
					OXI	
Staphylococcus auricularis		88.42	0.34	BGA 14	1	
Micrococcus sp.		8.35	-.03	NIT 90	89	
Staphylococcus carnosus subsp. utilis		3.23	-.12	TRE 92	1	
420211	**Good identification**					**420211**
Staphylococcus capitis subsp. capitis		100.0	0.37	BGA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
420240	**Low discrimination**						**420240**
					COA	PYR	
Staphylococcus auricularis		73.21	0.42	BGA 14	1	40	
Staphylococcus aureus subsp. anaerobius		24.41	0.18	NIT 90	99	1	
Staphylococcus haemolyticus		1.23	0.06	TRE 92	1	98	
420400	**Low discrimination**						**420400**
					NOV		
Staphylococcus carnosus subsp. utilis		60.36	0.44	BGA 1	1		
Staphylococcus auricularis		39.56	0.57		1		
Staphylococcus cohnii subsp. cohnii		0.08	0.05		99		
420401	**Low discrimination**						**420401**
Staphylococcus auricularis		84.79	0.37	BGA 14			
Staphylococcus carnosus subsp. utilis		11.76	0.03	NIT 90			
Staphylococcus schleiferi subsp. schleiferi		1.75	-.02	MNS 10			
420402	**Low discrimination**						**420402**
Staphylococcus auricularis		87.82	0.37	BGA 14			
Staphylococcus carnosus subsp. utilis		12.18	0.03	NIT 90			
420440	**Very good identification**						**420440**
					NOV		
Staphylococcus auricularis		98.95	0.64	BGA 14	1		
Staphylococcus carnosus subsp. utilis		0.65	0.03	NIT 90	1		
Staphylococcus cohnii subsp. cohnii		0.28	0.15		99		
420441	**Good identification**						**420441**
					NOV		
Staphylococcus auricularis		97.35	0.45	BGA 14	1		
Staphylococcus cohnii subsp. cohnii		2.65	0.16	NIT 90	99		
420442	**Good identification**						**420442**
Staphylococcus auricularis		98.84	0.45	BGA 14			
Aerococcus viridans		1.16	0.20	NIT 90			
420452	**Low discrimination**						**420452**
					NOV	PYR	
Aerococcus viridans		44.72	0.30	ARG 1	1	95	
Staphylococcus saprophyticus subsp. saprophyticus		42.48	0.02	ESL 90	99	7	
Staphylococcus auricularis		12.81	0.03		1	40	
420453	**Good identification**						**420453**
Aerococcus viridans		100.0	0.38	ARG 1			
420463	**Good identification**						**420463**
Aerococcus viridans		100.0	0.28	ARG 1			
420472	**Good identification**						**420472**
Aerococcus viridans		100.0	0.30	ARG 1			
420473	**Good identification**						**420473**
Aerococcus viridans		100.0	0.38	ARG 1			
420500	**Low discrimination**						**420500**
Staphylococcus auricularis		87.82	0.37	BGA 14			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. utilis		12.18	0.03	NIT 90			
420540	**Good identification**						**420540**
Staphylococcus auricularis		100.0	0.45	BGA 14			
420543	**Low discrimination**						**420543**
Aerococcus viridans		68.97	0.28	ARG 1			
Staphylococcus auricularis		31.03	0.05	ESL 90			
420552	**Low discrimination**						**420552**
					NOV	YEP	
Aerococcus viridans		81.22	0.30	ARG 1	1	1	
Staphylococcus kloosii		18.78	0.18	ESL 90	99	99	
420553	**Good identification**						**420553**
Aerococcus viridans		100.0	0.38	ARG 1			
420563	**Good identification**						**420563**
Aerococcus viridans		100.0	0.28	ARG 1			
420572	**Good identification**						**420572**
Aerococcus viridans		100.0	0.30	ARG 1			
420573	**Good identification**						**420573**
Aerococcus viridans		100.0	0.38	ARG 1			
420600	**Very good identification**						**420600**
Staphylococcus auricularis		98.05	0.56	BGA 14			
Staphylococcus carnosus subsp. utilis		1.64	0.03	NIT 90			
Staphylococcus warneri		0.31	0.13				
420601	**Acceptable identification**						**420601**
					OXI		
Staphylococcus auricularis		90.10	0.36	BGA 14	1		
Kocuria kristinae		9.90	0.10	NIT 90	99		
420602	**Acceptable identification**						**420602**
Staphylococcus auricularis		94.15	0.36	BGA 14			
Staphylococcus simulans		5.86	0.10	NIT 90			
420613	**Acceptable identification**						**420613**
Staphylococcus simulans		94.12	0.33	URE 98			
Aerococcus viridans		5.88	0.25	NIT 96			
420640	**Good identification**						**420640**
					PYR		
Staphylococcus auricularis		92.35	0.63	BGA 14	40		
Staphylococcus warneri		2.39	0.39	NIT 90	8		
Staphylococcus petrasii subsp. pragensis		2.21	0.13		99		
420641	**Low discrimination**						**420641**
					OXI		
Staphylococcus auricularis		70.23	0.44	BGA 14	1		
Kocuria kristinae		29.77	0.30	NIT 90	99		
420642	**Low discrimination**						**420642**
					PYR		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		80.81	0.44	BGA 14	40		
Staphylococcus haemolyticus		7.95	0.24	NIT 90	98		
Staphylococcus warneri		6.27	0.29	LAC 10	8		
420650	**Low discrimination**						**420650**
					PYR		
Staphylococcus warneri		37.78	0.36	URE 98	8		
Staphylococcus haemolyticus		23.02	0.25		98		
Staphylococcus auricularis		20.38	0.22		40		
420663	**Good identification**						**420663**
Aerococcus viridans		100.0	0.28	ARG 1			
420672	**Low discrimination**						**420672**
					NOV	YEP	
Aerococcus viridans		59.99	0.30	ARG 1	1	1	
Staphylococcus arlettae		40.01	0.15	ESL 90	99	99	
420673	**Low discrimination**						**420673**
					NOV	YEP	
Aerococcus viridans		72.52	0.38	ARG 1	1	1	
Staphylococcus arlettae		27.48	0.18	ESL 90	99	99	
420700	**Good identification**						**420700**
Staphylococcus auricularis		100.0	0.36	BGA 14			
420740	**Low discrimination**						**420740**
					PYR		
Staphylococcus auricularis		78.70	0.44	BGA 14	40		
Staphylococcus haemolyticus		15.52	0.30	NIT 90	98		
Staphylococcus warneri		5.78	0.29	GAL 10	8		
420763	**Good identification**						**420763**
Aerococcus viridans		100.0	0.28	ARG 1			
420772	**Low discrimination**						**420772**
					NOV	YEP	
Staphylococcus arlettae		79.08	0.31	ARG 1	99	99	
Aerococcus viridans		20.93	0.30	GLR 92	1	1	
420773	**Low discrimination**						**420773**
					NOV	YEP	
Staphylococcus arlettae		68.22	0.34	ARG 1	99	99	
Aerococcus viridans		31.78	0.38	GLR 92	1	1	
421053	**Good identification**						**421053**
Aerococcus viridans		100.0	0.31	ARG 1			
421073	**Good identification**						**421073**
Aerococcus viridans		100.0	0.31	ARG 1			
421153	**Good identification**						**421153**
Aerococcus viridans		100.0	0.31	ARG 1			
421173	**Good identification**						**421173**
Aerococcus viridans		100.0	0.31	ARG 1			
421253	**Good identification**						**421253**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.31	ARG 1			
421273	**Good identification**						**421273**
Aerococcus viridans		100.0	0.31	ARG 1			
421353	**Good identification**						**421353**
Aerococcus viridans		100.0	0.31	ARG 1			
421373	**Good identification**						**421373**
Aerococcus viridans		100.0	0.31	ARG 1			
421402	**Good identification**						**421402**
Aerococcus viridans		100.0	0.27	ARG 1			
421403	**Good identification**						**421403**
Aerococcus viridans		100.0	0.35	ARG 1			
421412	**Good identification**						**421412**
Aerococcus viridans		100.0	0.37	ARG 1			
421413	**Good identification**						**421413**
Aerococcus viridans		100.0	0.45	ARG 1			
421422	**Good identification**						**421422**
Aerococcus viridans		100.0	0.27	ARG 1			
421423	**Good identification**						**421423**
Aerococcus viridans		100.0	0.35	ARG 1			
421432	**Good identification**						**421432**
Aerococcus viridans		100.0	0.37	ARG 1			
421433	**Good identification**						**421433**
Aerococcus viridans		100.0	0.45	ARG 1			
421441	**Low discrimination**						**421441**
Aerococcus viridans		73.09	0.28	ARG 1			
Staphylococcus auricularis		26.91	0.03	LAC 90			
421442	**Acceptable identification**						**421442**
Aerococcus viridans		91.29	0.40	ARG 1			
Staphylococcus auricularis		8.71	0.03				
421443	**Good identification**						**421443**
Aerococcus viridans		100.0	0.47	ARG 1			
421450	**Low discrimination**						**421450**
					OXI	NOV	
Aerococcus viridans		66.64	0.30	ARG 1	1	1	
Macrococcus equipersicus		18.48	0.02	LAC 90	99	99	
Macrococcus bovicus		14.88	-.02		99	99	
421451	**Good identification**						**421451**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Aerococcus viridans		100.0	0.38	ARG 1		
421452	**Good identification**					**421452**
Aerococcus viridans		100.0	0.50	ARG 1		
421453	**Very good identification**					**421453**
Aerococcus viridans		100.0	0.57	ARG 1		
421461	**Good identification**					**421461**
Aerococcus viridans		100.0	0.28	ARG 1		
421462	**Good identification**					**421462**
Aerococcus viridans		100.0	0.40	ARG 1		
421463	**Good identification**					**421463**
Aerococcus viridans		100.0	0.47	ARG 1		
421470	**Good identification**					**421470**
Aerococcus viridans		100.0	0.30	ARG 1		
421471	**Good identification**					**421471**
Aerococcus viridans		100.0	0.38	ARG 1		
421472	**Good identification**					**421472**
Aerococcus viridans		100.0	0.50	ARG 1		
421473	**Very good identification**					**421473**
Aerococcus viridans		100.0	0.57	ARG 1		
421502	**Good identification**					**421502**
Aerococcus viridans		100.0	0.27	ARG 1		
421503	**Good identification**					**421503**
Aerococcus viridans		100.0	0.35	ARG 1		
421512	**Good identification**					**421512**
Aerococcus viridans		100.0	0.37	ARG 1		
421513	**Good identification**					**421513**
Aerococcus viridans		100.0	0.45	ARG 1		
421522	**Good identification**					**421522**
Aerococcus viridans		100.0	0.27	ARG 1		
421523	**Good identification**					**421523**
Aerococcus viridans		100.0	0.35	ARG 1		
421532	**Good identification**					**421532**
Aerococcus viridans		100.0	0.37	ARG 1		
421533	**Good identification**					**421533**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.45	ARG 1			
421541	**Good identification**						**421541**
Aerococcus viridans		100.0	0.28	ARG 1			
421542	**Good identification**						**421542**
Aerococcus viridans		100.0	0.40	ARG 1			
421543	**Good identification**						**421543**
Aerococcus viridans		100.0	0.47	ARG 1			
421550	**Low discrimination**						**421550**
					NOV	YEP	
Aerococcus viridans		82.41	0.30	ARG 1	1	1	
Staphylococcus kloosii		17.59	0.17	LAC 90	99	99	
421551	**Good identification**						**421551**
Aerococcus viridans		100.0	0.38	ARG 1			
421552	**Good identification**						**421552**
					NOV	YEP	
Aerococcus viridans		97.50	0.50	ARG 1	1	1	
Staphylococcus kloosii		2.50	0.18		99	99	
421553	**Very good identification**						**421553**
Aerococcus viridans		100.0	0.57	ARG 1			
421561	**Good identification**						**421561**
Aerococcus viridans		100.0	0.28	ARG 1			
421562	**Good identification**						**421562**
Aerococcus viridans		100.0	0.40	ARG 1			
421563	**Good identification**						**421563**
Aerococcus viridans		100.0	0.47	ARG 1			
421570	**Good identification**						**421570**
Aerococcus viridans		100.0	0.30	ARG 1			
421571	**Good identification**						**421571**
Aerococcus viridans		100.0	0.38	ARG 1			
421572	**Good identification**						**421572**
Aerococcus viridans		100.0	0.50	ARG 1			
421573	**Very good identification**						**421573**
Aerococcus viridans		100.0	0.57	ARG 1			
421602	**Good identification**						**421602**
Aerococcus viridans		100.0	0.27	ARG 1			
421603	**Good identification**						**421603**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Aerococcus viridans		100.0	0.35	ARG 1		
421612	**Good identification**					**421612**
Aerococcus viridans		100.0	0.37	ARG 1		
421613	**Good identification**					**421613**
Aerococcus viridans		100.0	0.45	ARG 1		
421622	**Good identification**					**421622**
Aerococcus viridans		100.0	0.27	ARG 1		
421623	**Good identification**					**421623**
Aerococcus viridans		100.0	0.35	ARG 1		
421632	**Good identification**					**421632**
Aerococcus viridans		100.0	0.37	ARG 1		
421633	**Good identification**					**421633**
Aerococcus viridans		100.0	0.45	ARG 1		
421642	**Acceptable identification**					**421642**
Aerococcus viridans		91.90	0.40	ARG 1		
Staphylococcus auricularis		8.10	0.03			
421643	**Acceptable identification**					**421643**
					OXI	
Aerococcus viridans		90.99	0.47	ARG 1	1	
Kocuria kristinae		9.01	0.05		99	
421650	**Good identification**					**421650**
Aerococcus viridans		100.0	0.30	ARG 1		
421651	**Good identification**					**421651**
Aerococcus viridans		100.0	0.38	ARG 1		
421652	**Good identification**					**421652**
Aerococcus viridans		100.0	0.50	ARG 1		
421653	**Very good identification**					**421653**
Aerococcus viridans		100.0	0.57	ARG 1		
421661	**Good identification**					**421661**
Aerococcus viridans		100.0	0.28	ARG 1		
421662	**Good identification**					**421662**
Aerococcus viridans		100.0	0.40	ARG 1		
421663	**Good identification**					**421663**
Aerococcus viridans		100.0	0.47	ARG 1		
421670	**Good identification**					**421670**
Aerococcus viridans		100.0	0.30	ARG 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
421671	**Good identification**						**421671**
Aerococcus viridans		100.0	0.38	ARG 1			
421672	**Acceptable identification**						**421672**
					NOV	YEP	
Aerococcus viridans		93.10	0.50	ARG 1	1	1	
Staphylococcus arlettae		6.90	0.15		99	99	
421673	**Very good identification**						**421673**
					NOV	YEP	
Aerococcus viridans		95.96	0.57	ARG 1	1	1	
Staphylococcus arlettae		4.04	0.18		99	99	
421702	**Good identification**						**421702**
Aerococcus viridans		100.0	0.27	ARG 1			
421703	**Good identification**						**421703**
Aerococcus viridans		100.0	0.35	ARG 1			
421712	**Good identification**						**421712**
Aerococcus viridans		100.0	0.37	ARG 1			
421713	**Low discrimination**						**421713**
					OXI	NOV	
Aerococcus viridans		89.36	0.45	ARG 1	1	1	
Staphylococcus lentus		10.64	0.12		99	99	
421722	**Good identification**						**421722**
Aerococcus viridans		100.0	0.27	ARG 1			
421723	**Good identification**						**421723**
Aerococcus viridans		100.0	0.35	ARG 1			
421732	**Good identification**						**421732**
Aerococcus viridans		100.0	0.37	ARG 1			
421733	**Low discrimination**						**421733**
					OXI	NOV	
Aerococcus viridans		89.36	0.45	ARG 1	1	1	
Staphylococcus lentus		10.64	0.12		99	99	
421742	**Good identification**						**421742**
Aerococcus viridans		100.0	0.40	ARG 1			
421743	**Good identification**						**421743**
					OXI		
Aerococcus viridans		96.80	0.47	ARG 1	1		
Kocuria kristinae		3.20	-.05		99		
421750	**Good identification**						**421750**
Aerococcus viridans		100.0	0.30	ARG 1			
421751	**Low discrimination**						**421751**
					OXI	NOV	
Aerococcus viridans		86.70	0.38	ARG 1	1	1	
Staphylococcus lentus		13.30	0.07	LAC 90	99	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
421752	**Good identification**							**421752**
Aerococcus viridans		100.0	0.50	ARG 1				
421753	**Good identification**							**421753**
					OXI	NOV		
Aerococcus viridans		94.30	0.57	ARG 1	1	1		
Staphylococcus lentus		5.70	0.18		99	99		
421761	**Good identification**							**421761**
Aerococcus viridans		100.0	0.28	ARG 1				
421762	**Good identification**							**421762**
Aerococcus viridans		100.0	0.40	ARG 1				
421763	**Good identification**							**421763**
Aerococcus viridans		100.0	0.47	ARG 1				
421770	**Good identification**							**421770**
Aerococcus viridans		100.0	0.30	ARG 1				
421771	**Low discrimination**							**421771**
					OXI	NOV		
Aerococcus viridans		86.70	0.38	ARG 1	1	1		
Staphylococcus lentus		13.30	0.07	LAC 90	99	99		
421772	**Low discrimination**							**421772**
					NOV	YEP		
Aerococcus viridans		70.43	0.50	ARG 1	1	1		
Staphylococcus arlettae		29.57	0.31		99	99		
421773	**Low discrimination**							**421773**
					OXI	NOV	YEP	
Aerococcus viridans		76.98	0.57	ARG 1	1	1	1	
Staphylococcus arlettae		18.36	0.34		1	99	99	
Staphylococcus lentus		4.65	0.18		99	99	1	
422000	**Low discrimination**							**422000**
Staphylococcus carnosus subsp. utilis		62.99	0.44	BGA 1				
Staphylococcus auricularis		32.31	0.54					
Kocuria varians		2.61	0.33					
422001	**Low discrimination**							**422001**
Staphylococcus auricularis		32.79	0.35	BGA 14				
Staphylococcus carnosus subsp. carnosus		32.67	0.31	TRE 92				
Staphylococcus capitis subsp. capitis		24.07	0.27	MNS 10				
422002	**Low discrimination**							**422002**
Staphylococcus auricularis		70.29	0.35	BGA 14				
Kocuria varians		14.42	0.22	TRE 92				
Staphylococcus carnosus subsp. utilis		12.46	0.03	LAC 10				
422003	**Good identification**							**422003**
Staphylococcus carnosus subsp. carnosus		97.29	0.43	PHS 99				
Staphylococcus auricularis		2.71	0.15					
422011	**Low discrimination**							**422011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. capitis		70.99	0.50	BGA 1		
Staphylococcus carnosus subsp. carnosus		29.01	0.43			
422013	**Very good identification**					**422013**
Staphylococcus carnosus subsp. carnosus		98.45	0.55	PHS 99		
Staphylococcus capitis subsp. capitis		1.23	0.15			
Staphylococcus simulans		0.32	0.07			
422040	**Very good identification**					**422040**
Staphylococcus auricularis		98.25	0.62	BGA 14		
Staphylococcus carnosus subsp. utilis		0.83	0.03	TRE 92		
Kocuria varians		0.75	0.20			
422041	**Good identification**					**422041**
Staphylococcus auricularis		100.0	0.42	BGA 14		
422042	**Good identification**					**422042**
Staphylococcus auricularis		98.11	0.42	BGA 14		
Kocuria varians		1.89	0.08	TRE 92		
422100	**Low discrimination**					**422100**
Kocuria varians		47.39	0.37	ARG 1		
Staphylococcus auricularis		43.45	0.35			
Staphylococcus carnosus subsp. utilis		7.70	0.03			
422103	**Good identification**					**422103**
Staphylococcus carnosus subsp. carnosus		97.39	0.31	PHS 99		
Kocuria varians		2.61	0.04			
422111	**Acceptable identification**					**422111**
Staphylococcus carnosus subsp. carnosus		91.00	0.31	PHS 99		
Staphylococcus capitis subsp. capitis		9.00	0.09			
422113	**Good identification**					**422113**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	PHS 99		
422120	**Good identification**					**422120**
Kocuria varians		100.0	0.27	ARG 1		
422140	**Low discrimination**					**422140**
					PYR	
Staphylococcus auricularis		89.28	0.42	BGA 14	40	
Kocuria varians		9.16	0.23	GAL 10	10	
Staphylococcus haemolyticus		1.56	0.07	TRE 92	98	
422200	**Very good identification**					**422200**
Staphylococcus auricularis		95.93	0.54	BGA 14		
Staphylococcus carnosus subsp. utilis		2.05	0.03	TRE 92		
Kocuria varians		1.15	0.15			
422201	**Low discrimination**					**422201**
Staphylococcus capitis subsp. capitis		59.63	0.32	BGA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		40.37	0.34	MAN 93		
422202	**Low discrimination**					**422202**
Staphylococcus auricularis		88.44	0.34	BGA 14		
Staphylococcus simulans		8.88	0.12	TRE 92		
Kocuria varians		2.68	0.04	LAC 10		
422211	**Very good identification**					**422211**
Staphylococcus capitis subsp. capitis		99.78	0.55	BGA 1		
Staphylococcus carnosus subsp. carnosus		0.22	0.02			
422213	**Low discrimination**					**422213**
Staphylococcus simulans		66.97	0.35	URE 98		
Staphylococcus capitis subsp. capitis		19.95	0.21	TRE 95		
Staphylococcus carnosus subsp. carnosus		8.69	0.14			
422240	**Low discrimination**					**422240**
					PYR	
Staphylococcus auricularis		84.31	0.61	BGA 14	40	
Staphylococcus haemolyticus		15.60	0.47	TRE 92	98	
Kocuria varians		0.10	0.02		10	
422241	**Good identification**					**422241**
Staphylococcus auricularis		95.10	0.42	BGA 14		
Staphylococcus haemolyticus		4.90	0.16	TRE 92		
422242	**Low discrimination**					**422242**
Staphylococcus auricularis		50.83	0.42	BGA 14		
Staphylococcus haemolyticus		47.60	0.42	TRE 92		
Staphylococcus simulans		0.98	0.05	LAC 10		
422250	**Acceptable identification**					**422250**
Staphylococcus haemolyticus		91.49	0.42	VPT 86		
Staphylococcus auricularis		8.51	0.20	TRE 93		
422252	**Low discrimination**					**422252**
Staphylococcus haemolyticus		86.29	0.37	VPT 86		
Staphylococcus simulans		12.13	0.17	TRE 93		
Staphylococcus auricularis		1.59	0.01			
422300	**Low discrimination**					**422300**
					PYR	
Staphylococcus auricularis		82.96	0.34	BGA 14	40	
Kocuria varians		13.37	0.19	GAL 10	10	
Staphylococcus haemolyticus		3.67	0.07	TRE 92	98	
422340	**Low discrimination**					**422340**
					PYR	
Staphylococcus haemolyticus		64.45	0.48	VPT 86	98	
Staphylococcus auricularis		34.33	0.42	TRE 93	40	
Staphylococcus epidermidis		0.71	0.06		1	
422342	**Low discrimination**					**422342**
					PYR	
Staphylococcus haemolyticus		85.70	0.43	VPT 86	98	
Staphylococcus auricularis		9.02	0.22	TRE 93	40	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		5.28	0.16		1	
422350	**Good identification**					**422350**
Staphylococcus haemolyticus		99.09	0.44	VPT 86		
Staphylococcus auricularis		0.91	0.01	TRE 93		
422352	**Good identification**					**422352**
Staphylococcus haemolyticus		100.0	0.38	VPT 86		
422400	**Low discrimination**					**422400**
Staphylococcus auricularis		52.89	0.76	BGA 14		
Staphylococcus carnosus subsp. utilis		47.08	0.59			
Kocuria varians		0.03	0.10			
422401	**Low discrimination**					**422401**
Staphylococcus auricularis		87.82	0.57	BGA 14		
Staphylococcus carnosus subsp. utilis		7.11	0.18	MNS 10		
Staphylococcus carnosus subsp. carnosus		3.26	0.23			
422402	**Good identification**					**422402**
Staphylococcus auricularis		91.77	0.57	BGA 14		
Staphylococcus carnosus subsp. utilis		7.43	0.18	LAC 10		
Staphylococcus simulans		0.59	0.10			
422403	**Low discrimination**					**422403**
Staphylococcus carnosus subsp. carnosus		52.86	0.35	PHS 99		
Staphylococcus auricularis		39.54	0.37			
Staphylococcus simulans		7.60	0.21			
422410	**Low discrimination**					**422410**
Staphylococcus auricularis		52.91	0.35	BGA 14		
Staphylococcus carnosus subsp. utilis		47.09	0.18	MAN 1		
422411	**Low discrimination**					**422411**
Staphylococcus carnosus subsp. carnosus		84.42	0.35	PHS 99		
Staphylococcus capitis subsp. capitis		9.84	0.15			
Staphylococcus auricularis		5.74	0.16			
422413	**Low discrimination**					**422413**
Staphylococcus carnosus subsp. carnosus		86.65	0.48	PHS 99		
Staphylococcus simulans		12.46	0.33			
Staphylococcus felis		0.89	0.01			
422420	**Low discrimination**					**422420**
Staphylococcus auricularis		52.91	0.35	BGA 14		
Staphylococcus carnosus subsp. utilis		47.09	0.18	XYL 1		
422440	**Very good identification**					**422440**
Staphylococcus auricularis		99.42	0.84	BGA 14		
Staphylococcus carnosus subsp. utilis		0.38	0.18			
Staphylococcus haemolyticus		0.20	0.29			
422441	**Very good identification**					**422441**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		100.0	0.64	BGA 14			
422442	**Very good identification**						**422442**
Staphylococcus auricularis		98.78	0.64	BGA 14			
Staphylococcus haemolyticus		1.00	0.24	LAC 10			
Staphylococcus simulans		0.12	0.03				
422443	**Good identification**						**422443**
Staphylococcus auricularis		96.43	0.45	BGA 14			
Staphylococcus simulans		3.57	0.14	MNS 10			
422450	**Low discrimination**						**422450**
					OXI	NOV	
Staphylococcus auricularis		88.85	0.43	BGA 14	1	1	
Staphylococcus haemolyticus		10.29	0.25	MAN 1	1	2	
Macrococcus bovicus		0.86	-.02		99	99	
422460	**Good identification**						**422460**
Staphylococcus auricularis		100.0	0.43	BGA 14			
422500	**Good identification**						**422500**
Staphylococcus auricularis		91.70	0.57	BGA 14			
Staphylococcus carnosus subsp. utilis		7.42	0.18	GAL 10			
Kocuria varians		0.66	0.14				
422501	**Low discrimination**						**422501**
Staphylococcus auricularis		78.81	0.37	BGA 14			
Staphylococcus schleiferi subsp. schleiferi		14.60	0.18	GAL 10			
Staphylococcus carnosus subsp. carnosus		6.59	0.11	MNS 10			
422502	**Good identification**						**422502**
Staphylococcus auricularis		95.15	0.37	BGA 14			
Staphylococcus piscifermentans		3.12	0.16	GAL 10			
Kocuria varians		1.72	0.02	LAC 10			
422513	**Low discrimination**						**422513**
Staphylococcus carnosus subsp. carnosus		65.94	0.35	PHS 99			
Staphylococcus condimenti		20.16	0.14				
Staphylococcus felis		9.07	0.12				
422540	**Very good identification**						**422540**
Staphylococcus auricularis		97.92	0.64	BGA 14			
Staphylococcus haemolyticus		1.98	0.30	GAL 10			
Staphylococcus piscifermentans		0.10	0.12				
422541	**Good identification**						**422541**
Staphylococcus auricularis		100.0	0.45	BGA 14			
422542	**Low discrimination**						**422542**
Staphylococcus auricularis		89.57	0.45	BGA 14			
Staphylococcus haemolyticus		9.17	0.25	GAL 10			
Staphylococcus piscifermentans		1.26	0.16	LAC 10			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
422600	**Very good identification**					**422600**
Staphylococcus auricularis		98.52	0.76	BGA 14		
Staphylococcus carnosus subsp. utilis		0.96	0.18			
Staphylococcus haemolyticus		0.50	0.29			
422601	**Very good identification**					**422601**
Staphylococcus auricularis		98.72	0.56	BGA 14		
Staphylococcus simulans		0.55	0.08	MNS 10		
Staphylococcus hyicus		0.47	-.01			
422602	**Low discrimination**					**422602**
Staphylococcus auricularis		83.66	0.56	BGA 14		
Staphylococcus simulans		13.87	0.38	LAC 10		
Staphylococcus haemolyticus		2.13	0.24			
422603	**Low discrimination**					**422603**
Staphylococcus simulans		76.81	0.49	URE 98		
Staphylococcus auricularis		15.37	0.36			
Staphylococcus hyicus		7.54	0.21			
422610	**Low discrimination**					**422610**
Staphylococcus auricularis		71.44	0.34	BGA 14		
Staphylococcus haemolyticus		20.92	0.25	MAN 1		
Staphylococcus simulans		5.27	0.10			
422612	**Good identification**					**422612**
Staphylococcus simulans		96.36	0.51	URE 98		
Staphylococcus haemolyticus		2.18	0.19			
Staphylococcus auricularis		1.47	0.15			
422613	**Very good identification**					**422613**
Staphylococcus simulans		98.95	0.62	URE 98		
Staphylococcus felis		0.60	0.07			
Staphylococcus carnosus subsp. carnosus		0.29	0.07			
422620	**Good identification**					**422620**
Staphylococcus auricularis		100.0	0.34	BGA 14		
422640	**Low discrimination**					**422640**
Staphylococcus auricularis		82.04	0.83	BGA 14		
Staphylococcus haemolyticus		17.53	0.70			
Staphylococcus petrasii subsp. jettensis		0.29	0.18			
422641	**Good identification**					**422641**
					OXI	
Staphylococcus auricularis		93.70	0.63	BGA 14	1	
Staphylococcus haemolyticus		5.57	0.39	MNS 10	1	
Kocuria kristinae		0.33	0.06		99	
422642	**Low discrimination**					**422642**
Staphylococcus haemolyticus		50.92	0.65	VPT 86		
Staphylococcus auricularis		47.07	0.63			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		1.50	0.31				
422643	**Low discrimination**						**422643**
Staphylococcus auricularis		43.58	0.44	BGA 14			
Staphylococcus simulans		41.93	0.42	MNS 10			
Staphylococcus haemolyticus		13.12	0.34	LAC 10			
422650	**Good identification**						**422650**
					PYR		
Staphylococcus haemolyticus		91.72	0.66	VPT 86	98		
Staphylococcus auricularis		7.38	0.42		40		
Staphylococcus warneri		0.65	0.29		8		
422651	**Low discrimination**						**422651**
Staphylococcus haemolyticus		70.78	0.35	VPT 86			
Staphylococcus auricularis		20.47	0.22	MNS 3			
Staphylococcus simulans		8.75	0.13				
422652	**Low discrimination**						**422652**
Staphylococcus haemolyticus		81.89	0.60	VPT 86			
Staphylococcus simulans		16.46	0.44				
Staphylococcus auricularis		1.30	0.22				
422653	**Good identification**						**422653**
					COA		
Staphylococcus simulans		94.86	0.54	URE 98	1		
Staphylococcus haemolyticus		4.36	0.29		1		
Staphylococcus intermedius		0.54	0.06		99		
422660	**Low discrimination**						**422660**
Staphylococcus auricularis		82.39	0.42	BGA 14			
Staphylococcus haemolyticus		17.61	0.29	XYL 1			
422700	**Good identification**						**422700**
Staphylococcus auricularis		94.78	0.56	BGA 14			
Staphylococcus haemolyticus		4.85	0.30	GAL 10			
Staphylococcus piscifermentans		0.37	0.16				
422701	**Low discrimination**						**422701**
					COA	PYR	
Staphylococcus auricularis		58.23	0.36	BGA 14	1	40	
Staphylococcus hyicus		28.55	0.21	GAL 10	24	1	
Staphylococcus intermedius		11.73	0.19	MNS 10	99	90	
422702	**Low discrimination**						**422702**
Staphylococcus auricularis		66.72	0.36	BGA 14			
Staphylococcus haemolyticus		17.27	0.25	GAL 10			
Staphylococcus simulans		8.66	0.17	LAC 10			
422703	**Low discrimination**						**422703**
					COA	PYR	
Staphylococcus hyicus		74.45	0.43	BGA 10	24	1	
Staphylococcus intermedius		16.33	0.35	PHS 92	99	90	
Staphylococcus simulans		5.74	0.27		1	70	
422711	**Good identification**						**422711**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus intermedius		100.0	0.28	URE 90			
422712	**Low discrimination**						**422712**
					COA		
Staphylococcus simulans		71.33	0.29	URE 98	1		
Staphylococcus haemolyticus		20.88	0.20	GAL 8	1		
Staphylococcus intermedius		4.64	0.03		99		
422713	**Low discrimination**						**422713**
					COA		
Staphylococcus intermedius		53.17	0.44	URE 90	99		
Staphylococcus simulans		27.63	0.40	PHS 97	1		
Staphylococcus condimenti		10.56	0.18		1		
422740	**Low discrimination**						**422740**
					PYR		
Staphylococcus haemolyticus		68.31	0.71	VPT 86	98		
Staphylococcus auricularis		31.50	0.63		40		
Staphylococcus warneri		0.11	0.21		8		
422741	**Low discrimination**						**422741**
					COA		
Staphylococcus auricularis		55.92	0.44	BGA 14	1		
Staphylococcus haemolyticus		33.76	0.40	GAL 10	1		
Staphylococcus intermedius		7.24	0.22	MNS 10	99		
422742	**Good identification**						**422742**
Staphylococcus haemolyticus		91.15	0.66	VPT 86			
Staphylococcus auricularis		8.30	0.44				
Staphylococcus simulans		0.21	0.09				
422743	**Low discrimination**						**422743**
					COA		
Staphylococcus intermedius		56.65	0.39	URE 90	99		
Staphylococcus haemolyticus		24.19	0.35	PHS 97	1		
Staphylococcus auricularis		7.92	0.24		1		
422750	**Very good identification**						**422750**
					NOV		
Staphylococcus haemolyticus		98.47	0.67	VPT 86	2		
Staphylococcus auricularis		0.78	0.22		1		
Staphylococcus saprophyticus subsp. bovis		0.55	0.18		99		
422751	**Low discrimination**						**422751**
					COA	NOV	
Staphylococcus haemolyticus		47.63	0.36	VPT 86	1	2	
Staphylococcus intermedius		47.02	0.31	MNS 3	99	1	
Staphylococcus saprophyticus subsp. bovis		2.15	0.05		1	99	
422752	**Very good identification**						**422752**
					COA		
Staphylococcus haemolyticus		97.50	0.62	VPT 86	1		
Staphylococcus simulans		1.51	0.22		1		
Staphylococcus intermedius		0.33	0.06		99		
422753	**Low discrimination**						**422753**
					COA		
Staphylococcus intermedius		78.36	0.48	URE 90	99		
Staphylococcus simulans		12.20	0.33	PHS 97	1		
Staphylococcus haemolyticus		7.27	0.30		1		
422760	**Low discrimination**						**422760**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus haemolyticus		68.44	0.30	VPT 86		
Staphylococcus auricularis		31.56	0.22	XYL 1		
423013	**Good identification**					**423013**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ESL 10		
423400	**Low discrimination**					**423400**
Staphylococcus auricularis		48.77	0.35	BGA 14		
Staphylococcus carnosus subsp. utilis		43.41	0.18	ESL 1		
Staphylococcus piscifermentans		7.83	0.28			
423402	**Low discrimination**					**423402**
Staphylococcus piscifermentans		68.42	0.32	URE 90		
Staphylococcus auricularis		31.58	0.16	BGA 1		
423413	**Good identification**					**423413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ESL 10		
423440	**Acceptable identification**					**423440**
Staphylococcus auricularis		93.57	0.43	BGA 14		
Staphylococcus piscifermentans		6.43	0.28	ESL 1		
423500	**Low discrimination**					**423500**
Staphylococcus piscifermentans		68.42	0.32	URE 90		
Staphylococcus auricularis		31.58	0.16	BGA 1		
423502	**Good identification**					**423502**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
423512	**Good identification**					**423512**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
423542	**Acceptable identification**					**423542**
Staphylococcus piscifermentans		92.61	0.36	URE 90		
Staphylococcus auricularis		7.39	0.03	BGA 1		
423552	**Good identification**					**423552**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
423600	**Low discrimination**					**423600**
Staphylococcus auricularis		79.32	0.34	BGA 14		
Staphylococcus piscifermentans		20.68	0.32	ESL 1		
423602	**Low discrimination**					**423602**
Staphylococcus piscifermentans		77.88	0.36	URE 90		
Staphylococcus auricularis		22.12	0.15	BGA 1		
423612	**Low discrimination**					**423612**
Staphylococcus piscifermentans		69.46	0.28	URE 90		
Staphylococcus simulans		30.54	0.10	BGA 1		
423640	**Low discrimination**					**423640**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		75.45	0.42	BGA 14			
Staphylococcus haemolyticus		16.12	0.29	ESL 1			
Staphylococcus piscifermentans		8.43	0.32				
423642	**Low discrimination**						**423642**
Staphylococcus piscifermentans		42.02	0.36	URE 90			
Staphylococcus haemolyticus		30.13	0.24	BGA 1			
Staphylococcus auricularis		27.85	0.22				
423652	**Low discrimination**						**423652**
Staphylococcus piscifermentans		45.87	0.28	URE 90			
Staphylococcus haemolyticus		45.07	0.19	BGA 1			
Staphylococcus simulans		9.06	0.03				
423700	**Low discrimination**						**423700**
Staphylococcus piscifermentans		65.59	0.36	URE 90			
Staphylococcus auricularis		18.63	0.15	BGA 1			
Rothia mucilaginosa		15.78	0.03				
423702	**Good identification**						**423702**
Staphylococcus piscifermentans		100.0	0.39	URE 90			
423710	**Good identification**						**423710**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
423711	**Good identification**						**423711**
Staphylococcus lentus		98.64	0.41	ARG 1			
Staphylococcus sciuri		1.36	0.01				
423712	**Low discrimination**						**423712**
					OXI	NOV	
Staphylococcus piscifermentans		78.68	0.32	URE 90	1	1	
Staphylococcus lentus		21.32	0.12	BGA 1	99	99	
423713	**Good identification**						**423713**
					OXI	COA	
Staphylococcus lentus		99.39	0.53	ARG 1	99	1	
Staphylococcus intermedius		0.61	0.03		1	99	
423731	**Good identification**						**423731**
Staphylococcus lentus		100.0	0.41	ARG 1			
423733	**Good identification**						**423733**
Staphylococcus lentus		100.0	0.53	ARG 1			
423740	**Low discrimination**						**423740**
Staphylococcus haemolyticus		39.26	0.30	VPT 86			
Staphylococcus piscifermentans		27.31	0.36	ESL 1			
Staphylococcus auricularis		18.10	0.22				
423742	**Low discrimination**						**423742**
Staphylococcus piscifermentans		62.97	0.39	URE 90			
Staphylococcus haemolyticus		33.94	0.25	BGA 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		3.09	0.03					
423751	**Good identification**							**423751**
Staphylococcus lentus		97.50	0.48	ARG 1				
Staphylococcus sciuri		2.50	0.13					
423752	**Low discrimination**							**423752**
					OXI	NOV		
Staphylococcus piscifermentans		43.69	0.32	URE 90	1	1		
Staphylococcus haemolyticus		32.28	0.20	BGA 1	1	2		
Staphylococcus lentus		24.03	0.18		99	99		
423753	**Very good identification**							**423753**
					OXI	COA	NOV	
Staphylococcus lentus		99.38	0.59	ARG 1	99	1	99	
Staphylococcus intermedius		0.45	0.06		1	99	1	
Aerococcus viridans		0.17	0.16		1	1	1	
423771	**Good identification**							**423771**
Staphylococcus lentus		98.11	0.48	ARG 1				
Staphylococcus sciuri		1.90	0.11					
423773	**Very good identification**							**423773**
					OXI	NOV		
Staphylococcus lentus		99.83	0.59	ARG 1	99	99		
Aerococcus viridans		0.17	0.16		1	1		
424001	**Low discrimination**							**424001**
Staphylococcus carnosus subsp. carnosus		53.45	0.31	NIT 99				
Staphylococcus schleiferi subsp. schleiferi		46.55	0.30					
424003	**Good identification**							**424003**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	NIT 99				
424011	**Good identification**							**424011**
Staphylococcus carnosus subsp. carnosus		96.40	0.43	NIT 99				
Staphylococcus capitis subsp. capitis		3.60	0.12					
424013	**Very good identification**							**424013**
Staphylococcus carnosus subsp. carnosus		100.0	0.55	NIT 99				
424101	**Low discrimination**							**424101**
Staphylococcus schleiferi subsp. schleiferi		77.70	0.30	VPT 99				
Staphylococcus carnosus subsp. carnosus		22.30	0.18	NIT 90				
424103	**Good identification**							**424103**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	NIT 99				
424111	**Good identification**							**424111**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	NIT 99				
424113	**Good identification**							**424113**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	NIT 99				
424240	**Very good identification**							**424240**
					COA			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus aureus subsp. anaerobius		99.97	0.59	BGA 1	99	
Staphylococcus auricularis		0.03	0.01		1	
424313	**Acceptable identification**					**424313**
					COA	
Staphylococcus intermedius		94.72	0.29	URE 90	99	
Staphylococcus carnosus subsp. carnosus		5.28	0.02	NIT 93	1	
424353	**Good identification**					**424353**
Staphylococcus intermedius		100.0	0.32	URE 90		
424401	**Low discrimination**					**424401**
Staphylococcus schleiferi subsp. schleiferi		85.91	0.39	VPT 99		
Staphylococcus carnosus subsp. carnosus		14.09	0.23	NIT 90		
424403	**Good identification**					**424403**
Staphylococcus carnosus subsp. carnosus		98.48	0.35	NIT 99		
Staphylococcus schleiferi subsp. schleiferi		1.52	-.02			
424411	**Good identification**					**424411**
Staphylococcus carnosus subsp. carnosus		98.48	0.35	NIT 99		
Staphylococcus schleiferi subsp. schleiferi		1.52	-.02			
424413	**Good identification**					**424413**
Staphylococcus carnosus subsp. carnosus		98.98	0.48	NIT 99		
Staphylococcus felis		1.02	0.01			
424450	**Good identification**					**424450**
Staphylococcus kloosii		100.0	0.52	ARG 1		
424452	**Good identification**					**424452**
Staphylococcus kloosii		100.0	0.53	ARG 1		
424470	**Good identification**					**424470**
Staphylococcus kloosii		100.0	0.40	ARG 1		
424472	**Good identification**					**424472**
Staphylococcus kloosii		100.0	0.40	ARG 1		
424501	**Good identification**					**424501**
Staphylococcus schleiferi subsp. schleiferi		96.06	0.39	VPT 99		
Staphylococcus carnosus subsp. carnosus		3.94	0.11	NIT 90		
424510	**Good identification**					**424510**
Staphylococcus kloosii		100.0	0.32	ARG 1		
424512	**Good identification**					**424512**
Staphylococcus kloosii		100.0	0.32	ARG 1		
424513	**Low discrimination**					**424513**
Staphylococcus carnosus subsp. carnosus		66.66	0.35	NIT 99		
Staphylococcus condimenti		20.38	0.14			
Staphylococcus felis		9.17	0.12			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
424550	**Very good identification**							**424550**
	Staphylococcus kloosii	100.0	0.58	ARG 1				
424552	**Very good identification**							**424552**
	Staphylococcus kloosii	100.0	0.59	ARG 1				
424570	**Good identification**							**424570**
	Staphylococcus kloosii	100.0	0.46	ARG 1				
424572	**Good identification**							**424572**
	Staphylococcus kloosii	100.0	0.46	ARG 1				
424650	**Good identification**							**424650**
	Staphylococcus kloosii	100.0	0.31	ARG 1				
424652	**Low discrimination**							**424652**
					NOV	YEP		
	Staphylococcus kloosii	73.62	0.32	ARG 1	99	99		
	Staphylococcus simulans	26.38	0.05	SUC 9	1	1		
424701	**Low discrimination**							**424701**
					COA	PYR		
	Staphylococcus intermedius	86.08	0.27	URE 90	99	90		
	Staphylococcus hyicus	10.01	0.02	NIT 93	24	1		
	Staphylococcus schleiferi subsp. schleiferi	3.92	-.02	LAC 86	1	37		
424703	**Low discrimination**							**424703**
					YEP	PYR	COA	
	Staphylococcus intermedius	78.84	0.43	URE 90	1	90	99	
	Staphylococcus hyicus	17.16	0.24	NIT 93	1	1	24	
	Staphylococcus chromogenes	2.42	0.10		95	45	1	
424711	**Good identification**							**424711**
	Staphylococcus intermedius	100.0	0.36	URE 90				
424713	**Low discrimination**							**424713**
					COA			
	Staphylococcus intermedius	87.91	0.52	URE 90	99			
	Staphylococcus condimenti	7.18	0.18	NIT 93	1			
	Staphylococcus felis	4.37	0.18		1			
424741	**Good identification**							**424741**
	Staphylococcus intermedius	100.0	0.30	URE 90				
424743	**Good identification**							**424743**
					COA	YEP		
	Staphylococcus intermedius	98.00	0.47	URE 90	99	1		
	Staphylococcus chromogenes	2.00	0.10	NIT 93	1	95		
424750	**Acceptable identification**							**424750**
					COA	NOV		
	Staphylococcus kloosii	90.27	0.37	ARG 1	1	99		
	Staphylococcus intermedius	9.73	-.02	SUC 9	99	1		
424751	**Good identification**							**424751**
	Staphylococcus intermedius	100.0	0.39	URE 90				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
424752	**Low discrimination**						**424752**
					COA	NOV	
Staphylococcus kloosii		62.07	0.38	ARG 1	1	99	
Staphylococcus intermedius		37.93	0.14	SUC 9	99	1	
424753	**Very good identification**						**424753**
					COA	YEP	
Staphylococcus intermedius		99.79	0.55	URE 90	99	1	
Staphylococcus chromogenes		0.21	-.01	NIT 93	1	95	
424772	**Acceptable identification**						**424772**
Staphylococcus arlettae		90.92	0.35	ARG 1			
Staphylococcus kloosii		9.08	0.26	GLR 92			
424773	**Low discrimination**						**424773**
					COA	NOV	
Staphylococcus arlettae		84.46	0.37	ARG 1	1	99	
Staphylococcus intermedius		15.54	0.14	GLR 92	99	1	
425013	**Good identification**						**425013**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ESL 10			
425413	**Good identification**						**425413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ESL 10			
425450	**Good identification**						**425450**
Staphylococcus kloosii		100.0	0.52	ARG 1			
425452	**Good identification**						**425452**
Staphylococcus kloosii		100.0	0.53	ARG 1			
425470	**Good identification**						**425470**
Staphylococcus kloosii		100.0	0.40	ARG 1			
425472	**Good identification**						**425472**
Staphylococcus kloosii		100.0	0.40	ARG 1			
425510	**Good identification**						**425510**
Staphylococcus kloosii		100.0	0.32	ARG 1			
425512	**Good identification**						**425512**
Staphylococcus kloosii		100.0	0.32	ARG 1			
425550	**Very good identification**						**425550**
Staphylococcus kloosii		100.0	0.58	ARG 1			
425552	**Very good identification**						**425552**
Staphylococcus kloosii		100.0	0.59	ARG 1			
425570	**Good identification**						**425570**
Staphylococcus kloosii		100.0	0.46	ARG 1			
425572	**Good identification**						**425572**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.46	ARG 1				
425650	**Good identification**							**425650**
Staphylococcus kloosii		100.0	0.31	ARG 1				
425652	**Good identification**							**425652**
Staphylococcus kloosii		100.0	0.32	ARG 1				
425750	**Good identification**							**425750**
Staphylococcus kloosii		100.0	0.37	ARG 1				
425752	**Good identification**							**425752**
Staphylococcus kloosii		100.0	0.38	ARG 1				
425772	**Acceptable identification**							**425772**
Staphylococcus arlettae		90.92	0.35	ARG 1				
Staphylococcus kloosii		9.08	0.26	GLR 92				
425773	**Acceptable identification**							**425773**
					OXI	NOV	YEP	
Staphylococcus arlettae		93.42	0.37	ARG 1	1	99	99	
Staphylococcus lentus		3.94	0.05	GLR 92	99	99	1	
Aerococcus viridans		2.64	0.16		1	1	1	
426000	**Low discrimination**							**426000**
Staphylococcus carnosus subsp. carnosus		74.17	0.31	MNS 99				
Staphylococcus carnosus subsp. utilis		13.19	0.03					
Staphylococcus auricularis		6.77	0.13					
426001	**Good identification**							**426001**
Staphylococcus carnosus subsp. carnosus		92.58	0.72					
Staphylococcus schleiferi subsp. schleiferi		7.33	0.49					
Staphylococcus capitis subsp. capitis		0.09	0.08					
426002	**Good identification**							**426002**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	MNS 99				
426003	**Very good identification**							**426003**
Staphylococcus carnosus subsp. carnosus		99.98	0.84					
Staphylococcus schleiferi subsp. schleiferi		0.02	0.08					
426010	**Good identification**							**426010**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	MNS 99				
426011	**Very good identification**							**426011**
Staphylococcus carnosus subsp. carnosus		99.68	0.84					
Staphylococcus capitis subsp. capitis		0.30	0.31					
Staphylococcus schleiferi subsp. schleiferi		0.02	0.08					
426012	**Very good identification**							**426012**
Staphylococcus carnosus subsp. carnosus		100.0	0.55	MNS 99				
426013	**Excellent identification**							**426013**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		99.99	0.96			
Staphylococcus felis		0.00	0.01			
Staphylococcus capitis subsp. capitis		0.00	-.04			
426021	**Acceptable identification**					**426021**
Staphylococcus carnosus subsp. carnosus		92.66	0.31	XYL 1		
Staphylococcus schleiferi subsp. schleiferi		7.34	0.08			
426023	**Good identification**					**426023**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	XYL 1		
426031	**Good identification**					**426031**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	XYL 1		
426033	**Very good identification**					**426033**
Staphylococcus carnosus subsp. carnosus		100.0	0.55	XYL 1		
426041	**Acceptable identification**					**426041**
Staphylococcus carnosus subsp. carnosus		90.68	0.31	MLT 1		
Staphylococcus schleiferi subsp. schleiferi		7.18	0.08			
Staphylococcus auricularis		2.15	0.01			
426043	**Good identification**					**426043**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	MLT 1		
426051	**Good identification**					**426051**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	MLT 1		
426053	**Very good identification**					**426053**
Staphylococcus carnosus subsp. carnosus		100.0	0.55	MLT 1		
426101	**Low discrimination**					**426101**
					COA	
Staphylococcus carnosus subsp. carnosus		75.89	0.59		1	
Staphylococcus schleiferi subsp. schleiferi		24.04	0.49		1	
Staphylococcus schleiferi subsp. coagulans		0.08	0.02		99	
426102	**Good identification**					**426102**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	MNS 99		
426103	**Very good identification**					**426103**
					COA	
Staphylococcus carnosus subsp. carnosus		99.63	0.72		1	
Staphylococcus schleiferi subsp. coagulans		0.16	0.18		99	
Staphylococcus schleiferi subsp. schleiferi		0.08	0.08		1	
426110	**Good identification**					**426110**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	MNS 99		
426111	**Very good identification**					**426111**
					COA	
Staphylococcus carnosus subsp. carnosus		99.90	0.72		1	
Staphylococcus schleiferi subsp. schleiferi		0.08	0.08		1	
Staphylococcus schleiferi subsp. coagulans		0.02	0.01		99	
426112	**Good identification**					**426112**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		100.0	0.43	MNS 99			
426113	**Very good identification**						**426113**
Staphylococcus carnosus subsp. carnosus		99.75	0.84				
Staphylococcus condimenti		0.13	0.14				
Staphylococcus felis		0.06	0.12				
426123	**Good identification**						**426123**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	XYL 1			
426131	**Good identification**						**426131**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	XYL 1			
426133	**Good identification**						**426133**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	XYL 1			
426143	**Acceptable identification**						**426143**
					COA		
Staphylococcus carnosus subsp. carnosus		94.88	0.31	MLT 1	1		
Staphylococcus intermedius		5.12	0.06		99		
426151	**Good identification**						**426151**
					COA		
Staphylococcus carnosus subsp. carnosus		97.68	0.31	MLT 1	1		
Staphylococcus intermedius		2.32	-.02		99		
426153	**Good identification**						**426153**
					COA		
Staphylococcus carnosus subsp. carnosus		96.48	0.43	MLT 1	1		
Staphylococcus intermedius		3.52	0.14		99		
426201	**Low discrimination**						**426201**
Staphylococcus carnosus subsp. carnosus		77.00	0.31	SUC 1			
Staphylococcus capitis subsp. capitis		13.02	0.14				
Staphylococcus schleiferi subsp. schleiferi		6.10	0.08				
426203	**Low discrimination**						**426203**
					YEP		
Staphylococcus carnosus subsp. carnosus		82.54	0.43	SUC 1	1		
Staphylococcus hyicus		11.98	0.21		1		
Staphylococcus chromogenes		2.16	0.10		95		
426211	**Low discrimination**						**426211**
Staphylococcus carnosus subsp. carnosus		64.04	0.43	SUC 1			
Staphylococcus capitis subsp. capitis		35.97	0.37				
426213	**Very good identification**						**426213**
Staphylococcus carnosus subsp. carnosus		95.63	0.55	SUC 1			
Staphylococcus simulans		2.48	0.25				
Staphylococcus felis		0.86	0.07				
426303	**Low discrimination**						**426303**
					COA	PYR	
Staphylococcus hyicus		54.27	0.43	BGA 10	24	1	
Staphylococcus intermedius		28.97	0.43	TRE 92	99	90	
Staphylococcus carnosus subsp. carnosus		8.13	0.31		1	72	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
426311	**Low discrimination**							**426311**
					COA	YEP		
Staphylococcus intermedius		59.85	0.36	URE 90	99	1		
Staphylococcus carnosus subsp. carnosus		38.16	0.31	TRE 93	1	1		
Staphylococcus chromogenes		1.12	-.01	LAC 86	1	95		
426313	**Low discrimination**							**426313**
					COA			
Staphylococcus intermedius		62.92	0.52	URE 90	99			
Staphylococcus carnosus subsp. carnosus		26.12	0.43	TRE 93	1			
Staphylococcus condimenti		5.14	0.18		1			
426341	**Low discrimination**							**426341**
					COA	YEP	PYR	
Staphylococcus intermedius		60.34	0.30	URE 90	99	1	90	
Staphylococcus epidermidis		32.42	0.27	TRE 93	1	1	1	
Staphylococcus chromogenes		7.25	0.10	LAC 86	1	95	45	
426342	**Acceptable identification**							**426342**
					COA	PYR		
Staphylococcus epidermidis		91.19	0.32	VPT 96	1	1		
Staphylococcus intermedius		6.18	0.06	URE 92	99	90		
Staphylococcus haemolyticus		2.64	0.02		1	98		
426343	**Low discrimination**							**426343**
					COA	YEP	PYR	
Staphylococcus intermedius		68.24	0.47	URE 90	99	1	90	
Staphylococcus epidermidis		18.90	0.37	TRE 93	1	1	1	
Staphylococcus chromogenes		12.00	0.30		1	95	45	
426351	**Good identification**							**426351**
					COA	YEP		
Staphylococcus intermedius		98.76	0.39	URE 90	99	1		
Staphylococcus chromogenes		1.24	-.01	TRE 93	1	95		
426353	**Good identification**							**426353**
					COA	YEP		
Staphylococcus intermedius		93.77	0.55	URE 90	99	1		
Staphylococcus delphini		4.25	0.18	TRE 93	99	1		
Staphylococcus chromogenes		1.72	0.18		1	95		
426400	**Low discrimination**							**426400**
Staphylococcus auricularis		39.60	0.35	BGA 14				
Staphylococcus carnosus subsp. utilis		35.25	0.18	PHS 1				
Staphylococcus carnosus subsp. carnosus		16.18	0.23					
426401	**Low discrimination**							**426401**
Staphylococcus carnosus subsp. carnosus		64.22	0.64					
Staphylococcus schleiferi subsp. schleiferi		35.60	0.59					
Staphylococcus auricularis		0.18	0.16					
426402	**Acceptable identification**							**426402**
Staphylococcus carnosus subsp. carnosus		92.40	0.35	MNS 99				
Staphylococcus auricularis		6.28	0.16					
Staphylococcus simulans		1.32	0.00					
426403	**Very good identification**							**426403**
Staphylococcus carnosus subsp. carnosus		99.17	0.76					

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus felis		0.61	0.25			
Staphylococcus schleiferi subsp. schleiferi		0.14	0.18			
426410	**Good identification**					**426410**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	MNS 99		
426411	**Very good identification**					**426411**
Staphylococcus carnosus subsp. carnosus		99.80	0.76			
Staphylococcus schleiferi subsp. schleiferi		0.14	0.18			
Staphylococcus felis		0.04	0.01			
426412	**Good identification**					**426412**
Staphylococcus carnosus subsp. carnosus		97.60	0.48	MNS 99		
Staphylococcus simulans		1.40	0.13			
Staphylococcus felis		1.00	0.01			
426413	**Very good identification**					**426413**
Staphylococcus carnosus subsp. carnosus		98.86	0.89			
Staphylococcus felis		1.02	0.42			
Staphylococcus condimenti		0.08	0.14			
426423	**Good identification**					**426423**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	XYL 1		
426431	**Good identification**					**426431**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	XYL 1		
426433	**Good identification**					**426433**
Staphylococcus carnosus subsp. carnosus		98.98	0.48	XYL 1		
Staphylococcus felis		1.02	0.01			
426440	**Good identification**					**426440**
Staphylococcus auricularis		100.0	0.43	BGA 14		
426443	**Good identification**					**426443**
Staphylococcus carnosus subsp. carnosus		96.23	0.35	MLT 1		
Staphylococcus simulans		2.07	0.04			
Staphylococcus auricularis		1.70	0.03			
426451	**Good identification**					**426451**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	MLT 1		
426453	**Good identification**					**426453**
Staphylococcus carnosus subsp. carnosus		96.92	0.48	MLT 1		
Staphylococcus simulans		2.09	0.16			
Staphylococcus felis		1.00	0.01			
426501	**Low discrimination**					**426501**
					COA	
Staphylococcus schleiferi subsp. schleiferi		68.55	0.59	VPT 99	1	
Staphylococcus carnosus subsp. carnosus		30.92	0.52		1	
Staphylococcus intermedius		0.23	0.09		99	
426503	**Low discrimination**					**426503**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		88.32	0.64				
Staphylococcus felis		7.26	0.36				
Staphylococcus hyicus		1.61	0.24				
426511	**Very good identification**						**426511**
Staphylococcus carnosus subsp. carnosus		97.23	0.64				
Staphylococcus condimenti		1.20	0.14				
Staphylococcus schleiferi subsp. schleiferi		0.54	0.18				
426512	**Low discrimination**						**426512**
Staphylococcus carnosus subsp. carnosus		69.29	0.35	MNS 99			
Staphylococcus condimenti		21.18	0.14				
Staphylococcus felis		9.53	0.12				
426513	**Low discrimination**						**426513**
Staphylococcus carnosus subsp. carnosus		68.91	0.76				
Staphylococcus condimenti		21.07	0.55				
Staphylococcus felis		9.48	0.53				
426533	**Low discrimination**						**426533**
Staphylococcus carnosus subsp. carnosus		69.29	0.35	XYL 1			
Staphylococcus condimenti		21.18	0.14				
Staphylococcus felis		9.53	0.12				
426543	**Low discrimination**						**426543**
					COA	YEP	
Staphylococcus intermedius		55.59	0.29	URE 90	99	1	
Staphylococcus carnosus subsp. carnosus		33.22	0.23	SUC 99	1	1	
Staphylococcus chromogenes		8.46	0.10		1	95	
426553	**Low discrimination**						**426553**
					COA		
Staphylococcus intermedius		43.00	0.38	URE 90	99		
Staphylococcus carnosus subsp. carnosus		38.01	0.35	SUC 99	1		
Staphylococcus condimenti		11.62	0.14		1		
426600	**Good identification**						**426600**
Staphylococcus auricularis		100.0	0.34	BGA 14			
426602	**Low discrimination**						**426602**
Staphylococcus simulans		72.40	0.29	URE 98			
Staphylococcus auricularis		13.23	0.15				
Staphylococcus hyicus		8.29	0.02				
426603	**Low discrimination**						**426603**
Staphylococcus hyicus		47.15	0.43	BGA 10			
Staphylococcus felis		14.95	0.31	GAL 92			
Staphylococcus simulans		13.93	0.39				
426611	**Low discrimination**						**426611**
					COA		
Staphylococcus carnosus subsp. carnosus		76.84	0.35	SUC 1	1		
Staphylococcus intermedius		9.43	0.18		99		
Staphylococcus felis		6.41	0.07		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
426612	**Acceptable identification**							**426612**
Staphylococcus simulans		91.73	0.41	URE 98				
Staphylococcus felis		5.57	0.07					
Staphylococcus carnosus subsp. carnosus		2.70	0.07					
426613	**Low discrimination**							**426613**
Staphylococcus felis		45.11	0.48	URE 99				
Staphylococcus simulans		25.13	0.52					
Staphylococcus carnosus subsp. carnosus		21.85	0.48					
426640	**Low discrimination**							**426640**
Staphylococcus auricularis		82.39	0.42	BGA 14				
Staphylococcus haemolyticus		17.61	0.29	PHS 1				
426643	**Low discrimination**							**426643**
					COA	YEP		
Staphylococcus chromogenes		41.13	0.32	URE 99	1	95		
Staphylococcus simulans		30.32	0.32	GAL 90	1	1		
Staphylococcus intermedius		24.56	0.29		99	1		
426652	**Low discrimination**							**426652**
Staphylococcus simulans		86.90	0.34	URE 98				
Staphylococcus haemolyticus		13.10	0.19					
426653	**Low discrimination**							**426653**
					COA	YEP		
Staphylococcus simulans		58.39	0.45	URE 98	1	1		
Staphylococcus intermedius		31.96	0.38		99	1		
Staphylococcus chromogenes		5.58	0.20		1	95		
426701	**Low discrimination**							**426701**
					YEP	PYR	COA	
Staphylococcus intermedius		48.17	0.50	URE 90	1	90	99	
Staphylococcus hyicus		41.72	0.43	LAC 86	1	1	24	
Staphylococcus chromogenes		7.51	0.32		95	45	1	
426703	**Low discrimination**							**426703**
					YEP	PYR	COA	
Staphylococcus hyicus		53.54	0.65	BGA 10	1	1	24	
Staphylococcus intermedius		33.02	0.66		1	90	99	
Staphylococcus chromogenes		7.54	0.51		95	45	1	
426711	**Good identification**							**426711**
					COA			
Staphylococcus intermedius		90.85	0.59	URE 90	99			
Staphylococcus condimenti		3.43	0.18	LAC 86	1			
Staphylococcus felis		2.09	0.18		1			
426712	**Low discrimination**							**426712**
					COA			
Staphylococcus intermedius		46.28	0.34	URE 90	99			
Staphylococcus condimenti		28.15	0.18	MNS 99	1			
Staphylococcus felis		17.14	0.18		1			
426713	**Low discrimination**							**426713**
					COA			
Staphylococcus intermedius		49.33	0.75	URE 90	99			
Staphylococcus condimenti		30.00	0.59		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus felis		18.27	0.59		1			
426733	**Low discrimination**							**426733**
					COA			
Staphylococcus intermedius		49.94	0.34	URE 90	99			
Staphylococcus condimenti		30.37	0.18	XYL 1	1			
Staphylococcus felis		18.50	0.18		1			
426740	**Low discrimination**							**426740**
					COA			
Staphylococcus haemolyticus		57.90	0.30	VPT 86	1			
Staphylococcus auricularis		26.70	0.22	PHS 1	1			
Staphylococcus intermedius		12.42	0.12		99			
426741	**Low discrimination**							**426741**
					YEP	PYR	COA	
Staphylococcus intermedius		89.91	0.53	URE 90	1	90	99	
Staphylococcus chromogenes		9.35	0.32	LAC 86	95	45	1	
Staphylococcus hyicus		0.52	0.02		1	1	24	
426742	**Low discrimination**							**426742**
					COA	YEP		
Staphylococcus intermedius		58.52	0.29	URE 90	99	1		
Staphylococcus haemolyticus		24.99	0.25	MNS 99	1	20		
Staphylococcus chromogenes		8.91	0.10		1	95		
426743	**Low discrimination**							**426743**
					YEP	PYR	COA	
Staphylococcus intermedius		85.82	0.70	URE 90	1	90	99	
Staphylococcus chromogenes		13.07	0.51		95	45	1	
Staphylococcus hyicus		0.94	0.24		1	1	24	
426751	**Very good identification**							**426751**
					COA	YEP		
Staphylococcus intermedius		98.93	0.62	URE 90	99	1		
Staphylococcus chromogenes		1.07	0.20	LAC 86	1	95		
426752	**Low discrimination**							**426752**
					COA			
Staphylococcus intermedius		86.50	0.38	URE 90	99			
Staphylococcus haemolyticus		8.02	0.20	MNS 99	1			
Staphylococcus simulans		4.11	0.12		1			
426753	**Very good identification**							**426753**
					COA	PYR	YEP	
Staphylococcus intermedius		96.79	0.79	URE 90	99	90	1	
Staphylococcus chromogenes		1.54	0.40		1	45	95	
Staphylococcus aureus subsp. aureus		0.85	0.22		99	10	84	
426763	**Low discrimination**							**426763**
					COA	PYR	YEP	
Staphylococcus intermedius		68.21	0.29	URE 90	99	90	1	
Staphylococcus lutrae		21.40	0.12	XYL 1	99	10	1	
Staphylococcus chromogenes		10.39	0.10		1	45	95	
426773	**Low discrimination**							**426773**
					OXI	PYR	COA	
Staphylococcus intermedius		79.47	0.38	URE 90	1	90	99	
Staphylococcus lutrae		17.90	0.18	XYL 1	1	10	99	
Staphylococcus lentus		1.37	0.05		99	6	1	
427001	**Good identification**							**427001**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		99.29	0.52	ESL 10		
Staphylococcus schleiferi subsp. schleiferi		0.72	0.08			
427003	**Very good identification**					**427003**
Staphylococcus carnosus subsp. carnosus		100.0	0.64	ESL 10		
427011	**Very good identification**					**427011**
Staphylococcus carnosus subsp. carnosus		100.0	0.64	ESL 10		
427012	**Good identification**					**427012**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ESL 10		
427013	**Very good identification**					**427013**
Staphylococcus carnosus subsp. carnosus		100.0	0.77	ESL 10		
427033	**Good identification**					**427033**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ESL 10		
427053	**Good identification**					**427053**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ESL 10		
427101	**Good identification**					**427101**
					COA	
Staphylococcus carnosus subsp. carnosus		96.38	0.40	ESL 10	1	
Staphylococcus schleiferi subsp. schleiferi		2.78	0.08		1	
Staphylococcus schleiferi subsp. coagulans		0.84	0.01		99	
427103	**Good identification**					**427103**
					COA	
Staphylococcus carnosus subsp. carnosus		98.68	0.52	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		1.33	0.18		99	
427111	**Good identification**					**427111**
					COA	
Staphylococcus carnosus subsp. carnosus		99.80	0.52	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		0.20	0.01		99	
427113	**Very good identification**					**427113**
					COA	
Staphylococcus carnosus subsp. carnosus		99.69	0.64	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		0.31	0.17		99	
427213	**Good identification**					**427213**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ESL 10		
427401	**Good identification**					**427401**
Staphylococcus carnosus subsp. carnosus		95.20	0.44	ESL 10		
Staphylococcus schleiferi subsp. schleiferi		4.80	0.18			
427403	**Very good identification**					**427403**
Staphylococcus carnosus subsp. carnosus		100.0	0.57	ESL 10		
427411	**Very good identification**					**427411**
Staphylococcus carnosus subsp. carnosus		100.0	0.57	ESL 10		
427412	**Acceptable identification**					**427412**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		91.35	0.28	ESL 10		
Staphylococcus piscifermentans		8.65	0.21	MNS 99		
427413	**Very good identification**					**427413**
Staphylococcus carnosus subsp. carnosus		99.91	0.69	ESL 10		
Staphylococcus felis		0.09	0.01			
427433	**Good identification**					**427433**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ESL 10		
427442	**Good identification**					**427442**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
427453	**Good identification**					**427453**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ESL 10		
427500	**Good identification**					**427500**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
427501	**Low discrimination**					**427501**
Staphylococcus carnosus subsp. carnosus		83.23	0.32	ESL 10		
Staphylococcus schleiferi subsp. schleiferi		16.78	0.18			
427502	**Low discrimination**					**427502**
Staphylococcus piscifermentans		84.14	0.32	URE 90		
Staphylococcus carnosus subsp. carnosus		15.87	0.03	BGA 1		
427503	**Good identification**					**427503**
Staphylococcus carnosus subsp. carnosus		99.26	0.44	ESL 10		
Staphylococcus felis		0.74	-.06			
427511	**Good identification**					**427511**
Staphylococcus carnosus subsp. carnosus		100.0	0.44	ESL 10		
427513	**Very good identification**					**427513**
Staphylococcus carnosus subsp. carnosus		96.13	0.57	ESL 10		
Staphylococcus condimenti		2.67	0.14			
Staphylococcus felis		1.20	0.12			
427540	**Good identification**					**427540**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
427542	**Good identification**					**427542**
Staphylococcus piscifermentans		100.0	0.32	URE 90		
427600	**Good identification**					**427600**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
427602	**Good identification**					**427602**
Staphylococcus piscifermentans		100.0	0.32	URE 90		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
427613	**Low discrimination**							**427613**
Staphylococcus carnosus subsp. carnosus		77.38	0.28	ESL 10				
Staphylococcus felis		14.53	0.07	SUC 1				
Staphylococcus simulans		8.09	0.11					
427640	**Low discrimination**							**427640**
Staphylococcus piscifermentans		88.06	0.28	URE 90				
Staphylococcus auricularis		11.94	0.01	BGA 1				
427642	**Good identification**							**427642**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
427700	**Good identification**							**427700**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
427702	**Good identification**							**427702**
Staphylococcus piscifermentans		100.0	0.36	URE 90				
427711	**Low discrimination**							**427711**
					OXI	COA	NOV	
Staphylococcus lentus		60.44	0.29	ARG 1	99	1	99	
Staphylococcus intermedius		27.62	0.18		1	99	1	
Staphylococcus carnosus subsp. carnosus		6.25	0.03		1	1	1	
427712	**Good identification**							**427712**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
427713	**Low discrimination**							**427713**
					OXI	COA	NOV	
Staphylococcus lentus		37.01	0.40	ARG 1	99	1	99	
Staphylococcus intermedius		29.30	0.34		1	99	1	
Staphylococcus condimenti		17.82	0.18		1	1	1	
427731	**Acceptable identification**							**427731**
Staphylococcus lentus		93.37	0.29	ARG 1				
Staphylococcus sciuri		6.63	0.04					
427733	**Good identification**							**427733**
Staphylococcus lentus		100.0	0.40	ARG 1				
427740	**Good identification**							**427740**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
427742	**Good identification**							**427742**
Staphylococcus piscifermentans		100.0	0.36	URE 90				
427743	**Low discrimination**							**427743**
					OXI	COA	YEP	
Staphylococcus intermedius		83.41	0.29	URE 90	1	99	1	
Staphylococcus chromogenes		12.70	0.10	ESL 1	1	1	95	
Staphylococcus lentus		3.89	0.05		99	1	1	
427751	**Low discrimination**							**427751**
					OXI	COA		
Staphylococcus lentus		66.14	0.35	ARG 1	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		22.33	0.21		1	99		
Staphylococcus sciuri		11.53	0.18		99	1		
427752	**Low discrimination**							**427752**
					OXI	NOV		
Staphylococcus piscifermentans		82.90	0.28	URE 90	1	1		
Staphylococcus lentus		17.10	0.05	BGA 1	99	99		
427753	**Low discrimination**							**427753**
					OXI	COA	YEP	
Staphylococcus lentus		62.41	0.47	ARG 1	99	1	1	
Staphylococcus intermedius		36.51	0.38		1	99	1	
Staphylococcus chromogenes		0.58	-.01		1	1	95	
427771	**Low discrimination**							**427771**
					OXI			
Staphylococcus lentus		63.93	0.35	ARG 1	99			
Staphylococcus gallinarum		27.66	0.20		1			
Staphylococcus sciuri		8.41	0.15		99			
427773	**Low discrimination**							**427773**
					OXI			
Staphylococcus lentus		76.99	0.47	ARG 1	99			
Staphylococcus gallinarum		23.01	0.28		1			
432400	**Low discrimination**							**432400**
Staphylococcus auricularis		52.91	0.35	ORN 1				
Staphylococcus carnosus subsp. utilis		47.09	0.18	BGA 14				
432440	**Good identification**							**432440**
Staphylococcus auricularis		100.0	0.43	ORN 1				
432600	**Good identification**							**432600**
Staphylococcus auricularis		100.0	0.34	ORN 1				
432640	**Low discrimination**							**432640**
Staphylococcus auricularis		82.39	0.42	ORN 1				
Staphylococcus haemolyticus		17.61	0.29	BGA 14				
432740	**Low discrimination**							**432740**
Staphylococcus haemolyticus		68.44	0.30	VPT 86				
Staphylococcus auricularis		31.56	0.22	ORN 1				
436001	**Acceptable identification**							**436001**
Staphylococcus carnosus subsp. carnosus		92.66	0.31	ORN 1				
Staphylococcus schleiferi subsp. schleiferi		7.34	0.08					
436003	**Good identification**							**436003**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	ORN 1				
436011	**Good identification**							**436011**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	ORN 1				
436013	**Very good identification**							**436013**
Staphylococcus carnosus subsp. carnosus		100.0	0.55	ORN 1				
436103	**Good identification**							**436103**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		100.0	0.31	ORN 1			
436111	**Good identification**						**436111**
Staphylococcus carnosus subsp. carnosus		100.0	0.31	ORN 1			
436113	**Good identification**						**436113**
Staphylococcus carnosus subsp. carnosus		100.0	0.43	ORN 1			
436403	**Good identification**						**436403**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	ORN 1			
436411	**Good identification**						**436411**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	ORN 1			
436413	**Good identification**						**436413**
Staphylococcus carnosus subsp. carnosus		98.98	0.48	ORN 1			
Staphylococcus felis		1.02	0.01				
436513	**Low discrimination**						**436513**
Staphylococcus carnosus subsp. carnosus		69.29	0.35	ORN 1			
Staphylococcus condimenti		21.18	0.14				
Staphylococcus felis		9.53	0.12				
436713	**Low discrimination**						**436713**
					COA		
Staphylococcus intermedius		49.94	0.34	URE 90	99		
Staphylococcus condimenti		30.37	0.18	ORN 1	1		
Staphylococcus felis		18.50	0.18		1		
436743	**Low discrimination**						**436743**
					COA	YEP	
Staphylococcus intermedius		86.79	0.29	URE 90	99	1	
Staphylococcus chromogenes		13.21	0.10	ORN 1	1	95	
436753	**Good identification**						**436753**
					COA	YEP	
Staphylococcus intermedius		98.44	0.38	URE 90	99	1	
Staphylococcus chromogenes		1.56	-.01	ORN 1	1	95	
437013	**Good identification**						**437013**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	ORN 1			
437413	**Good identification**						**437413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	ORN 1			
440000	**Low discrimination**						**440000**
					OXI		
Staphylococcus carnosus subsp. utilis		50.04	0.29	GLR 1	1		
Kocuria rosea		29.23	0.33		35		
Micrococcus sp.		13.20	0.18		89		
440011	**Good identification**						**440011**
Staphylococcus capitis subsp. capitis		100.0	0.31	GLR 1			
440040	**Low discrimination**						**440040**
					OXI		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus auricularis		77.53	0.27	GLR 3	1			
Micrococcus sp.		11.60	-.05	NIT 90	89			
Staphylococcus carnosus subsp. utilis		5.91	-.12	TRE 92	1			
440211	**Good identification**							**440211**
Staphylococcus capitis subsp. capitis		100.0	0.37	GLR 1				
440242	**Low discrimination**							**440242**
					PYR			
Staphylococcus warneri		72.43	0.28	URE 98	8			
Staphylococcus auricularis		18.32	0.07	TRE 95	40			
Staphylococcus haemolyticus		9.26	0.02		98			
440250	**Low discrimination**							**440250**
					OXI	PYR	NOV	
Staphylococcus warneri		86.83	0.35	URE 98	1	8	1	
Macrococcus bovicus		7.83	0.02	TRE 95	99	1	99	
Staphylococcus haemolyticus		5.34	0.02		1	98	2	
440340	**Low discrimination**							**440340**
					PYR			
Staphylococcus warneri		65.04	0.27	URE 98	8			
Staphylococcus haemolyticus		17.60	0.08	TRE 95	98			
Staphylococcus auricularis		17.36	0.07		40			
440400	**Low discrimination**							**440400**
					NOV			
Staphylococcus carnosus subsp. utilis		86.84	0.44	GLR 1	1			
Staphylococcus auricularis		10.81	0.42		1			
Staphylococcus cohnii subsp. cohnii		2.21	0.31		99			
440401	**Low discrimination**							**440401**
					NOV			
Staphylococcus cohnii subsp. cohnii		53.47	0.32	ARG 1	99			
Staphylococcus auricularis		26.90	0.22		1			
Staphylococcus carnosus subsp. utilis		19.64	0.03		1			
440410	**Low discrimination**							**440410**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		66.17	0.33	ARG 1	1	99	12	
Staphylococcus carnosus subsp. utilis		20.69	0.03		1	1	10	
Macrococcus bovicus		10.57	0.10		99	99	99	
440411	**Good identification**							**440411**
					NOV			
Staphylococcus cohnii subsp. cohnii		96.15	0.34	ARG 1	99			
Staphylococcus capitis subsp. capitis		3.85	-.04		1			
440440	**Low discrimination**							**440440**
					NOV			
Staphylococcus auricularis		70.56	0.49	GLR 3	1			
Staphylococcus cohnii subsp. cohnii		19.52	0.41	NIT 90	99			
Staphylococcus warneri		7.47	0.38		1			
440441	**Low discrimination**							**440441**
					NOV			
Staphylococcus cohnii subsp. cohnii		72.96	0.42	ARG 1	99			
Staphylococcus auricularis		27.04	0.30		1			
440442	**Low discrimination**							**440442**
Staphylococcus auricularis		75.91	0.30	GLR 3				

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		24.09	0.28	NIT 90				
440450	**Low discrimination**							**440450**
					OXI	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		73.32	0.43	ARG 1	1	99	12	
Staphylococcus warneri		15.95	0.35		1	1	43	
Macrococcus bovicus		8.63	0.18		99	99	99	
440451	**Good identification**							**440451**
Staphylococcus cohnii subsp. cohnii		100.0	0.44	ARG 1				
440540	**Low discrimination**							**440540**
					NOV			
Staphylococcus auricularis		63.40	0.30	GLR 3	1			
Staphylococcus warneri		19.06	0.27	NIT 90	1			
Staphylococcus cohnii subsp. cohnii		17.54	0.22	GAL 10	99			
440600	**Low discrimination**							**440600**
Staphylococcus auricularis		72.34	0.41	GLR 3				
Staphylococcus warneri		19.35	0.38	NIT 90				
Staphylococcus carnosus subsp. utilis		6.36	0.03					
440610	**Low discrimination**							**440610**
					OXI	NOV		
Staphylococcus warneri		52.21	0.35	URE 98	1	1		
Staphylococcus pasteuri		44.77	0.19	MLT 95	1	1		
Macrococcus bovicus		3.03	-.02		99	99		
440640	**Low discrimination**							**440640**
					PYR			
Staphylococcus warneri		60.20	0.64	URE 98	8			
Staphylococcus auricularis		27.64	0.49		40			
Staphylococcus petrasii subsp. pragensis		3.49	0.13		99			
440641	**Low discrimination**							**440641**
					OXI	PYR		
Staphylococcus auricularis		68.14	0.29	GLR 3	1	40		
Staphylococcus warneri		27.26	0.29	NIT 90	1	8		
Kocuria kristinae		4.61	-.02	MNS 10	99	93		
440642	**Low discrimination**							**440642**
					PYR			
Staphylococcus warneri		76.93	0.54	URE 98	8			
Staphylococcus auricularis		11.78	0.29		40			
Staphylococcus haemolyticus		6.88	0.25		98			
440650	**Low discrimination**							**440650**
					PYR			
Staphylococcus warneri		74.98	0.61	URE 98	8			
Staphylococcus pasteuri		20.79	0.35		1			
Staphylococcus haemolyticus		3.22	0.26		98			
440652	**Low discrimination**							**440652**
					PYR			
Staphylococcus warneri		69.92	0.51	URE 98	8			
Staphylococcus pasteuri		23.75	0.27		1			
Staphylococcus haemolyticus		5.07	0.20		98			
440672	**Acceptable identification**							**440672**
					NOV	PYR		
Staphylococcus arlettae		94.42	0.37	ARG 1	99	1		

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		2.93	0.10		1	8		
Aerococcus viridans		2.66	0.16		1	95		
440673	**Good identification**							**440673**
					NOV	YEP		
Staphylococcus arlettae		95.28	0.40	ARG 1	99	99		
Aerococcus viridans		4.72	0.24		1	1		
440703	**Good identification**							**440703**
Staphylococcus hyicus		100.0	0.35	NIT 99				
440740	**Low discrimination**							**440740**
					PYR			
Staphylococcus warneri		74.03	0.54	URE 98	8			
Staphylococcus haemolyticus		14.00	0.31		98			
Staphylococcus auricularis		11.96	0.29		40			
440742	**Low discrimination**							**440742**
					PYR			
Staphylococcus warneri		72.83	0.44	URE 98	8			
Staphylococcus haemolyticus		23.25	0.26		98			
Staphylococcus auricularis		3.92	0.09		40			
440750	**Low discrimination**							**440750**
					OXI	PYR	NOV	
Staphylococcus warneri		85.98	0.51	URE 98	1	8	1	
Staphylococcus haemolyticus		13.19	0.27		1	98	2	
Macrococcus bovicus		0.83	-.02		99	1	99	
440752	**Low discrimination**							**440752**
					NOV	PYR		
Staphylococcus warneri		74.24	0.41	URE 98	1	8		
Staphylococcus haemolyticus		19.22	0.22		2	98		
Staphylococcus arlettae		4.39	0.12		99	1		
440772	**Good identification**							**440772**
					NOV	YEP		
Staphylococcus arlettae		99.51	0.53	ARG 1	99	99		
Aerococcus viridans		0.49	0.16		1	1		
440773	**Very good identification**							**440773**
					NOV	YEP		
Staphylococcus arlettae		99.13	0.55	ARG 1	99	99		
Aerococcus viridans		0.87	0.24		1	1		
441413	**Good identification**							**441413**
Aerococcus viridans		100.0	0.31	ARG 1				
441433	**Good identification**							**441433**
Aerococcus viridans		100.0	0.31	ARG 1				
441443	**Good identification**							**441443**
Aerococcus viridans		100.0	0.34	ARG 1				
441452	**Good identification**							**441452**
Aerococcus viridans		100.0	0.36	ARG 1				
441453	**Good identification**							**441453**
Aerococcus viridans		100.0	0.43	ARG 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
441463	**Good identification**						**441463**
Aerococcus viridans		100.0	0.34	ARG 1			
441472	**Good identification**						**441472**
Aerococcus viridans		100.0	0.36	ARG 1			
441473	**Good identification**						**441473**
Aerococcus viridans		100.0	0.43	ARG 1			
441513	**Good identification**						**441513**
Aerococcus viridans		100.0	0.31	ARG 1			
441533	**Good identification**						**441533**
Aerococcus viridans		100.0	0.31	ARG 1			
441543	**Good identification**						**441543**
Aerococcus viridans		100.0	0.34	ARG 1			
441552	**Low discrimination**						**441552**
					NOV	YEP	
Aerococcus viridans		88.97	0.36	ARG 1	1	1	
Staphylococcus kloosii		11.03	0.18		99	99	
441553	**Good identification**						**441553**
Aerococcus viridans		100.0	0.43	ARG 1			
441563	**Good identification**						**441563**
Aerococcus viridans		100.0	0.34	ARG 1			
441572	**Low discrimination**						**441572**
					NOV	YEP	
Aerococcus viridans		81.56	0.36	ARG 1	1	1	
Staphylococcus arlettae		18.44	0.12		99	99	
441573	**Low discrimination**						**441573**
					NOV	YEP	
Aerococcus viridans		88.62	0.43	ARG 1	1	1	
Staphylococcus arlettae		11.38	0.14		99	99	
441613	**Good identification**						**441613**
Aerococcus viridans		100.0	0.31	ARG 1			
441633	**Good identification**						**441633**
Aerococcus viridans		100.0	0.31	ARG 1			
441643	**Good identification**						**441643**
Aerococcus viridans		100.0	0.34	ARG 1			
441652	**Low discrimination**						**441652**
					PYR		
Aerococcus viridans		89.10	0.36	ARG 1	95		
Staphylococcus warneri		10.90	0.10		8		
441653	**Good identification**						**441653**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Aerococcus viridans		100.0	0.43	ARG 1			
441663	**Good identification**						**441663**
Aerococcus viridans		100.0	0.34	ARG 1			
441672	**Low discrimination**						**441672**
					NOV	YEP	
Staphylococcus arlettae		79.80	0.37	ARG 1	99	99	
Aerococcus viridans		20.20	0.36		1	1	
441673	**Low discrimination**						**441673**
					NOV	YEP	
Staphylococcus arlettae		69.18	0.40	ARG 1	99	99	
Aerococcus viridans		30.82	0.43		1	1	
441713	**Good identification**						**441713**
Aerococcus viridans		100.0	0.31	ARG 1			
441733	**Low discrimination**						**441733**
					NOV	YEP	
Aerococcus viridans		66.06	0.31	ARG 1	1	1	
Staphylococcus arlettae		33.94	0.14		99	99	
441743	**Good identification**						**441743**
Aerococcus viridans		100.0	0.34	ARG 1			
441752	**Low discrimination**						**441752**
					NOV	YEP	
Aerococcus viridans		81.56	0.36	ARG 1	1	1	
Staphylococcus arlettae		18.44	0.12		99	99	
441753	**Low discrimination**						**441753**
					NOV	YEP	
Aerococcus viridans		88.62	0.43	ARG 1	1	1	
Staphylococcus arlettae		11.38	0.14		99	99	
441763	**Low discrimination**						**441763**
					NOV	YEP	
Aerococcus viridans		72.18	0.34	ARG 1	1	1	
Staphylococcus arlettae		27.82	0.14		99	99	
441772	**Good identification**						**441772**
					NOV	YEP	
Staphylococcus arlettae		95.72	0.53	ARG 1	99	99	
Aerococcus viridans		4.28	0.36		1	1	
441773	**Good identification**						**441773**
					NOV	YEP	
Staphylococcus arlettae		92.71	0.55	ARG 1	99	99	
Aerococcus viridans		7.29	0.43		1	1	
442000	**Low discrimination**						**442000**
Kocuria rosea		47.40	0.53	ARG 1			
Staphylococcus carnosus subsp. utilis		47.33	0.44				
Staphylococcus auricularis		4.61	0.40				
442010	**Low discrimination**						**442010**
Kocuria rosea		87.62	0.33	ARG 1			
Staphylococcus carnosus subsp. utilis		7.96	0.03	MAN 10			
Staphylococcus capitis subsp. capitis		4.42	0.09				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
442011	**Good identification**					**442011**
Staphylococcus capitis subsp. capitis		98.96	0.50	GLR 1		
Kocuria rosea		1.04	0.07			
442020	**Good identification**					**442020**
Kocuria rosea		97.70	0.45	ARG 1		
Staphylococcus carnosus subsp. utilis		2.30	0.03			
442040	**Acceptable identification**					**442040**
Staphylococcus auricularis		90.98	0.47	GLR 3		
Kocuria rosea		4.05	0.12	TRE 92		
Staphylococcus carnosus subsp. utilis		4.04	0.03			
442041	**Good identification**					**442041**
Staphylococcus auricularis		100.0	0.27	GLR 3		
442042	**Good identification**					**442042**
Staphylococcus auricularis		95.07	0.27	GLR 3		
Staphylococcus haemolyticus		4.93	0.02	TRE 92		
442140	**Acceptable identification**					**442140**
Staphylococcus auricularis		90.58	0.27	GLR 3		
Staphylococcus haemolyticus		9.42	0.08	GAL 10		
442200	**Low discrimination**					**442200**
Kocuria rosea		52.08	0.33	ARG 1		
Staphylococcus auricularis		42.10	0.39	SUC 10		
Staphylococcus carnosus subsp. utilis		4.73	0.03			
442201	**Low discrimination**					**442201**
Staphylococcus capitis subsp. capitis		74.88	0.32	GLR 1		
Staphylococcus hyicus		9.85	0.11	MAN 93		
Staphylococcus auricularis		9.63	0.19			
442203	**Good identification**					**442203**
Staphylococcus hyicus		96.29	0.33	PHS 92		
Staphylococcus simulans		2.41	0.07	GAL 92		
Staphylococcus capitis subsp. capitis		1.30	-.03	TRE 92		
442211	**Very good identification**					**442211**
Staphylococcus capitis subsp. capitis		100.0	0.55	GLR 1		
442240	**Low discrimination**					**442240**
					PYR	
Staphylococcus haemolyticus		50.83	0.48	VPT 86	98	
Staphylococcus auricularis		46.27	0.46	TRE 93	40	
Staphylococcus warneri		2.90	0.30		8	
442241	**Low discrimination**					**442241**
Staphylococcus auricularis		76.58	0.27	GLR 3		
Staphylococcus haemolyticus		23.42	0.17	TRE 92		
442242	**Low discrimination**					**442242**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus haemolyticus		82.39	0.43	VPT 86	98	
Staphylococcus auricularis		14.82	0.27	TRE 93	40	
Staphylococcus warneri		2.79	0.20		8	
442250	**Acceptable identification**					**442250**
					PYR	
Staphylococcus haemolyticus		90.29	0.44	VPT 86	98	
Staphylococcus warneri		6.36	0.27	TRE 93	8	
Staphylococcus pasteuri		1.93	0.02		1	
442252	**Acceptable identification**					**442252**
					PYR	
Staphylococcus haemolyticus		94.12	0.38	VPT 86	98	
Staphylococcus warneri		3.93	0.17	TRE 93	8	
Staphylococcus simulans		1.95	0.01		70	
442301	**Good identification**					**442301**
Staphylococcus hyicus		100.0	0.33	PHS 92		
442303	**Very good identification**					**442303**
Staphylococcus hyicus		100.0	0.55	PHS 92		
442340	**Acceptable identification**					**442340**
					PYR	
Staphylococcus haemolyticus		90.44	0.49	VPT 86	98	
Staphylococcus auricularis		8.11	0.27	TRE 93	40	
Staphylococcus warneri		1.45	0.20		8	
442342	**Good identification**					**442342**
					PYR	
Staphylococcus haemolyticus		97.35	0.44	VPT 86	98	
Staphylococcus auricularis		1.73	0.07	TRE 93	40	
Staphylococcus warneri		0.92	0.10		8	
442350	**Good identification**					**442350**
					PYR	
Staphylococcus haemolyticus		98.07	0.45	VPT 86	98	
Staphylococcus warneri		1.94	0.17	TRE 93	8	
442352	**Good identification**					**442352**
Staphylococcus haemolyticus		100.0	0.39	VPT 86		
442400	**Low discrimination**					**442400**
Staphylococcus carnosus subsp. utilis		82.26	0.59	GLR 1		
Staphylococcus auricularis		17.56	0.61			
Kocuria rosea		0.16	0.12			
442401	**Low discrimination**					**442401**
					NOV	
Staphylococcus auricularis		69.32	0.42	GLR 3	1	
Staphylococcus carnosus subsp. utilis		29.53	0.18	MNS 10	1	
Staphylococcus cohnii subsp. cohnii		1.15	0.09		99	
442402	**Low discrimination**					**442402**
Staphylococcus auricularis		69.51	0.42	GLR 3		
Staphylococcus carnosus subsp. utilis		29.61	0.18	LAC 10		
Staphylococcus piscifermentans		0.89	0.12			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
442440	**Very good identification**					**442440**
Staphylococcus auricularis		96.20	0.69	GLR 3		
Staphylococcus carnosus subsp. utilis		1.95	0.18			
Staphylococcus haemolyticus		1.14	0.30			
442441	**Good identification**					**442441**
					NOV	
Staphylococcus auricularis		97.79	0.49	GLR 3	1	
Staphylococcus cohnii subsp. cohnii		2.21	0.19	MNS 10	99	
442442	**Acceptable identification**					**442442**
					PYR	
Staphylococcus auricularis		92.55	0.49	GLR 3	40	
Staphylococcus haemolyticus		5.54	0.25	LAC 10	98	
Staphylococcus warneri		1.40	0.20		8	
442443	**Good identification**					**442443**
Staphylococcus auricularis		100.0	0.30	GLR 3		
442450	**Low discrimination**					**442450**
					PYR	
Staphylococcus auricularis		39.98	0.28	GLR 3	40	
Staphylococcus haemolyticus		27.51	0.26	MAN 1	98	
Staphylococcus warneri		14.44	0.27		8	
442460	**Good identification**					**442460**
Staphylococcus auricularis		100.0	0.28	GLR 3		
442500	**Low discrimination**					**442500**
Staphylococcus auricularis		69.51	0.42	GLR 3		
Staphylococcus carnosus subsp. utilis		29.61	0.18	GAL 10		
Staphylococcus piscifermentans		0.89	0.12			
442503	**Good identification**					**442503**
Staphylococcus hyicus		99.03	0.35	PHS 92		
Staphylococcus auricularis		0.97	0.02	SUC 99		
442540	**Low discrimination**					**442540**
					PYR	
Staphylococcus auricularis		87.73	0.49	GLR 3	40	
Staphylococcus haemolyticus		10.54	0.31	GAL 10	98	
Staphylococcus warneri		1.26	0.20		8	
442541	**Good identification**					**442541**
Staphylococcus auricularis		100.0	0.30	GLR 3		
442542	**Low discrimination**					**442542**
Staphylococcus auricularis		57.97	0.30	GLR 3		
Staphylococcus haemolyticus		35.24	0.26	GAL 10		
Staphylococcus piscifermentans		4.29	0.16	LAC 10		
442550	**Low discrimination**					**442550**
					PYR	
Staphylococcus haemolyticus		77.50	0.27	VPT 86	98	
Staphylococcus warneri		11.40	0.17	SUC 99	8	
Staphylococcus auricularis		11.10	0.08		40	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
442600	**Good identification**						**442600**
Staphylococcus auricularis		90.68	0.61	GLR 3			
Staphylococcus carnosus subsp. utilis		4.65	0.18				
Staphylococcus haemolyticus		2.71	0.30				
442601	**Low discrimination**						**442601**
Staphylococcus hyicus		50.21	0.33	PHS 92			
Staphylococcus auricularis		49.11	0.41	GAL 92			
Staphylococcus capitis subsp. capitis		0.68	-.03	LAC 92			
442602	**Low discrimination**						**442602**
Staphylococcus auricularis		66.59	0.41	GLR 3			
Staphylococcus haemolyticus		10.08	0.25	LAC 10			
Staphylococcus simulans		9.69	0.22				
442603	**Very good identification**						**442603**
Staphylococcus hyicus		95.16	0.55	PHS 92			
Staphylococcus simulans		3.94	0.33	GAL 92			
Staphylococcus auricularis		0.90	0.21				
442610	**Low discrimination**						**442610**
					PYR	YEP	
Staphylococcus pasteuri		53.89	0.27	VPT 86	1	86	
Staphylococcus haemolyticus		20.95	0.26	URE 99	98	20	
Staphylococcus auricularis		12.05	0.20	MLT 86	40	1	
442612	**Low discrimination**						**442612**
					PYR	YEP	
Staphylococcus simulans		67.31	0.35	URE 98	70	1	
Staphylococcus pasteuri		19.21	0.19	BGA 90	1	86	
Staphylococcus haemolyticus		10.29	0.20		98	20	
442613	**Acceptable identification**						**442613**
Staphylococcus simulans		94.25	0.46	URE 98			
Staphylococcus hyicus		5.75	0.14	BGA 90			
442640	**Low discrimination**						**442640**
					PYR		
Staphylococcus haemolyticus		52.06	0.71	VPT 86	98		
Staphylococcus auricularis		41.03	0.68		40		
Staphylococcus warneri		4.26	0.56		8		
442641	**Low discrimination**						**442641**
					PYR		
Staphylococcus auricularis		72.87	0.49	GLR 3	40		
Staphylococcus haemolyticus		25.74	0.40	MNS 10	98		
Staphylococcus warneri		1.39	0.21		8		
442642	**Low discrimination**						**442642**
					PYR		
Staphylococcus haemolyticus		81.60	0.66	VPT 86	98		
Staphylococcus auricularis		12.70	0.49		40		
Staphylococcus warneri		3.95	0.46		8		
442643	**Low discrimination**						**442643**
Staphylococcus haemolyticus		43.12	0.35	VPT 86			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		24.11	0.29	MNS 3		
Staphylococcus simulans		20.35	0.26			
442650	**Low discrimination**					**442650**
					PYR	
Staphylococcus haemolyticus		78.50	0.67	VPT 86	98	
Staphylococcus pasteuri		12.53	0.43		1	
Staphylococcus warneri		7.91	0.53		8	
442651	**Low discrimination**					**442651**
					PYR	
Staphylococcus haemolyticus		85.67	0.36	VPT 86	98	
Staphylococcus warneri		5.70	0.19	MNS 3	8	
Staphylococcus pasteuri		4.47	0.02		1	
442652	**Low discrimination**					**442652**
					PYR	
Staphylococcus haemolyticus		82.78	0.62	VPT 86	98	
Staphylococcus pasteuri		9.59	0.35		1	
Staphylococcus warneri		4.94	0.44		8	
442653	**Low discrimination**					**442653**
					PYR	
Staphylococcus simulans		75.57	0.38	URE 98	70	
Staphylococcus haemolyticus		23.51	0.30	BGA 90	98	
Staphylococcus warneri		0.93	0.09		8	
442660	**Low discrimination**					**442660**
					PYR	
Staphylococcus haemolyticus		53.48	0.30	VPT 86	98	
Staphylococcus auricularis		42.15	0.27	XYL 1	40	
Staphylococcus warneri		4.37	0.15		8	
442700	**Low discrimination**					**442700**
					PYR	
Staphylococcus auricularis		67.60	0.41	GLR 3	40	
Staphylococcus haemolyticus		20.52	0.31	GAL 10	98	
Staphylococcus hyicus		8.03	0.14		1	
442701	**Very good identification**					**442701**
Staphylococcus hyicus		99.06	0.55	PHS 92		
Staphylococcus auricularis		0.94	0.21	LAC 92		
442702	**Low discrimination**					**442702**
					PYR	
Staphylococcus hyicus		80.17	0.35	PHS 92	1	
Staphylococcus haemolyticus		10.02	0.26	MNS 99	98	
Staphylococcus auricularis		6.52	0.21		40	
442703	**Very good identification**					**442703**
Staphylococcus hyicus		99.96	0.76	PHS 92		
Staphylococcus simulans		0.03	0.11			
Staphylococcus auricularis		0.01	0.02			
442710	**Low discrimination**					**442710**
					PYR	
Staphylococcus haemolyticus		87.18	0.27	VPT 86	98	
Staphylococcus warneri		12.82	0.17	MLT 99	8	
442713	**Low discrimination**					**442713**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus hyicus		88.97	0.35	PHS 92		
Staphylococcus simulans		11.04	0.24	MAN 1		
442723	**Good identification**					**442723**
Staphylococcus hyicus		100.0	0.35	PHS 92		
442740	**Good identification**					**442740**
					PYR	
Staphylococcus haemolyticus		90.81	0.73	VPT 86	98	
Staphylococcus auricularis		7.05	0.49		40	
Staphylococcus warneri		2.08	0.46		8	
442741	**Low discrimination**					**442741**
					PYR	
Staphylococcus haemolyticus		70.33	0.41	VPT 86	98	
Staphylococcus auricularis		19.62	0.29	MNS 3	40	
Staphylococcus hyicus		8.99	0.14		1	
442742	**Very good identification**					**442742**
					PYR	
Staphylococcus haemolyticus		96.94	0.67	VPT 86	98	
Staphylococcus auricularis		1.49	0.29		40	
Staphylococcus warneri		1.32	0.36		8	
442743	**Low discrimination**					**442743**
					PYR	
Staphylococcus hyicus		70.53	0.35	PHS 92	1	
Staphylococcus haemolyticus		27.00	0.36	MLT 1	98	
Staphylococcus auricularis		1.49	0.09		40	
442750	**Very good identification**					**442750**
					PYR	
Staphylococcus haemolyticus		96.92	0.68	VPT 86	98	
Staphylococcus warneri		2.74	0.43		8	
Staphylococcus pasteuri		0.14	0.02		1	
442751	**Good identification**					**442751**
Staphylococcus haemolyticus		100.0	0.37	VPT 86		
442752	**Very good identification**					**442752**
					PYR	
Staphylococcus haemolyticus		98.01	0.63	VPT 86	98	
Staphylococcus warneri		1.64	0.33		8	
Staphylococcus simulans		0.22	0.06		70	
442753	**Low discrimination**					**442753**
					PYR	
Staphylococcus haemolyticus		77.35	0.32	VPT 86	98	
Staphylococcus simulans		19.17	0.17	MNS 3	70	
Staphylococcus hyicus		3.48	-.06		1	
442760	**Acceptable identification**					**442760**
Staphylococcus haemolyticus		92.80	0.31	VPT 86		
Staphylococcus auricularis		7.21	0.08	XYL 1		
442770	**Good identification**					**442770**
Staphylococcus haemolyticus		100.0	0.27	VPT 86		
443402	**Acceptable identification**					**443402**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		91.94	0.32	URE 90		
Staphylococcus auricularis		8.06	0.01	GLR 1		
443440	**Low discrimination**					**443440**
Staphylococcus auricularis		73.42	0.28	GLR 3		
Staphylococcus piscifermentans		26.58	0.28	ESL 1		
443442	**Low discrimination**					**443442**
Staphylococcus piscifermentans		83.01	0.32	URE 90		
Staphylococcus auricularis		16.99	0.08	GLR 1		
443500	**Acceptable identification**					**443500**
Staphylococcus piscifermentans		91.94	0.32	URE 90		
Staphylococcus auricularis		8.06	0.01	GLR 1		
443502	**Good identification**					**443502**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
443512	**Good identification**					**443512**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
443540	**Low discrimination**					**443540**
Staphylococcus piscifermentans		83.01	0.32	URE 90		
Staphylococcus auricularis		16.99	0.08	GLR 1		
443542	**Good identification**					**443542**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
443552	**Good identification**					**443552**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
443600	**Low discrimination**					**443600**
Staphylococcus piscifermentans		57.85	0.32	URE 90		
Staphylococcus auricularis		42.15	0.20	GLR 1		
443602	**Good identification**					**443602**
Staphylococcus piscifermentans		100.0	0.36	URE 90		
443612	**Good identification**					**443612**
Staphylococcus piscifermentans		100.0	0.28	URE 90		
443640	**Low discrimination**					**443640**
Staphylococcus haemolyticus		42.86	0.30	VPT 86		
Staphylococcus auricularis		33.77	0.27	ESL 1		
Staphylococcus piscifermentans		19.87	0.32			
443642	**Low discrimination**					**443642**
Staphylococcus piscifermentans		51.68	0.36	URE 90		
Staphylococcus haemolyticus		41.81	0.25	GLR 1		
Staphylococcus auricularis		6.51	0.08			
443651	**Good identification**					**443651**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus sciuri		100.0	0.32	ARG 1			
443671	**Good identification**						**443671**
Staphylococcus sciuri		100.0	0.30	ARG 1			
443700	**Good identification**						**443700**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
443702	**Good identification**						**443702**
Staphylococcus piscifermentans		95.32	0.39	URE 90			
Staphylococcus hyicus		4.68	-.06	GLR 1			
443703	**Good identification**						**443703**
Staphylococcus hyicus		100.0	0.35	ESL 1			
443710	**Low discrimination**						**443710**
					OXI	NOV	
Staphylococcus piscifermentans		76.21	0.28	URE 90	1	1	
Staphylococcus sciuri		23.80	0.07	GLR 1	99	99	
443712	**Good identification**						**443712**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
443740	**Low discrimination**						**443740**
Staphylococcus haemolyticus		59.03	0.31	VPT 86			
Staphylococcus piscifermentans		36.39	0.36	ESL 1			
Staphylococcus auricularis		4.58	0.08				
443742	**Low discrimination**						**443742**
Staphylococcus piscifermentans		62.18	0.39	URE 90			
Staphylococcus haemolyticus		37.82	0.26	GLR 1			
443750	**Low discrimination**						**443750**
					OXI	NOV	
Staphylococcus haemolyticus		50.55	0.27	VPT 86	1	2	
Staphylococcus sciuri		26.71	0.19	ESL 1	99	99	
Staphylococcus piscifermentans		22.74	0.28		1	1	
443751	**Good identification**						**443751**
Staphylococcus sciuri		97.61	0.38	ARG 1			
Staphylococcus lentus		2.39	0.07	GLR 14			
443752	**Low discrimination**						**443752**
					OXI	NOV	
Staphylococcus piscifermentans		51.00	0.32	URE 90	1	1	
Staphylococcus haemolyticus		42.50	0.22	GLR 1	1	2	
Staphylococcus sciuri		6.50	0.03		99	99	
443771	**Acceptable identification**						**443771**
					OXI		
Staphylococcus sciuri		92.34	0.35	ARG 1	99		
Staphylococcus equorum		4.66	0.06	GLR 14	1		
Staphylococcus lentus		3.00	0.07		99		
444240	**Very good identification**						**444240**
Staphylococcus aureus subsp. anaerobius		100.0	0.59	GLR 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
444303	**Good identification**					**444303**
Staphylococcus hyicus		100.0	0.35	NIT 99		
444412	**Good identification**					**444412**
Staphylococcus kloosii		100.0	0.27	ARG 1		
444450	**Low discrimination**					**444450**
					YEP	
Staphylococcus kloosii		87.03	0.52	ARG 1	99	
Staphylococcus cohnii subsp. cohnii		12.97	0.23		12	
444452	**Very good identification**					**444452**
Staphylococcus kloosii		100.0	0.53	ARG 1		
444470	**Good identification**					**444470**
Staphylococcus kloosii		100.0	0.40	ARG 1		
444472	**Good identification**					**444472**
Staphylococcus kloosii		100.0	0.41	ARG 1		
444510	**Good identification**					**444510**
Staphylococcus kloosii		100.0	0.32	ARG 1		
444512	**Good identification**					**444512**
Staphylococcus kloosii		100.0	0.33	ARG 1		
444550	**Very good identification**					**444550**
Staphylococcus kloosii		100.0	0.58	ARG 1		
444552	**Very good identification**					**444552**
Staphylococcus kloosii		100.0	0.59	ARG 1		
444570	**Good identification**					**444570**
Staphylococcus kloosii		100.0	0.46	ARG 1		
444572	**Acceptable identification**					**444572**
Staphylococcus kloosii		90.04	0.47	ARG 1		
Staphylococcus arlettae		9.96	0.15			
444603	**Good identification**					**444603**
Staphylococcus hyicus		100.0	0.35	NIT 99		
444650	**Low discrimination**					**444650**
					NOV	
Staphylococcus kloosii		61.12	0.32	ARG 1	99	
Staphylococcus warneri		38.89	0.20	SUC 9	1	
444652	**Low discrimination**					**444652**
					NOV	
Staphylococcus kloosii		83.63	0.32	ARG 1	99	
Staphylococcus warneri		16.37	0.10	SUC 9	1	
444672	**Good identification**					**444672**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus arlettae		97.43	0.41	ARG 1		
Staphylococcus kloosii		2.57	0.20			
444673	**Good identification**					**444673**
Staphylococcus arlettae		100.0	0.43	ARG 1		
444701	**Good identification**					**444701**
Staphylococcus hyicus		100.0	0.35	NIT 99		
444703	**Very good identification**					**444703**
Staphylococcus hyicus		100.0	0.57	NIT 99		
444750	**Acceptable identification**					**444750**
					NOV	
Staphylococcus kloosii		90.62	0.38	ARG 1	99	
Staphylococcus warneri		9.38	0.10	SUC 9	1	
444752	**Low discrimination**					**444752**
Staphylococcus kloosii		78.16	0.38	ARG 1		
Staphylococcus arlettae		21.85	0.15	SUC 9		
444772	**Very good identification**					**444772**
Staphylococcus arlettae		99.11	0.56	ARG 1		
Staphylococcus kloosii		0.90	0.26			
444773	**Very good identification**					**444773**
Staphylococcus arlettae		100.0	0.59	ARG 1		
445412	**Good identification**					**445412**
Staphylococcus kloosii		100.0	0.27	ARG 1		
445450	**Good identification**					**445450**
Staphylococcus kloosii		100.0	0.52	ARG 1		
445452	**Very good identification**					**445452**
Staphylococcus kloosii		100.0	0.53	ARG 1		
445470	**Good identification**					**445470**
Staphylococcus kloosii		100.0	0.40	ARG 1		
445472	**Good identification**					**445472**
Staphylococcus kloosii		100.0	0.41	ARG 1		
445510	**Good identification**					**445510**
Staphylococcus kloosii		100.0	0.32	ARG 1		
445512	**Good identification**					**445512**
Staphylococcus kloosii		100.0	0.33	ARG 1		
445550	**Very good identification**					**445550**
Staphylococcus kloosii		100.0	0.58	ARG 1		
445552	**Very good identification**					**445552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		100.0	0.59	ARG 1		
445570	**Good identification**					**445570**
Staphylococcus kloosii		100.0	0.46	ARG 1		
445572	**Acceptable identification**					**445572**
Staphylococcus kloosii		90.04	0.47	ARG 1		
Staphylococcus arlettae		9.96	0.15			
445650	**Good identification**					**445650**
Staphylococcus kloosii		100.0	0.32	ARG 1		
445652	**Good identification**					**445652**
Staphylococcus kloosii		100.0	0.32	ARG 1		
445672	**Good identification**					**445672**
Staphylococcus arlettae		97.43	0.41	ARG 1		
Staphylococcus kloosii		2.57	0.20			
445673	**Good identification**					**445673**
Staphylococcus arlettae		100.0	0.43	ARG 1		
445750	**Good identification**					**445750**
Staphylococcus kloosii		100.0	0.38	ARG 1		
445752	**Low discrimination**					**445752**
Staphylococcus kloosii		78.16	0.38	ARG 1		
Staphylococcus arlettae		21.85	0.15	SUC 9		
445772	**Very good identification**					**445772**
Staphylococcus arlettae		99.11	0.56	ARG 1		
Staphylococcus kloosii		0.90	0.26			
445773	**Very good identification**					**445773**
Staphylococcus arlettae		99.71	0.59	ARG 1		
Staphylococcus gallinarum		0.29	-.04			
446011	**Low discrimination**					**446011**
Staphylococcus capitis subsp. capitis		73.13	0.31	GLR 1		
Staphylococcus carnosus subsp. carnosus		26.87	0.23	PHS 11		
446013	**Good identification**					**446013**
Staphylococcus carnosus subsp. carnosus		98.63	0.36	BGA 90		
Staphylococcus capitis subsp. capitis		1.37	-.04	GLR 1		
446103	**Good identification**					**446103**
Staphylococcus hyicus		96.14	0.35	SUC 99		
Staphylococcus carnosus subsp. carnosus		3.86	0.11	TRE 92		
446201	**Acceptable identification**					**446201**
Staphylococcus hyicus		92.45	0.33	GAL 92		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. capitis		7.56	0.14	TRE 92		
446203	**Very good identification**					**446203**
Staphylococcus hyicus		100.0	0.55	GAL 92		
446211	**Good identification**					**446211**
Staphylococcus capitis subsp. capitis		100.0	0.37	GLR 1		
446301	**Very good identification**					**446301**
Staphylococcus hyicus		100.0	0.55	TRE 92		
446302	**Good identification**					**446302**
Staphylococcus hyicus		100.0	0.35	TRE 92		
446303	**Very good identification**					**446303**
					YEP	
Staphylococcus hyicus		99.99	0.76	TRE 92	1	
Staphylococcus chromogenes		0.01	-.04		95	
446313	**Good identification**					**446313**
Staphylococcus hyicus		100.0	0.35	TRE 92		
446323	**Good identification**					**446323**
Staphylococcus hyicus		100.0	0.35	TRE 92		
446343	**Good identification**					**446343**
					YEP	
Staphylococcus hyicus		97.12	0.35	TRE 92	1	
Staphylococcus epidermidis		2.10	0.09	MLT 1	1	
Staphylococcus chromogenes		0.78	-.04		95	
446403	**Acceptable identification**					**446403**
Staphylococcus hyicus		93.55	0.35	GAL 92		
Staphylococcus carnosus subsp. carnosus		6.45	0.16	SUC 99		
446413	**Acceptable identification**					**446413**
Staphylococcus carnosus subsp. carnosus		90.21	0.28	BGA 90		
Staphylococcus simulans		6.48	0.07	GLR 1		
Staphylococcus hyicus		3.31	-.06			
446440	**Good identification**					**446440**
Staphylococcus auricularis		100.0	0.28	GLR 3		
446501	**Acceptable identification**					**446501**
Staphylococcus hyicus		94.58	0.35	SUC 99		
Staphylococcus schleiferi subsp. schleiferi		5.42	0.14	LAC 92		
446503	**Very good identification**					**446503**
Staphylococcus hyicus		99.85	0.57	SUC 99		
Staphylococcus carnosus subsp. carnosus		0.15	0.03			
446601	**Very good identification**					**446601**
Staphylococcus hyicus		100.0	0.55	GAL 92		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
446602	**Good identification**					**446602**
Staphylococcus hyicus		95.63	0.35	GAL 92		
Staphylococcus simulans		3.40	0.13	MNS 99		
Staphylococcus piscifermentans		0.97	0.12			
446603	**Very good identification**					**446603**
					YEP	
Staphylococcus hyicus		99.87	0.76	GAL 92	1	
Staphylococcus simulans		0.12	0.23		1	
Staphylococcus chromogenes		0.01	-.02		95	
446613	**Low discrimination**					**446613**
Staphylococcus hyicus		67.36	0.35	GAL 92		
Staphylococcus simulans		32.03	0.36	MAN 1		
Staphylococcus felis		0.62	-.09			
446620	**Good identification**					**446620**
Staphylococcus muscae		100.0	0.39	ARG 1		
446623	**Good identification**					**446623**
Staphylococcus hyicus		100.0	0.35	GAL 92		
446640	**Low discrimination**					**446640**
					PYR	
Staphylococcus haemolyticus		53.48	0.30	VPT 86	98	
Staphylococcus auricularis		42.15	0.27	PHS 1	40	
Staphylococcus warneri		4.37	0.15		8	
446643	**Acceptable identification**					**446643**
					YEP	
Staphylococcus hyicus		94.02	0.35	GAL 92	1	
Staphylococcus simulans		5.02	0.16	MLT 1	1	
Staphylococcus chromogenes		0.96	-.02		95	
446653	**Good identification**					**446653**
Staphylococcus simulans		95.49	0.29	URE 98		
Staphylococcus hyicus		4.52	-.06	BGA 90		
446700	**Good identification**					**446700**
Staphylococcus hyicus		99.00	0.35	MNS 99		
Staphylococcus piscifermentans		1.01	0.12	LAC 92		
446701	**Very good identification**					**446701**
					YEP	
Staphylococcus hyicus		99.99	0.76	LAC 92	1	
Staphylococcus chromogenes		0.01	-.02		95	
446702	**Very good identification**					**446702**
Staphylococcus hyicus		99.87	0.57	MNS 99		
Staphylococcus piscifermentans		0.13	0.16			
446703	**Excellent identification**					**446703**
					YEP	
Staphylococcus hyicus		99.99	0.98		1	
Staphylococcus chromogenes		0.01	0.18		95	
Staphylococcus simulans		0.00	0.02		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
446711	**Good identification**							**446711**
Staphylococcus hyicus		100.0	0.35	MAN 1				
446713	**Very good identification**							**446713**
Staphylococcus hyicus		99.03	0.57	MAN 1				
Staphylococcus simulans		0.36	0.14					
Staphylococcus felis		0.26	0.02					
446721	**Good identification**							**446721**
Staphylococcus hyicus		100.0	0.35	XYL 1				
446723	**Very good identification**							**446723**
Staphylococcus hyicus		100.0	0.57	XYL 1				
446740	**Low discrimination**							**446740**
Staphylococcus haemolyticus		85.56	0.31	VPT 86				
Staphylococcus auricularis		6.64	0.08	PHS 1				
Staphylococcus piscifermentans		3.91	0.12					
446741	**Good identification**							**446741**
					YEP			
Staphylococcus hyicus		98.99	0.35	MLT 1	1			
Staphylococcus chromogenes		1.01	-.02	LAC 92	95			
446743	**Very good identification**							**446743**
					YEP	PYR	COA	
Staphylococcus hyicus		99.14	0.57	MLT 1	1	1	24	
Staphylococcus chromogenes		0.80	0.18		95	45	1	
Staphylococcus intermedius		0.07	-.03		1	90	99	
446750	**Good identification**							**446750**
Staphylococcus haemolyticus		100.0	0.27	VPT 86				
447303	**Good identification**							**447303**
Staphylococcus hyicus		100.0	0.35	ESL 1				
447402	**Good identification**							**447402**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
447442	**Good identification**							**447442**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
447500	**Good identification**							**447500**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
447502	**Good identification**							**447502**
Staphylococcus piscifermentans		100.0	0.32	URE 90				
447540	**Good identification**							**447540**
Staphylococcus piscifermentans		100.0	0.28	URE 90				
447542	**Good identification**							**447542**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		100.0	0.32	URE 90			
447600	**Good identification**						**447600**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
447602	**Acceptable identification**						**447602**
Staphylococcus piscifermentans		90.04	0.32	URE 90			
Staphylococcus hyicus		9.96	-.06	GLR 1			
447603	**Good identification**						**447603**
Staphylococcus hyicus		100.0	0.35	ESL 1			
447640	**Good identification**						**447640**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
447642	**Good identification**						**447642**
Staphylococcus piscifermentans		100.0	0.32	URE 90			
447651	**Good identification**						**447651**
Staphylococcus sciuri		100.0	0.37	ARG 1			
447671	**Good identification**						**447671**
Staphylococcus sciuri		100.0	0.34	ARG 1			
447700	**Acceptable identification**						**447700**
Staphylococcus piscifermentans		90.04	0.32	URE 90			
Staphylococcus hyicus		9.96	-.06	GLR 1			
447701	**Good identification**						**447701**
					OXI	NOV	
Staphylococcus hyicus		99.23	0.35	ESL 1	1	1	
Staphylococcus sciuri		0.78	0.00	LAC 92	99	99	
447702	**Low discrimination**						**447702**
Staphylococcus piscifermentans		54.12	0.36	URE 90			
Staphylococcus hyicus		45.88	0.16	GLR 1			
447703	**Very good identification**						**447703**
Staphylococcus hyicus		100.0	0.57	ESL 1			
447711	**Good identification**						**447711**
					OXI	NOV	
Staphylococcus sciuri		96.16	0.31	ARG 1	99	99	
Staphylococcus hyicus		3.84	-.06	GLR 14	1	1	
447712	**Good identification**						**447712**
Staphylococcus piscifermentans		100.0	0.28	URE 90			
447731	**Good identification**						**447731**
Staphylococcus sciuri		100.0	0.28	ARG 1			
447740	**Good identification**						**447740**
Staphylococcus piscifermentans		100.0	0.32	URE 90			

Code		ID	T	tests	Differentiation tests		Code
Taxon		score	index	against			
447742	**Good identification**						**447742**
Staphylococcus piscifermentans		100.0	0.36	URE 90			
447751	**Good identification**						**447751**
Staphylococcus sciuri		100.0	0.43	ARG 1			
447752	**Low discrimination**						**447752**
					OXI	NOV	
Staphylococcus piscifermentans		75.44	0.28	URE 90	1	1	
Staphylococcus sciuri		24.56	0.08	GLR 1	99	99	
447753	**Low discrimination**						**447753**
					OXI		
Staphylococcus sciuri		87.62	0.27	ARG 1	99		
Staphylococcus lentus		6.86	0.05	GLR 14	99		
Staphylococcus gallinarum		5.52	-.04	LAC 14	1		
447771	**Low discrimination**						**447771**
					OXI		
Staphylococcus sciuri		62.74	0.40	ARG 1	99		
Staphylococcus gallinarum		34.47	0.29	GLR 14	1		
Staphylococcus equorum		2.79	0.09		1		
447773	**Low discrimination**						**447773**
					OXI		
Staphylococcus gallinarum		79.80	0.37	URE 92	1		
Staphylococcus sciuri		9.66	0.24	ARG 1	99		
Staphylococcus equorum		6.79	0.18		1		
452440	**Good identification**						**452440**
Staphylococcus auricularis		100.0	0.28	ORN 1			
452640	**Low discrimination**						**452640**
					PYR		
Staphylococcus haemolyticus		53.48	0.30	VPT 86	98		
Staphylococcus auricularis		42.15	0.27	ORN 1	40		
Staphylococcus warneri		4.37	0.15		8		
452703	**Good identification**						**452703**
Staphylococcus hyicus		100.0	0.35	ORN 1			
452740	**Acceptable identification**						**452740**
Staphylococcus haemolyticus		92.80	0.31	VPT 86			
Staphylococcus auricularis		7.21	0.08	ORN 1			
452750	**Good identification**						**452750**
Staphylococcus haemolyticus		100.0	0.27	VPT 86			
456303	**Good identification**						**456303**
Staphylococcus hyicus		100.0	0.35	ORN 1			
456603	**Good identification**						**456603**
Staphylococcus hyicus		100.0	0.35	ORN 1			
456701	**Good identification**						**456701**
Staphylococcus hyicus		100.0	0.35	ORN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
456703	**Very good identification**						**456703**
Staphylococcus hyicus		100.0	0.57	ORN 1			
460440	**Low discrimination**						**460440**
Staphylococcus auricularis		86.02	0.33	BGA 14			
Staphylococcus warneri		13.98	0.25	GLR 3			
460613	**Good identification**						**460613**
Staphylococcus simulans		100.0	0.37	URE 98			
460640	**Low discrimination**						**460640**
					PYR		
Staphylococcus warneri		73.97	0.51	URE 98	8		
Staphylococcus auricularis		22.12	0.32		40		
Staphylococcus haemolyticus		3.92	0.18		98		
460642	**Low discrimination**						**460642**
					PYR		
Staphylococcus warneri		80.20	0.42	URE 98	8		
Staphylococcus auricularis		7.99	0.13		40		
Staphylococcus haemolyticus		7.17	0.13		98		
460650	**Good identification**						**460650**
					PYR		
Staphylococcus warneri		95.88	0.49	URE 98	8		
Staphylococcus haemolyticus		4.12	0.13		98		
460652	**Low discrimination**						**460652**
					PYR		
Staphylococcus warneri		71.83	0.39	URE 98	8		
Staphylococcus simulans		22.97	0.19		70		
Staphylococcus haemolyticus		5.21	0.08		98		
460653	**Acceptable identification**						**460653**
Staphylococcus simulans		93.86	0.30	URE 98			
Aerococcus viridans		6.14	0.22	NIT 96			
460672	**Good identification**						**460672**
Staphylococcus arlettae		100.0	0.37	ARG 1			
460673	**Good identification**						**460673**
					NOV	YEP	
Staphylococcus arlettae		96.11	0.40	ARG 1	99	99	
Aerococcus viridans		3.89	0.22		1	1	
460740	**Low discrimination**						**460740**
					PYR		
Staphylococcus warneri		77.26	0.41	URE 98	8		
Staphylococcus haemolyticus		14.61	0.19		98		
Staphylococcus auricularis		8.13	0.13		40		
460742	**Low discrimination**						**460742**
					PYR		
Staphylococcus warneri		75.80	0.31	URE 98	8		
Staphylococcus haemolyticus		24.20	0.14		98		
460750	**Low discrimination**						**460750**
					PYR		
Staphylococcus warneri		86.70	0.38	URE 98	8		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		13.30	0.14		98		
460752	**Low discrimination**						**460752**
					NOV	PYR	
Staphylococcus warneri		66.88	0.28	URE 98	1	8	
Staphylococcus haemolyticus		17.31	0.09		2	98	
Staphylococcus arlettae		15.81	0.12		99	1	
460772	**Good identification**						**460772**
Staphylococcus arlettae		100.0	0.53	ARG 1			
460773	**Very good identification**						**460773**
					NOV	YEP	
Staphylococcus arlettae		98.76	0.55	ARG 1	99	99	
Aerococcus viridans		0.71	0.22		1	1	
Staphylococcus xylosus		0.53	-.01		99	40	
461413	**Good identification**						**461413**
Aerococcus viridans		100.0	0.29	ARG 1			
461433	**Good identification**						**461433**
Aerococcus viridans		100.0	0.29	ARG 1			
461443	**Good identification**						**461443**
Aerococcus viridans		100.0	0.32	ARG 1			
461452	**Good identification**						**461452**
Aerococcus viridans		100.0	0.34	ARG 1			
461453	**Acceptable identification**						**461453**
					NOV		
Aerococcus viridans		94.07	0.42	ARG 1	1		
Staphylococcus cohnii subsp. urealyticum		5.93	0.06		99		
461463	**Good identification**						**461463**
Aerococcus viridans		100.0	0.32	ARG 1			
461472	**Good identification**						**461472**
Aerococcus viridans		100.0	0.34	ARG 1			
461473	**Good identification**						**461473**
Aerococcus viridans		100.0	0.42	ARG 1			
461513	**Good identification**						**461513**
Aerococcus viridans		100.0	0.29	ARG 1			
461533	**Good identification**						**461533**
Aerococcus viridans		100.0	0.29	ARG 1			
461543	**Good identification**						**461543**
Aerococcus viridans		100.0	0.32	ARG 1			
461552	**Low discrimination**						**461552**
					NOV	YEP	
Aerococcus viridans		86.84	0.34	ARG 1	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		13.16	0.18		99	99	
461553	**Good identification**						**461553**
Aerococcus viridans		100.0	0.42	ARG 1			
461563	**Good identification**						**461563**
Aerococcus viridans		100.0	0.32	ARG 1			
461572	**Low discrimination**						**461572**
					NOV	YEP	
Aerococcus viridans		78.35	0.34	ARG 1	1	1	
Staphylococcus arlettae		21.65	0.12		99	99	
461573	**Low discrimination**						**461573**
					NOV	YEP	
Aerococcus viridans		86.43	0.42	ARG 1	1	1	
Staphylococcus arlettae		13.57	0.14		99	99	
461613	**Good identification**						**461613**
Aerococcus viridans		100.0	0.29	ARG 1			
461633	**Good identification**						**461633**
Aerococcus viridans		100.0	0.29	ARG 1			
461643	**Good identification**						**461643**
Aerococcus viridans		100.0	0.32	ARG 1			
461652	**Good identification**						**461652**
Aerococcus viridans		100.0	0.34	ARG 1			
461653	**Good identification**						**461653**
Aerococcus viridans		100.0	0.42	ARG 1			
461663	**Good identification**						**461663**
Aerococcus viridans		100.0	0.32	ARG 1			
461672	**Low discrimination**						**461672**
					NOV	YEP	
Staphylococcus arlettae		82.84	0.37	ARG 1	99	99	
Aerococcus viridans		17.16	0.34		1	1	
461673	**Low discrimination**						**461673**
					NOV	YEP	
Staphylococcus arlettae		73.29	0.40	ARG 1	99	99	
Aerococcus viridans		26.72	0.42		1	1	
461713	**Good identification**						**461713**
Aerococcus viridans		100.0	0.29	ARG 1			
461733	**Low discrimination**						**461733**
					NOV	YEP	
Aerococcus viridans		61.42	0.29	ARG 1	1	1	
Staphylococcus arlettae		38.58	0.14		99	99	
461743	**Good identification**						**461743**
Aerococcus viridans		100.0	0.32	ARG 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
461752	**Low discrimination**						**461752**
					NOV	YEP	
Aerococcus viridans		78.35	0.34	ARG 1	1	1	
Staphylococcus arlettae		21.65	0.12		99	99	
461753	**Low discrimination**						**461753**
					NOV	YEP	
Aerococcus viridans		86.43	0.42	ARG 1	1	1	
Staphylococcus arlettae		13.57	0.14		99	99	
461763	**Low discrimination**						**461763**
					NOV	YEP	
Aerococcus viridans		67.98	0.32	ARG 1	1	1	
Staphylococcus arlettae		32.02	0.14		99	99	
461772	**Good identification**						**461772**
					NOV	YEP	
Staphylococcus arlettae		96.47	0.53	ARG 1	99	99	
Aerococcus viridans		3.53	0.34		1	1	
461773	**Good identification**						**461773**
					NOV	YEP	
Staphylococcus arlettae		93.96	0.55	ARG 1	99	99	
Aerococcus viridans		6.04	0.42		1	1	
462040	**Good identification**						**462040**
Staphylococcus auricularis		100.0	0.31	BGA 14			
462203	**Low discrimination**						**462203**
Staphylococcus simulans		67.01	0.27	URE 98			
Staphylococcus hyicus		32.99	0.13	TRE 95			
462212	**Good identification**						**462212**
Staphylococcus simulans		100.0	0.28	URE 98			
462213	**Good identification**						**462213**
Staphylococcus simulans		100.0	0.39	URE 98			
462240	**Low discrimination**						**462240**
					PYR		
Staphylococcus haemolyticus		60.61	0.36	VPT 86	98		
Staphylococcus auricularis		35.93	0.30	TRE 93	40		
Staphylococcus warneri		3.46	0.18		8		
462242	**Low discrimination**						**462242**
Staphylococcus haemolyticus		81.51	0.31	VPT 86			
Staphylococcus auricularis		9.55	0.11	TRE 93			
Staphylococcus simulans		8.95	0.09				
462250	**Acceptable identification**						**462250**
					PYR		
Staphylococcus haemolyticus		93.42	0.31	VPT 86	98		
Staphylococcus warneri		6.58	0.15	TRE 93	8		
462253	**Good identification**						**462253**
Staphylococcus simulans		100.0	0.32	URE 98			
462303	**Good identification**						**462303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus hyicus		98.49	0.35	BGA 10		
Staphylococcus simulans		1.51	0.05	PHS 92		
462340	**Acceptable identification**					**462340**
Staphylococcus haemolyticus		94.48	0.37	VPT 86		
Staphylococcus auricularis		5.52	0.11	TRE 93		
462342	**Good identification**					**462342**
Staphylococcus haemolyticus		100.0	0.32	VPT 86		
462350	**Good identification**					**462350**
Staphylococcus haemolyticus		100.0	0.32	VPT 86		
462352	**Acceptable identification**					**462352**
Staphylococcus haemolyticus		94.55	0.27	VPT 86		
Staphylococcus simulans		5.45	-.01	TRE 93		
462400	**Low discrimination**					**462400**
Staphylococcus auricularis		77.48	0.45	BGA 14		
Staphylococcus carnosus subsp. utilis		22.52	0.18	GLR 3		
462413	**Good identification**					**462413**
Staphylococcus simulans		95.53	0.37	URE 98		
Staphylococcus carnosus subsp. carnosus		4.48	0.07	SUC 96		
462440	**Good identification**					**462440**
					PYR	
Staphylococcus auricularis		97.48	0.53	BGA 14	40	
Staphylococcus haemolyticus		1.77	0.18	GLR 3	98	
Staphylococcus warneri		0.75	0.18		8	
462441	**Good identification**					**462441**
Staphylococcus auricularis		100.0	0.33	BGA 14		
462442	**Low discrimination**					**462442**
Staphylococcus auricularis		86.84	0.33	BGA 14		
Staphylococcus haemolyticus		7.99	0.13	GLR 3		
Staphylococcus simulans		5.17	0.06	LAC 10		
462453	**Good identification**					**462453**
Staphylococcus simulans		100.0	0.30	URE 98		
462540	**Low discrimination**					**462540**
Staphylococcus auricularis		84.43	0.33	BGA 14		
Staphylococcus haemolyticus		15.57	0.19	GLR 3		
462600	**Acceptable identification**					**462600**
					PYR	
Staphylococcus auricularis		93.07	0.44	BGA 14	40	
Staphylococcus haemolyticus		4.28	0.18	GLR 3	98	
Staphylococcus warneri		1.82	0.18		8	
462602	**Low discrimination**					**462602**
Staphylococcus simulans		85.96	0.42	URE 98		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		10.69	0.25			
Staphylococcus haemolyticus		2.49	0.13			
462603	**Low discrimination**					**462603**
Staphylococcus simulans		76.80	0.53	URE 98		
Staphylococcus hyicus		22.89	0.35			
Staphylococcus auricularis		0.32	0.05			
462612	**Very good identification**					**462612**
Staphylococcus simulans		99.58	0.54	URE 98		
Staphylococcus haemolyticus		0.42	0.08			
462613	**Very good identification**					**462613**
Staphylococcus simulans		99.93	0.65	URE 98		
Staphylococcus hyicus		0.08	-.06			
462640	**Low discrimination**					**462640**
					PYR	
Staphylococcus haemolyticus		62.70	0.59	VPT 86	98	
Staphylococcus auricularis		32.17	0.52		40	
Staphylococcus warneri		5.12	0.44		8	
462641	**Low discrimination**					**462641**
Staphylococcus auricularis		61.57	0.32	BGA 14		
Staphylococcus haemolyticus		33.40	0.28	GLR 3		
Staphylococcus simulans		3.23	0.05	MNS 10		
462642	**Low discrimination**					**462642**
Staphylococcus haemolyticus		76.53	0.54	VPT 86		
Staphylococcus simulans		12.01	0.35			
Staphylococcus auricularis		7.76	0.32			
462643	**Acceptable identification**					**462643**
Staphylococcus simulans		92.00	0.46	URE 98		
Staphylococcus haemolyticus		5.41	0.23			
Staphylococcus auricularis		1.97	0.13			
462650	**Low discrimination**					**462650**
					PYR	
Staphylococcus haemolyticus		89.12	0.54	VPT 86	98	
Staphylococcus warneri		8.98	0.41		8	
Staphylococcus auricularis		0.79	0.11		40	
462652	**Low discrimination**					**462652**
					PYR	
Staphylococcus simulans		50.22	0.47	URE 98	70	
Staphylococcus haemolyticus		46.98	0.49		98	
Staphylococcus warneri		2.81	0.31		8	
462653	**Very good identification**					**462653**
Staphylococcus simulans		99.14	0.58	URE 98		
Staphylococcus haemolyticus		0.86	0.18			
462701	**Good identification**					**462701**
Staphylococcus hyicus		98.63	0.35	BGA 10		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		1.37	0.05	PHS 92			
462703	**Very good identification**						**462703**
					COA	PYR	
Staphylococcus hyicus		97.38	0.57	BGA 10	24	1	
Staphylococcus simulans		2.47	0.31	PHS 92	1	70	
Staphylococcus intermedius		0.15	0.04		99	90	
462712	**Acceptable identification**						**462712**
Staphylococcus simulans		94.78	0.33	URE 98			
Staphylococcus haemolyticus		5.22	0.09	GAL 8			
462713	**Low discrimination**						**462713**
					COA	PYR	
Staphylococcus simulans		87.78	0.43	URE 98	1	70	
Staphylococcus hyicus		8.74	0.16	GAL 8	24	1	
Staphylococcus intermedius		3.48	0.13		99	90	
462740	**Good identification**						**462740**
					PYR		
Staphylococcus haemolyticus		93.16	0.60	VPT 86	98		
Staphylococcus auricularis		4.71	0.32		40		
Staphylococcus warneri		2.13	0.34		8		
462741	**Low discrimination**						**462741**
					PYR		
Staphylococcus haemolyticus		80.75	0.29	VPT 86	98		
Staphylococcus auricularis		14.67	0.13	MNS 3	40		
Staphylococcus hyicus		4.59	-.06		1		
462742	**Very good identification**						**462742**
					PYR		
Staphylococcus haemolyticus		96.56	0.55	VPT 86	98		
Staphylococcus warneri		1.31	0.24		8		
Staphylococcus simulans		1.17	0.13		70		
462750	**Very good identification**						**462750**
					NOV	PYR	
Staphylococcus haemolyticus		96.79	0.55	VPT 86	2	98	
Staphylococcus warneri		2.73	0.31		1	8	
Staphylococcus saprophyticus subsp. bovis		0.48	0.05		99	99	
462752	**Acceptable identification**						**462752**
					PYR		
Staphylococcus haemolyticus		90.98	0.50	VPT 86	98		
Staphylococcus simulans		7.50	0.25		70		
Staphylococcus warneri		1.52	0.21		8		
462753	**Low discrimination**						**462753**
					COA		
Staphylococcus simulans		79.29	0.36	URE 98	1		
Staphylococcus intermedius		10.50	0.16	GAL 8	99		
Staphylococcus haemolyticus		8.89	0.19		1		
464412	**Good identification**						**464412**
Staphylococcus kloosii		100.0	0.27	ARG 1			
464450	**Good identification**						**464450**
Staphylococcus kloosii		100.0	0.52	ARG 1			
464452	**Very good identification**						**464452**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.53	ARG 1			
464470	**Good identification**						**464470**
Staphylococcus kloosii		100.0	0.40	ARG 1			
464472	**Good identification**						**464472**
Staphylococcus kloosii		100.0	0.41	ARG 1			
464510	**Good identification**						**464510**
Staphylococcus kloosii		100.0	0.32	ARG 1			
464512	**Good identification**						**464512**
Staphylococcus kloosii		100.0	0.33	ARG 1			
464550	**Very good identification**						**464550**
Staphylococcus kloosii		100.0	0.58	ARG 1			
464552	**Very good identification**						**464552**
Staphylococcus kloosii		100.0	0.59	ARG 1			
464570	**Good identification**						**464570**
Staphylococcus kloosii		100.0	0.46	ARG 1			
464572	**Acceptable identification**						**464572**
Staphylococcus kloosii		90.04	0.47	ARG 1			
Staphylococcus arlettae		9.96	0.15				
464613	**Good identification**						**464613**
Staphylococcus simulans		100.0	0.27	URE 98			
464650	**Good identification**						**464650**
Staphylococcus kloosii		100.0	0.32	ARG 1			
464652	**Low discrimination**						**464652**
					NOV	YEP	
Staphylococcus kloosii		65.95	0.32	ARG 1	99	99	
Staphylococcus simulans		34.05	0.09	SUC 9	1	1	
464672	**Good identification**						**464672**
Staphylococcus arlettae		97.43	0.41	ARG 1			
Staphylococcus kloosii		2.57	0.20				
464673	**Good identification**						**464673**
Staphylococcus arlettae		100.0	0.43	ARG 1			
464703	**Good identification**						**464703**
					PYR		
Staphylococcus hyicus		96.97	0.38	BGA 10	1		
Staphylococcus intermedius		3.03	0.12	NIT 99	90		
464750	**Good identification**						**464750**
Staphylococcus kloosii		100.0	0.38	ARG 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
464752	**Low discrimination**					**464752**
Staphylococcus kloosii		78.16	0.38	ARG 1		
Staphylococcus arlettae		21.85	0.15	SUC 9		
464772	**Very good identification**					**464772**
Staphylococcus arlettae		99.11	0.56	ARG 1		
Staphylococcus kloosii		0.90	0.26			
464773	**Very good identification**					**464773**
Staphylococcus arlettae		100.0	0.59	ARG 1		
465412	**Good identification**					**465412**
Staphylococcus kloosii		100.0	0.27	ARG 1		
465450	**Good identification**					**465450**
Staphylococcus kloosii		100.0	0.52	ARG 1		
465452	**Very good identification**					**465452**
Staphylococcus kloosii		100.0	0.53	ARG 1		
465470	**Good identification**					**465470**
Staphylococcus kloosii		100.0	0.40	ARG 1		
465472	**Good identification**					**465472**
Staphylococcus kloosii		100.0	0.41	ARG 1		
465510	**Good identification**					**465510**
Staphylococcus kloosii		100.0	0.32	ARG 1		
465512	**Good identification**					**465512**
Staphylococcus kloosii		100.0	0.33	ARG 1		
465550	**Very good identification**					**465550**
Staphylococcus kloosii		100.0	0.58	ARG 1		
465552	**Very good identification**					**465552**
Staphylococcus kloosii		100.0	0.59	ARG 1		
465570	**Good identification**					**465570**
Staphylococcus kloosii		100.0	0.46	ARG 1		
465572	**Acceptable identification**					**465572**
Staphylococcus kloosii		90.04	0.47	ARG 1		
Staphylococcus arlettae		9.96	0.15			
465650	**Good identification**					**465650**
Staphylococcus kloosii		100.0	0.32	ARG 1		
465652	**Good identification**					**465652**
Staphylococcus kloosii		100.0	0.32	ARG 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
465672	**Good identification**					**465672**
	Staphylococcus arlettae	97.43	0.41	ARG 1		
	Staphylococcus kloosii	2.57	0.20			
465673	**Good identification**					**465673**
	Staphylococcus arlettae	100.0	0.43	ARG 1		
465750	**Good identification**					**465750**
	Staphylococcus kloosii	100.0	0.38	ARG 1		
465752	**Low discrimination**					**465752**
	Staphylococcus kloosii	78.16	0.38	ARG 1		
	Staphylococcus arlettae	21.85	0.15	SUC 9		
465772	**Very good identification**					**465772**
	Staphylococcus arlettae	99.11	0.56	ARG 1		
	Staphylococcus kloosii	0.90	0.26			
465773	**Very good identification**					**465773**
	Staphylococcus arlettae	100.0	0.59	ARG 1		
466001	**Acceptable identification**					**466001**
	Staphylococcus carnosus subsp. carnosus	92.66	0.31	GLR 1		
	Staphylococcus schleiferi subsp. schleiferi	7.34	0.08			
466003	**Good identification**					**466003**
	Staphylococcus carnosus subsp. carnosus	99.34	0.43	GLR 1		
	Staphylococcus hyicus	0.66	-.06			
466011	**Good identification**					**466011**
	Staphylococcus carnosus subsp. carnosus	100.0	0.43	GLR 1		
466013	**Very good identification**					**466013**
	Staphylococcus carnosus subsp. carnosus	99.84	0.55	GLR 1		
	Staphylococcus simulans	0.16	0.01			
466103	**Low discrimination**					**466103**
	Staphylococcus carnosus subsp. carnosus	76.50	0.31	GLR 1		
	Staphylococcus hyicus	23.50	0.16			
466111	**Good identification**					**466111**
	Staphylococcus carnosus subsp. carnosus	100.0	0.31	GLR 1		
466113	**Good identification**					**466113**
	Staphylococcus carnosus subsp. carnosus	100.0	0.43	GLR 1		
466203	**Acceptable identification**					**466203**
	Staphylococcus hyicus	93.10	0.35	BGA 10		
	Staphylococcus simulans	5.48	0.17	GAL 92		
	Staphylococcus carnosus subsp. carnosus	1.42	0.02	TRE 92		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
466213	**Low discrimination**					**466213**
Staphylococcus simulans		76.78	0.29	URE 98		
Staphylococcus carnosus subsp. carnosus		19.93	0.14	TRE 95		
Staphylococcus hyicus		3.29	-.06			
466301	**Good identification**					**466301**
Staphylococcus hyicus		100.0	0.35	BGA 10		
466303	**Very good identification**					**466303**
					PYR	
Staphylococcus hyicus		99.64	0.57	BGA 10	1	
Staphylococcus intermedius		0.36	0.12	TRE 92	90	
466403	**Low discrimination**					**466403**
Staphylococcus carnosus subsp. carnosus		79.93	0.35	GLR 1		
Staphylococcus hyicus		14.32	0.16			
Staphylococcus simulans		5.75	0.15			
466411	**Good identification**					**466411**
Staphylococcus carnosus subsp. carnosus		100.0	0.35	GLR 1		
466413	**Acceptable identification**					**466413**
Staphylococcus carnosus subsp. carnosus		92.41	0.48	GLR 1		
Staphylococcus simulans		6.64	0.27			
Staphylococcus felis		0.95	0.01			
466503	**Low discrimination**					**466503**
Staphylococcus hyicus		88.39	0.38	BGA 10		
Staphylococcus carnosus subsp. carnosus		10.73	0.23	SUC 99		
Staphylococcus felis		0.88	-.06			
466513	**Low discrimination**					**466513**
Staphylococcus carnosus subsp. carnosus		66.12	0.35	GLR 1		
Staphylococcus condimenti		20.22	0.14			
Staphylococcus felis		9.10	0.12			
466601	**Good identification**					**466601**
Staphylococcus hyicus		98.88	0.35	BGA 10		
Staphylococcus simulans		1.12	0.02	GAL 92		
466602	**Low discrimination**					**466602**
Staphylococcus simulans		74.20	0.32	URE 98		
Staphylococcus hyicus		25.80	0.16			
466603	**Good identification**					**466603**
Staphylococcus hyicus		91.08	0.57	BGA 10		
Staphylococcus simulans		8.86	0.43	GAL 92		
Staphylococcus felis		0.06	-.10			
466612	**Good identification**					**466612**
Staphylococcus simulans		100.0	0.45	URE 98		
466613	**Very good identification**					**466613**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		95.48	0.55	URE 98			
Staphylococcus hyicus		2.48	0.16				
Staphylococcus felis		1.15	0.07				
466643	**Low discrimination**						**466643**
					COA	PYR	
Staphylococcus simulans		80.14	0.36	URE 98	1	70	
Staphylococcus hyicus		18.53	0.16		24	1	
Staphylococcus intermedius		1.34	-.03		99	90	
466652	**Good identification**						**466652**
Staphylococcus simulans		97.24	0.37	URE 98			
Staphylococcus haemolyticus		2.76	0.08				
466653	**Good identification**						**466653**
					COA		
Staphylococcus simulans		98.88	0.48	URE 98	1		
Staphylococcus intermedius		1.12	0.06		99		
466701	**Very good identification**						**466701**
					PYR		
Staphylococcus hyicus		99.22	0.57	BGA 10	1		
Staphylococcus intermedius		0.78	0.19	LAC 92	90		
466702	**Good identification**						**466702**
Staphylococcus hyicus		97.87	0.38	BGA 10			
Staphylococcus simulans		2.13	0.10	MNS 99			
466703	**Very good identification**						**466703**
					COA	PYR	
Staphylococcus hyicus		99.46	0.79	BGA 10	24	1	
Staphylococcus intermedius		0.42	0.35		99	90	
Staphylococcus simulans		0.07	0.21		1	70	
466711	**Low discrimination**						**466711**
					PYR		
Staphylococcus intermedius		67.72	0.28	URE 90	90		
Staphylococcus hyicus		32.28	0.16	GLR 3	1		
466713	**Low discrimination**						**466713**
					COA	PYR	
Staphylococcus intermedius		40.33	0.44	URE 90	99	90	
Staphylococcus hyicus		35.99	0.38	GLR 3	24	1	
Staphylococcus simulans		10.48	0.34		1	70	
466723	**Good identification**						**466723**
Staphylococcus hyicus		100.0	0.38	BGA 10			
466743	**Low discrimination**						**466743**
					COA	PYR	
Staphylococcus hyicus		59.33	0.38	BGA 10	24	1	
Staphylococcus intermedius		36.89	0.39	MLT 1	99	90	
Staphylococcus simulans		1.94	0.14		1	70	
466751	**Good identification**						**466751**
Staphylococcus intermedius		100.0	0.31	URE 90			
466753	**Acceptable identification**						**466753**
					COA	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus intermedius		91.83	0.48	URE 90	99	90	
Staphylococcus simulans		7.15	0.26	GLR 3	1	70	
Staphylococcus hyicus		0.55	-.04		24	1	
467013	**Good identification**						**467013**
Staphylococcus carnosus subsp. carnosus		100.0	0.36	GLR 1			
467413	**Good identification**						**467413**
Staphylococcus carnosus subsp. carnosus		100.0	0.28	GLR 1			
467703	**Good identification**						**467703**
Staphylococcus hyicus		100.0	0.38	BGA 10			
467771	**Acceptable identification**						**467771**
Staphylococcus gallinarum		91.93	0.27	URE 92			
Staphylococcus equorum		8.07	0.08	ARG 1			
467773	**Low discrimination**						**467773**
					YEP		
Staphylococcus gallinarum		86.60	0.35	URE 92	78		
Staphylococcus equorum		7.99	0.16	ARG 1	1		
Staphylococcus arlettae		3.97	0.18		99		
476703	**Good identification**						**476703**
Staphylococcus hyicus		100.0	0.38	ORN 1			
500000	**Low discrimination**						**500000**
					OXI		
Staphylococcus carnosus subsp. utilis		34.35	0.29	VPT 1	1		
Micrococcus sp.		27.73	0.28		89		
Dermacoccus nishinomiyaensis		16.76	0.33		93		
500001	**Acceptable identification**						**500001**
Staphylococcus capitis subsp. capitis		94.16	0.53	NIT 89			
Staphylococcus schleiferi subsp. schleiferi		4.04	0.26	MAN 93			
Staphylococcus auricularis		0.65	0.14				
500010	**Low discrimination**						**500010**
					NOV	YEP	
Staphylococcus capitis subsp. capitis		87.46	0.35	NIT 89	1	1	
Kocuria rosea		5.33	0.07	MNS 99	1	1	
Macrococcus bovicus		2.92	0.07		99	99	
500011	**Very good identification**						**500011**
					NOV		
Staphylococcus capitis subsp. capitis		99.98	0.76	NIT 89	1		
Staphylococcus cohnii subsp. cohnii		0.02	0.07		99		
500013	**Good identification**						**500013**
Staphylococcus capitis subsp. capitis		99.27	0.41	NIT 89			
Staphylococcus carnosus subsp. carnosus		0.73	-.02	LAC 2			
500031	**Good identification**						**500031**
Staphylococcus capitis subsp. capitis		100.0	0.35	NIT 89			
500040	**Low discrimination**						**500040**
					OXI		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		72.10	0.41	VPT 12	1		
Staphylococcus petrasii subsp. jettensis		12.41	0.10	NIT 90	1		
Micrococcus sp.		7.49	0.05	TRE 92	89		
500051	**Acceptable identification**						**500051**
					NOV		
Staphylococcus capitis subsp. capitis		93.89	0.35	NIT 89	1		
Staphylococcus cohnii subsp. cohnii		4.94	0.17	MLT 1	99		
Staphylococcus capitis subsp. ureolyticus		1.17	-.14		1		
500111	**Good identification**						**500111**
Staphylococcus capitis subsp. capitis		100.0	0.35	NIT 89			
500200	**Low discrimination**						**500200**
					OXI		
Staphylococcus auricularis		38.83	0.32	VPT 12	1		
Staphylococcus petrasii subsp. jettensis		16.89	0.10	NIT 90	1		
Micrococcus sp.		13.41	0.07	TRE 92	89		
500201	**Very good identification**						**500201**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		98.93	0.58	NIT 89	1	1	
Kocuria kristinae		0.73	0.13	MAN 93	99	93	
Staphylococcus auricularis		0.34	0.13		1	40	
500210	**Good identification**						**500210**
Staphylococcus capitis subsp. capitis		100.0	0.40	NIT 89			
500211	**Very good identification**						**500211**
Staphylococcus capitis subsp. capitis		99.99	0.81	NIT 89			
Staphylococcus capitis subsp. ureolyticus		0.01	-.14				
500213	**Good identification**						**500213**
Staphylococcus capitis subsp. capitis		97.63	0.47	NIT 89			
Staphylococcus capitis subsp. ureolyticus		2.37	0.04	LAC 2			
500231	**Good identification**						**500231**
Staphylococcus capitis subsp. capitis		100.0	0.40	NIT 89			
500240	**Low discrimination**						**500240**
Staphylococcus petrasii subsp. jettensis		69.13	0.51	TRE 99			
Staphylococcus petrasii subsp. pragensis		23.41	0.39				
Staphylococcus auricularis		3.75	0.40				
500241	**Low discrimination**						**500241**
					OXI	PYR	
Kocuria kristinae		61.65	0.33	ARG 1	99	93	
Staphylococcus petrasii subsp. jettensis		12.51	0.10	TRE 95	1	99	
Staphylococcus capitis subsp. capitis		9.40	0.17		1	1	
500242	**Low discrimination**						**500242**
Staphylococcus petrasii subsp. jettensis		86.19	0.36	TRE 99			
Staphylococcus haemolyticus		7.34	0.30				
Staphylococcus auricularis		2.94	0.20				
500250	**Low discrimination**						**500250**
					PYR		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		45.19	0.30	NIT 99	98		
Staphylococcus petrasii subsp. jettensis		29.08	0.10	TRE 93	99		
Staphylococcus warneri		9.91	0.24		8		
500251	**Low discrimination**						**500251**
					OXI	PYR	
Staphylococcus capitis subsp. capitis		84.67	0.40	NIT 89	1	1	
Staphylococcus capitis subsp. ureolyticus		13.59	0.15	MLT 1	1	1	
Kocuria kristinae		1.74	0.04		99	93	
500253	**Good identification**						**500253**
Staphylococcus capitis subsp. ureolyticus		98.30	0.33	URE 99			
Staphylococcus capitis subsp. capitis		1.71	0.05	NIT 95			
500311	**Good identification**						**500311**
Staphylococcus capitis subsp. capitis		100.0	0.40	NIT 89			
500340	**Low discrimination**						**500340**
Staphylococcus haemolyticus		53.38	0.36	NIT 99			
Staphylococcus petrasii subsp. jettensis		17.89	0.10	TRE 93			
Staphylococcus auricularis		10.66	0.20				
500342	**Low discrimination**						**500342**
					PYR		
Staphylococcus haemolyticus		51.97	0.31	NIT 99	98		
Staphylococcus epidermidis		38.31	0.26	TRE 93	1		
Staphylococcus petrasii subsp. jettensis		5.46	-.05		99		
500343	**Low discrimination**						**500343**
					OXI	PYR	
Staphylococcus epidermidis		84.84	0.32	URE 92	1	1	
Kocuria kristinae		15.17	0.10	NIT 97	99	93	
500350	**Acceptable identification**						**500350**
					PYR		
Staphylococcus haemolyticus		94.21	0.31	NIT 99	98		
Staphylococcus warneri		5.79	0.13	TRE 93	8		
500400	**Low discrimination**						**500400**
					NOV		
Staphylococcus carnosus subsp. utilis		58.85	0.44	VPT 1	1		
Staphylococcus auricularis		32.31	0.55		1		
Staphylococcus cohnii subsp. cohnii		7.24	0.45		99		
500401	**Low discrimination**						**500401**
					NOV		
Staphylococcus cohnii subsp. cohnii		41.52	0.46	ARG 1	99		
Staphylococcus schleiferi subsp. schleiferi		30.82	0.36		1		
Staphylococcus auricularis		19.00	0.35		1		
500402	**Low discrimination**						**500402**
Staphylococcus auricularis		80.55	0.35	VPT 12			
Staphylococcus carnosus subsp. utilis		13.34	0.03	NIT 90			
Staphylococcus petrasii subsp. jettensis		4.47	-.05	LAC 10			
500410	**Low discrimination**						**500410**
					NOV		
Staphylococcus cohnii subsp. cohnii		87.24	0.47	ARG 1	99		
Staphylococcus carnosus subsp. utilis		5.63	0.03		1		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		3.09	0.14		1		
500411	**Low discrimination**						**500411**
					NOV		
Staphylococcus capitis subsp. capitis		55.10	0.41	NIT 89	1		
Staphylococcus cohnii subsp. cohnii		44.90	0.48	TRE 2	99		
500440	**Low discrimination**						**500440**
Staphylococcus petrasii subsp. jettensis		44.79	0.51	SUC 99			
Staphylococcus auricularis		30.24	0.63				
Staphylococcus petrasii subsp. pragensis		15.17	0.39				
500441	**Low discrimination**						**500441**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		69.17	0.56	ARG 1	99	1	
Staphylococcus auricularis		23.33	0.43		1	40	
Staphylococcus petrasii subsp. jettensis		3.14	0.10		1	99	
500442	**Low discrimination**						**500442**
Staphylococcus petrasii subsp. jettensis		67.72	0.36	SUC 99			
Staphylococcus auricularis		28.79	0.43				
Staphylococcus petrasii subsp. pragensis		1.31	-.02				
500450	**Low discrimination**						**500450**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		89.02	0.58	ARG 1	99	1	
Staphylococcus petrasii subsp. jettensis		3.44	0.10		1	99	
Staphylococcus auricularis		2.32	0.22		1	40	
500451	**Very good identification**						**500451**
					NOV		
Staphylococcus cohnii subsp. cohnii		99.35	0.58	ARG 1	99		
Staphylococcus capitis subsp. capitis		0.39	0.00		1		
Staphylococcus auricularis		0.27	0.02		1		
500500	**Low discrimination**						**500500**
					NOV		
Staphylococcus auricularis		71.96	0.35	VPT 12	1		
Staphylococcus cohnii subsp. cohnii		16.13	0.25	NIT 90	99		
Staphylococcus carnosus subsp. utilis		11.92	0.03	GAL 10	1		
500501	**Low discrimination**						**500501**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		82.09	0.36	NIT 90	1		
Staphylococcus cohnii subsp. cohnii		12.29	0.26	PHS 99	99		
Staphylococcus auricularis		5.62	0.16		1		
500510	**Low discrimination**						**500510**
					OXI	YEP	
Staphylococcus cohnii subsp. cohnii		88.72	0.28	ARG 1	1	12	
Macrococcus bovicus		11.28	0.03	GAL 10	99	99	
500511	**Low discrimination**						**500511**
					NOV		
Staphylococcus cohnii subsp. cohnii		89.96	0.28	ARG 1	99		
Staphylococcus capitis subsp. capitis		10.04	0.00	GAL 10	1		
500540	**Low discrimination**						**500540**
					NOV	PYR	
Staphylococcus auricularis		59.34	0.43	VPT 12	1	40	
Staphylococcus cohnii subsp. cohnii		18.05	0.36	NIT 90	99	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests				Code
Staphylococcus petrasii subsp. jettensis		7.99	0.10	GAL 10	1	99			
500541	**Low discrimination**								**500541**
					OXI	NOV			
Staphylococcus cohnii subsp. cohnii		66.93	0.36	ARG 1	1	99			
Staphylococcus auricularis		22.57	0.23	GAL 10	1	1			
Kocuria kristinae		8.63	0.08		99	1			
500550	**Low discrimination**								**500550**
					OXI	PYR	NOV	YEP	
Staphylococcus cohnii subsp. cohnii		80.71	0.38	ARG 1	1	1	99	12	
Macrococcus bovicus		7.56	0.10	GAL 10	99	1	99	99	
Staphylococcus haemolyticus		6.60	0.13		1	98	2	20	
500551	**Good identification**								**500551**
					NOV				
Staphylococcus cohnii subsp. cohnii		96.53	0.39	ARG 1	99				
Staphylococcus caprae		3.48	-.07	GAL 10	1				
500600	**Low discrimination**								**500600**
Staphylococcus petrasii subsp. jettensis		61.69	0.51	MLT 99					
Staphylococcus petrasii subsp. pragensis		20.89	0.39						
Staphylococcus auricularis		16.47	0.54						
500601	**Low discrimination**								**500601**
					OXI	PYR			
Kocuria kristinae		63.55	0.39	ARG 1	99	93			
Staphylococcus auricularis		17.94	0.35	MLT 90	1	40			
Staphylococcus capitis subsp. capitis		9.27	0.23		1	1			
500602	**Low discrimination**								**500602**
Staphylococcus petrasii subsp. jettensis		82.79	0.36	MLT 99					
Staphylococcus auricularis		13.92	0.35						
Staphylococcus petrasii subsp. pragensis		1.61	-.02						
500603	**Low discrimination**								**500603**
					OXI				
Kocuria kristinae		83.80	0.27	ARG 1	99				
Staphylococcus auricularis		10.52	0.15	MLT 90	1				
Staphylococcus petrasii subsp. jettensis		5.69	-.05		1				
500611	**Good identification**								**500611**
					OXI	NOV	PYR		
Staphylococcus capitis subsp. capitis		97.47	0.47	NIT 89	1	1	1		
Kocuria kristinae		2.10	0.11	TRE 2	99	1	93		
Staphylococcus cohnii subsp. cohnii		0.43	0.07		1	99	1		
500640	**Low discrimination**								**500640**
Staphylococcus petrasii subsp. jettensis		74.09	0.92						
Staphylococcus petrasii subsp. pragensis		25.09	0.80						
Staphylococcus auricularis		0.47	0.62						
500641	**Low discrimination**								**500641**
					OXI				
Staphylococcus petrasii subsp. jettensis		40.45	0.51	MNS 1	1				
Kocuria kristinae		38.27	0.59		99				
Staphylococcus petrasii subsp. pragensis		13.70	0.39		1				
500642	**Very good identification**								**500642**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. jettensis		96.43	0.77				
Staphylococcus petrasii subsp. pragensis		1.87	0.39				
Staphylococcus haemolyticus		1.10	0.53				
500643	**Low discrimination**						**500643**
					OXI		
Kocuria kristinae		48.68	0.47	ARG 1	99		
Staphylococcus petrasii subsp. jettensis		36.33	0.36		1		
Staphylococcus petrasii subsp. petrasii		11.44	0.34		1		
500650	**Low discrimination**						**500650**
Staphylococcus petrasii subsp. jettensis		61.23	0.51	MAN 1			
Staphylococcus petrasii subsp. pragensis		20.74	0.39				
Staphylococcus haemolyticus		12.77	0.53				
500651	**Low discrimination**						**500651**
					OXI		
Kocuria kristinae		50.66	0.31	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		14.73	0.20	MAN 4	1		
Staphylococcus petrasii subsp. jettensis		12.98	0.10		1		
500652	**Low discrimination**						**500652**
					NOV	PYR	
Staphylococcus petrasii subsp. jettensis		50.41	0.36	MAN 1	1	99	
Staphylococcus haemolyticus		33.51	0.48		2	98	
Staphylococcus saprophyticus subsp. saprophyticus		8.03	0.22		99	7	
500660	**Low discrimination**						**500660**
Staphylococcus petrasii subsp. jettensis		74.10	0.51	XYL 1			
Staphylococcus petrasii subsp. pragensis		25.10	0.39				
Staphylococcus auricularis		0.47	0.21				
500662	**Good identification**						**500662**
Staphylococcus petrasii subsp. jettensis		96.64	0.36	XYL 1			
Staphylococcus petrasii subsp. pragensis		1.87	-.02				
Staphylococcus haemolyticus		1.11	0.12				
500700	**Low discrimination**						**500700**
Staphylococcus auricularis		60.88	0.35	VPT 12			
Staphylococcus petrasii subsp. jettensis		20.72	0.10	NIT 90			
Staphylococcus haemolyticus		8.30	0.18	GAL 10			
500701	**Low discrimination**						**500701**
					OXI		
Kocuria kristinae		88.19	0.30	ARG 1	99		
Staphylococcus auricularis		8.30	0.15	MLT 90	1		
Rothia mucilaginosa		1.76	-.09		1		
500740	**Low discrimination**						**500740**
Staphylococcus petrasii subsp. jettensis		53.94	0.51	GAL 1			
Staphylococcus haemolyticus		21.60	0.59				
Staphylococcus petrasii subsp. pragensis		18.27	0.39				
500741	**Low discrimination**						**500741**
					OXI		
Kocuria kristinae		75.20	0.49	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		16.40	0.36		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		2.95	0.28		1		
500742	**Low discrimination**						**500742**
					PYR		
Staphylococcus haemolyticus		51.57	0.54	NIT 99	98		
Staphylococcus petrasii subsp. jettensis		40.41	0.36		99		
Staphylococcus warneri		2.18	0.33		8		
500743	**Low discrimination**						**500743**
					OXI		
Kocuria kristinae		62.83	0.37	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		29.52	0.30		1		
Staphylococcus haemolyticus		5.55	0.23		1		
500750	**Low discrimination**						**500750**
					PYR		
Staphylococcus haemolyticus		86.59	0.54	NIT 99	98		
Staphylococcus warneri		7.61	0.40		8		
Staphylococcus petrasii subsp. jettensis		3.72	0.10		99		
500752	**Acceptable identification**						**500752**
					NOV	PYR	
Staphylococcus haemolyticus		91.56	0.49	NIT 99	2	98	
Staphylococcus warneri		4.77	0.30		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		1.69	0.00		99	7	
501011	**Good identification**						**501011**
Staphylococcus capitis subsp. capitis		100.0	0.35	ESL 1			
501211	**Good identification**						**501211**
Staphylococcus capitis subsp. capitis		100.0	0.40	ESL 1			
501601	**Good identification**						**501601**
Kocuria kristinae		100.0	0.27	ARG 1			
501640	**Low discrimination**						**501640**
Staphylococcus petrasii subsp. jettensis		73.97	0.51	ESL 1			
Staphylococcus petrasii subsp. pragensis		25.05	0.39				
Staphylococcus auricularis		0.47	0.21				
501641	**Acceptable identification**						**501641**
					OXI		
Kocuria kristinae		93.94	0.47	ARG 1	99		
Staphylococcus petrasii subsp. jettensis		4.01	0.10		1		
Staphylococcus petrasii subsp. pragensis		1.36	-.02		1		
501642	**Good identification**						**501642**
Staphylococcus petrasii subsp. jettensis		96.64	0.36	ESL 1			
Staphylococcus petrasii subsp. pragensis		1.87	-.02				
Staphylococcus haemolyticus		1.11	0.12				
501643	**Good identification**						**501643**
					OXI		
Kocuria kristinae		97.07	0.34	ARG 1	99		
Staphylococcus petrasii subsp. jettensis		2.93	-.05		1		
501741	**Low discrimination**						**501741**
					OXI		
Kocuria kristinae		83.70	0.37	ARG 1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Rothia mucilaginosa		15.57	0.18		1	
Staphylococcus petrasii subsp. petrasii		0.74	-.05		1	
502000	**Low discrimination**					**502000**
Staphylococcus carnosus subsp. utilis		47.98	0.44	VPT	1	
Kocuria rosea		24.39	0.47			
Staphylococcus auricularis		20.61	0.53			
502001	**Good identification**					**502001**
Staphylococcus capitis subsp. capitis		94.06	0.71	MAN	93	
Staphylococcus schleiferi subsp. schleiferi		4.49	0.46			
Staphylococcus auricularis		0.72	0.33			
502002	**Low discrimination**					**502002**
Staphylococcus auricularis		68.23	0.33	VPT	12	
Staphylococcus carnosus subsp. utilis		14.44	0.03	TRE	92	
Kocuria varians		8.16	0.15	LAC	10	
502003	**Low discrimination**					**502003**
Staphylococcus capitis subsp. capitis		73.31	0.37	MAN	93	
Staphylococcus carnosus subsp. carnosus		21.89	0.27	LAC	2	
Staphylococcus auricularis		3.07	0.14			
502010	**Good identification**					**502010**
Staphylococcus capitis subsp. capitis		92.16	0.53	MNS	99	
Kocuria rosea		6.24	0.27			
Staphylococcus carnosus subsp. utilis		1.12	0.03			
502011	**Excellent identification**					**502011**
Staphylococcus capitis subsp. capitis		99.95	0.95			
Staphylococcus carnosus subsp. carnosus		0.05	0.27			
Staphylococcus schleiferi subsp. schleiferi		0.00	0.04			
502013	**Good identification**					**502013**
Staphylococcus capitis subsp. capitis		91.38	0.60	LAC	2	
Staphylococcus carnosus subsp. carnosus		8.22	0.39			
Staphylococcus capitis subsp. ureolyticus		0.40	0.02			
502020	**Acceptable identification**					**502020**
Kocuria rosea		91.00	0.39	ARG	1	
Staphylococcus carnosus subsp. utilis		4.22	0.03			
Kocuria varians		2.97	0.17			
502021	**Low discrimination**					**502021**
Staphylococcus capitis subsp. capitis		81.27	0.30	MAN	93	
Kocuria rosea		14.85	0.13	XYL	1	
Staphylococcus schleiferi subsp. schleiferi		3.88	0.04			
502031	**Very good identification**					**502031**
Staphylococcus capitis subsp. capitis		99.85	0.53	XYL	1	
Kocuria rosea		0.15	-.07			
502040	**Low discrimination**					**502040**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		89.70	0.60	VPT 12			
Staphylococcus haemolyticus		4.54	0.35	TRE 92			
Staphylococcus petrasii subsp. jettensis		4.00	0.18				
502041	**Low discrimination**						**502041**
Staphylococcus auricularis		62.33	0.41	VPT 12			
Staphylococcus capitis subsp. capitis		35.12	0.30	TRE 92			
Staphylococcus schleiferi subsp. schleiferi		1.68	0.04	MNS 10			
502042	**Low discrimination**						**502042**
Staphylococcus auricularis		74.05	0.41	VPT 12			
Staphylococcus haemolyticus		18.97	0.30	TRE 92			
Staphylococcus petrasii subsp. jettensis		5.25	0.02	LAC 10			
502050	**Low discrimination**						**502050**
					PYR		
Staphylococcus haemolyticus		61.63	0.30	SUC 99	98		
Staphylococcus auricularis		20.95	0.19	TRE 93	40		
Staphylococcus capitis subsp. capitis		17.42	0.12		1		
502051	**Very good identification**						**502051**
Staphylococcus capitis subsp. capitis		97.17	0.53	MLT 1			
Staphylococcus capitis subsp. ureolyticus		2.83	0.13				
502053	**Acceptable identification**						**502053**
Staphylococcus capitis subsp. ureolyticus		90.58	0.31	URE 99			
Staphylococcus capitis subsp. capitis		8.65	0.19	SUC 96			
Staphylococcus carnosus subsp. carnosus		0.78	-.02				
502100	**Low discrimination**						**502100**
Staphylococcus auricularis		44.91	0.33	VPT 12			
Kocuria varians		28.56	0.30	GAL 10			
Staphylococcus carnosus subsp. utilis		9.50	0.03	TRE 92			
502101	**Low discrimination**						**502101**
Staphylococcus schleiferi subsp. schleiferi		80.25	0.46	PHS 99			
Staphylococcus capitis subsp. capitis		16.97	0.30				
Staphylococcus auricularis		1.43	0.14				
502111	**Very good identification**						**502111**
Staphylococcus capitis subsp. capitis		98.53	0.53	GAL 1			
Staphylococcus carnosus subsp. carnosus		1.12	0.15				
Staphylococcus schleiferi subsp. schleiferi		0.35	0.04				
502113	**Low discrimination**						**502113**
					COA		
Staphylococcus carnosus subsp. carnosus		63.94	0.27	BGA 90	1		
Staphylococcus capitis subsp. capitis		28.74	0.19	PHS 99	1		
Staphylococcus schleiferi subsp. coagulans		7.33	0.12		99		
502140	**Low discrimination**						**502140**
					OXI	NOV	
Staphylococcus auricularis		56.91	0.41	VPT 12	1	1	
Staphylococcus haemolyticus		29.22	0.36	GAL 10	1	2	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus caseolyticus		9.22	0.14	TRE 92	99	99	
502142	**Low discrimination**						**502142**
					OXI	NOV	
Staphylococcus haemolyticus		39.60	0.31	SUC 99	1	2	
Macrococcus caseolyticus		33.31	0.18	TRE 93	99	99	
Staphylococcus auricularis		15.24	0.21		1	1	
502150	**Good identification**						**502150**
Staphylococcus haemolyticus		100.0	0.31	SUC 99			
502200	**Low discrimination**						**502200**
Staphylococcus auricularis		58.22	0.52	VPT 12			
Staphylococcus capitis subsp. capitis		17.11	0.36	TRE 92			
Kocuria rosea		8.29	0.27				
502201	**Very good identification**						**502201**
Staphylococcus capitis subsp. capitis		99.56	0.77	MAN 93			
Staphylococcus auricularis		0.38	0.32				
Staphylococcus schleiferi subsp. schleiferi		0.03	0.04				
502202	**Low discrimination**						**502202**
Staphylococcus auricularis		52.32	0.32	VPT 12			
Staphylococcus haemolyticus		33.87	0.30	TRE 92			
Staphylococcus petrasii subsp. jettensis		9.37	0.02	LAC 10			
502203	**Acceptable identification**						**502203**
Staphylococcus capitis subsp. capitis		94.06	0.42	MAN 93			
Staphylococcus capitis subsp. ureolyticus		2.97	0.02	LAC 2			
Staphylococcus auricularis		1.96	0.13				
502210	**Very good identification**						**502210**
					PYR		
Staphylococcus capitis subsp. capitis		97.28	0.59	MNS 99	1		
Staphylococcus haemolyticus		1.87	0.30		98		
Kocuria rosea		0.39	0.07		1		
502211	**Excellent identification**						**502211**
Staphylococcus capitis subsp. capitis		99.98	1.00				
Staphylococcus capitis subsp. ureolyticus		0.02	0.13				
502213	**Good identification**						**502213**
Staphylococcus capitis subsp. capitis		94.57	0.65	LAC 2			
Staphylococcus capitis subsp. ureolyticus		5.39	0.31				
Staphylococcus carnosus subsp. carnosus		0.05	-.02				
502221	**Good identification**						**502221**
Staphylococcus capitis subsp. capitis		98.92	0.36	MAN 93			
Kocuria rosea		1.08	-.07	XYL 1			
502231	**Very good identification**						**502231**
Staphylococcus capitis subsp. capitis		100.0	0.59	XYL 1			
502240	**Low discrimination**						**502240**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus haemolyticus		47.71	0.76	TRE 93		
Staphylococcus petrasii subsp. jettensis		42.07	0.59			
Staphylococcus auricularis		8.79	0.60			
502241	**Low discrimination**					**502241**
					PYR	
Staphylococcus haemolyticus		28.24	0.45	TRE 93	98	
Staphylococcus capitis subsp. capitis		21.20	0.36	MNS 3	1	
Staphylococcus auricularis		18.70	0.40		40	
502242	**Low discrimination**					**502242**
					PYR	
Staphylococcus haemolyticus		72.86	0.71	TRE 93	98	
Staphylococcus petrasii subsp. jettensis		20.16	0.43		99	
Staphylococcus epidermidis		3.49	0.42		1	
502243	**Low discrimination**					**502243**
					PYR	
Staphylococcus epidermidis		45.70	0.47	URE 92	1	
Staphylococcus capitis subsp. ureolyticus		32.41	0.31		1	
Staphylococcus haemolyticus		15.88	0.40		98	
502250	**Very good identification**					**502250**
					PYR	
Staphylococcus haemolyticus		97.31	0.71	TRE 93	98	
Staphylococcus petrasii subsp. jettensis		1.48	0.18		99	
Staphylococcus capitis subsp. capitis		0.52	0.18		1	
502251	**Low discrimination**					**502251**
					PYR	
Staphylococcus capitis subsp. capitis		69.59	0.59	MLT 1	1	
Staphylococcus capitis subsp. ureolyticus		26.22	0.41		1	
Staphylococcus haemolyticus		4.10	0.40		98	
502252	**Very good identification**					**502252**
					PYR	
Staphylococcus haemolyticus		96.72	0.66	TRE 93	98	
Staphylococcus capitis subsp. ureolyticus		2.52	0.18		1	
Staphylococcus petrasii subsp. jettensis		0.46	0.02		99	
502253	**Very good identification**					**502253**
					PYR	
Staphylococcus capitis subsp. ureolyticus		98.04	0.59	URE 99	1	
Staphylococcus haemolyticus		1.18	0.35		98	
Staphylococcus capitis subsp. capitis		0.72	0.24		1	
502260	**Low discrimination**					**502260**
Staphylococcus haemolyticus		48.40	0.35	TRE 93		
Staphylococcus petrasii subsp. jettensis		42.68	0.18	XYL 1		
Staphylococcus auricularis		8.92	0.19			
502262	**Low discrimination**					**502262**
					PYR	
Staphylococcus haemolyticus		75.50	0.30	TRE 93	98	
Staphylococcus petrasii subsp. jettensis		20.89	0.02	XYL 1	99	
Staphylococcus epidermidis		3.62	0.01		1	
502270	**Good identification**					**502270**
Staphylococcus haemolyticus		100.0	0.30	TRE 93		
502300	**Low discrimination**					**502300**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus haemolyticus		53.03	0.36	TRE 93	98	
Staphylococcus auricularis		40.86	0.32	MLT 99	40	
Kocuria varians		3.84	0.13		10	
502301	**Low discrimination**					**502301**
Staphylococcus capitis subsp. capitis		88.24	0.36	GAL 1		
Staphylococcus auricularis		3.71	0.13	MAN 93		
Staphylococcus epidermidis		3.44	0.11			
502302	**Low discrimination**					**502302**
					PYR	
Staphylococcus haemolyticus		71.79	0.31	TRE 93	98	
Staphylococcus epidermidis		17.28	0.16	MLT 99	1	
Staphylococcus auricularis		10.93	0.13		40	
502310	**Low discrimination**					**502310**
					PYR	
Staphylococcus haemolyticus		68.23	0.31	TRE 93	98	
Staphylococcus capitis subsp. capitis		31.77	0.18	MLT 99	1	
502311	**Very good identification**					**502311**
Staphylococcus capitis subsp. capitis		100.0	0.59	GAL 1		
502340	**Good identification**					**502340**
					PYR	
Staphylococcus haemolyticus		92.63	0.77	TRE 93	98	
Staphylococcus epidermidis		3.96	0.47		1	
Staphylococcus auricularis		1.68	0.40		40	
502341	**Low discrimination**					**502341**
					PYR	
Staphylococcus epidermidis		68.21	0.52	URE 92	1	
Staphylococcus haemolyticus		26.56	0.46		98	
Staphylococcus petrasii subsp. petrasii		1.80	0.14		99	
502342	**Low discrimination**					**502342**
					PYR	
Staphylococcus haemolyticus		79.05	0.72	TRE 93	98	
Staphylococcus epidermidis		19.03	0.57		1	
Staphylococcus hominis subsp. hominis		0.68	0.31		1	
502343	**Good identification**					**502343**
					PYR	
Staphylococcus epidermidis		92.77	0.63	URE 92	1	
Staphylococcus haemolyticus		6.42	0.41		98	
Staphylococcus petrasii subsp. petrasii		0.42	0.09		99	
502350	**Very good identification**					**502350**
					PYR	
Staphylococcus haemolyticus		99.82	0.72	TRE 93	98	
Staphylococcus hominis subsp. hominis		0.11	0.12		1	
Staphylococcus epidermidis		0.07	0.06		1	
502351	**Low discrimination**					**502351**
					PYR	
Staphylococcus haemolyticus		78.97	0.41	TRE 93	98	
Staphylococcus capitis subsp. capitis		12.01	0.18	MNS 3	1	
Staphylococcus capitis subsp. ureolyticus		4.53	0.00		1	
502352	**Very good identification**					**502352**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus haemolyticus		99.26	0.67	TRE 93	98	
Staphylococcus epidermidis		0.41	0.16		1	
Staphylococcus hominis subsp. hominis		0.22	0.14		1	
502353	**Low discrimination**					**502353**
					PYR	
Staphylococcus haemolyticus		44.31	0.36	TRE 93	98	
Staphylococcus capitis subsp. ureolyticus		33.10	0.18	MNS 3	1	
Staphylococcus caprae		11.57	0.11		75	
502360	**Good identification**					**502360**
					PYR	
Staphylococcus haemolyticus		95.90	0.36	TRE 93	98	
Staphylococcus epidermidis		4.10	0.06	XYL 1	1	
502362	**Low discrimination**					**502362**
					PYR	
Staphylococcus haemolyticus		80.60	0.31	TRE 93	98	
Staphylococcus epidermidis		19.41	0.16	XYL 1	1	
502370	**Good identification**					**502370**
Staphylococcus haemolyticus		100.0	0.31	TRE 93		
502400	**Low discrimination**					**502400**
Staphylococcus carnosus subsp. utilis		51.13	0.59	VPT 1		
Staphylococcus auricularis		48.11	0.75			
Staphylococcus petrasii subsp. jettensis		0.44	0.18			
502401	**Low discrimination**					**502401**
Staphylococcus schleiferi subsp. schleiferi		54.51	0.55	PHS 99		
Staphylococcus auricularis		33.61	0.55			
Staphylococcus capitis subsp. capitis		7.76	0.37			
502402	**Low discrimination**					**502402**
Staphylococcus auricularis		89.22	0.55	VPT 12		
Staphylococcus carnosus subsp. utilis		8.62	0.18	LAC 10		
Staphylococcus petrasii subsp. jettensis		1.28	0.02			
502403	**Low discrimination**					**502403**
Staphylococcus auricularis		68.69	0.35	VPT 12		
Staphylococcus carnosus subsp. carnosus		18.27	0.19	MNS 10		
Staphylococcus schleiferi subsp. schleiferi		10.13	0.14	LAC 10		
502411	**Very good identification**					**502411**
					NOV	
Staphylococcus capitis subsp. capitis		97.48	0.60	TRE 2	1	
Staphylococcus carnosus subsp. carnosus		0.94	0.19		1	
Staphylococcus cohnii subsp. cohnii		0.74	0.25		99	
502413	**Low discrimination**					**502413**
Staphylococcus carnosus subsp. carnosus		65.37	0.32	BGA 90		
Staphylococcus capitis subsp. capitis		34.63	0.25	PHS 99		
502440	**Low discrimination**					**502440**
Staphylococcus auricularis		68.59	0.82	VPT 12		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. jettensis		26.34	0.59				
Staphylococcus haemolyticus		4.01	0.58				
502441	**Good identification**						**502441**
					NOV	PYR	
Staphylococcus auricularis		91.00	0.63	VPT 12	1	40	
Staphylococcus petrasii subsp. jettensis		3.18	0.18	MNS 10	1	99	
Staphylococcus cohnii subsp. cohnii		2.26	0.33		99	1	
502442	**Low discrimination**						**502442**
Staphylococcus auricularis		51.84	0.63	VPT 12			
Staphylococcus petrasii subsp. jettensis		31.62	0.43	LAC 10			
Staphylococcus haemolyticus		15.34	0.53				
502443	**Low discrimination**						**502443**
Staphylococcus auricularis		82.98	0.43	VPT 12			
Staphylococcus haemolyticus		6.83	0.22	MNS 10			
Staphylococcus petrasii subsp. petrasii		5.59	0.12	LAC 10			
502450	**Low discrimination**						**502450**
Staphylococcus haemolyticus		66.14	0.53	SUC 99			
Staphylococcus auricularis		19.46	0.41				
Staphylococcus petrasii subsp. jettensis		7.47	0.18				
502451	**Low discrimination**						**502451**
					NOV		
Staphylococcus cohnii subsp. cohnii		47.79	0.35	ARG 1	99		
Staphylococcus capitis subsp. capitis		20.11	0.19	NIT 7	1		
Staphylococcus auricularis		15.29	0.22		1		
502452	**Acceptable identification**						**502452**
Staphylococcus haemolyticus		91.44	0.48	SUC 99			
Staphylococcus auricularis		5.32	0.22				
Staphylococcus petrasii subsp. jettensis		3.24	0.02				
502460	**Low discrimination**						**502460**
Staphylococcus auricularis		69.33	0.41	VPT 12			
Staphylococcus petrasii subsp. jettensis		26.62	0.18	XYL 1			
Staphylococcus haemolyticus		4.05	0.17				
502500	**Low discrimination**						**502500**
					OXI	NOV	
Staphylococcus auricularis		85.70	0.55	VPT 12	1	1	
Staphylococcus carnosus subsp. utilis		8.28	0.18	GAL 10	1	1	
Macrococcus caseolyticus		2.82	0.14		99	99	
502501	**Good identification**						**502501**
Staphylococcus schleiferi subsp. schleiferi		93.33	0.55	PHS 99			
Staphylococcus auricularis		6.39	0.35				
Rothia mucilaginosa		0.14	-.09				
502502	**Low discrimination**						**502502**
					OXI	NOV	
Staphylococcus auricularis		64.37	0.35	VPT 12	1	1	
Macrococcus caseolyticus		28.55	0.18	GAL 10	99	99	
Staphylococcus haemolyticus		4.56	0.13	LAC 10	1	2	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
502540	**Low discrimination**						**502540**
					OXI	NOV	
Macrococcus caseolyticus		46.04	0.55	ARG 1	99	99	
Staphylococcus auricularis		33.03	0.63		1	1	
Staphylococcus haemolyticus		19.59	0.59		1	2	
502541	**Low discrimination**						**502541**
					OXI	NOV	
Staphylococcus auricularis		67.34	0.43	VPT 12	1	1	
Staphylococcus haemolyticus		11.12	0.28	GAL 10	1	2	
Macrococcus caseolyticus		8.54	0.14	MNS 10	99	99	
502542	**Low discrimination**						**502542**
					OXI	NOV	
Macrococcus caseolyticus		82.14	0.59	ARG 1	99	99	
Staphylococcus haemolyticus		13.11	0.54		1	2	
Staphylococcus auricularis		4.36	0.43		1	1	
502550	**Good identification**						**502550**
					OXI	NOV	
Staphylococcus haemolyticus		92.20	0.54	SUC 99	1	2	
Macrococcus caseolyticus		3.73	0.14		99	99	
Staphylococcus auricularis		2.67	0.22		1	1	
502552	**Low discrimination**						**502552**
					OXI	NOV	
Staphylococcus haemolyticus		76.15	0.49	SUC 99	1	2	
Staphylococcus caprae		15.20	0.22		1	1	
Macrococcus caseolyticus		8.21	0.18		99	99	
502553	**Good identification**						**502553**
Staphylococcus caprae		97.93	0.39	URE 90			
Staphylococcus haemolyticus		2.07	0.18	PHS 99			
502600	**Low discrimination**						**502600**
Staphylococcus auricularis		46.56	0.74	VPT 12			
Staphylococcus petrasii subsp. jettensis		45.20	0.59				
Staphylococcus haemolyticus		6.88	0.58				
502601	**Low discrimination**						**502601**
					PYR		
Staphylococcus auricularis		59.56	0.54	VPT 12	40		
Staphylococcus capitis subsp. capitis		27.67	0.42	MNS 10	1		
Staphylococcus petrasii subsp. jettensis		5.26	0.18		99		
502602	**Low discrimination**						**502602**
Staphylococcus petrasii subsp. jettensis		46.76	0.43	MLT 99			
Staphylococcus auricularis		30.33	0.54				
Staphylococcus haemolyticus		22.69	0.53				
502603	**Low discrimination**						**502603**
Staphylococcus auricularis		57.10	0.35	VPT 12			
Staphylococcus haemolyticus		11.89	0.22	MNS 10			
Staphylococcus petrasii subsp. petrasii		9.72	0.12	LAC 10			
502610	**Low discrimination**						**502610**
Staphylococcus haemolyticus		75.28	0.53	MLT 99			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		8.76	0.33			
Staphylococcus petrasii subsp. jettensis		8.51	0.18			
502611	**Very good identification**					**502611**
					PYR	
Staphylococcus capitis subsp. capitis		99.39	0.65	TRE 2	1	
Staphylococcus haemolyticus		0.39	0.22		98	
Staphylococcus auricularis		0.16	0.13		40	
502612	**Acceptable identification**					**502612**
Staphylococcus haemolyticus		93.74	0.48	MLT 99		
Staphylococcus petrasii subsp. jettensis		3.32	0.02			
Staphylococcus auricularis		2.16	0.13			
502613	**Low discrimination**					**502613**
					PYR	
Staphylococcus capitis subsp. capitis		81.07	0.30	TRE 2	1	
Staphylococcus haemolyticus		8.74	0.17	LAC 2	98	
Staphylococcus simulans		7.90	0.14		70	
502620	**Low discrimination**					**502620**
Staphylococcus auricularis		47.20	0.33	VPT 12		
Staphylococcus petrasii subsp. jettensis		45.82	0.18	XYL 1		
Staphylococcus haemolyticus		6.97	0.17			
502640	**Low discrimination**					**502640**
Staphylococcus petrasii subsp. jettensis		83.78	1.00			
Staphylococcus haemolyticus		12.75	0.99			
Staphylococcus auricularis		2.03	0.82			
502641	**Low discrimination**					**502641**
Staphylococcus petrasii subsp. jettensis		45.85	0.59	MNS 1		
Staphylococcus haemolyticus		21.37	0.68			
Staphylococcus petrasii subsp. petrasii		18.25	0.59			
502642	**Low discrimination**					**502642**
Staphylococcus petrasii subsp. jettensis		66.48	0.85			
Staphylococcus haemolyticus		32.25	0.94			
Staphylococcus auricularis		1.02	0.62			
502643	**Low discrimination**					**502643**
Staphylococcus haemolyticus		37.24	0.63	MNS 3		
Staphylococcus petrasii subsp. petrasii		30.44	0.53			
Staphylococcus petrasii subsp. jettensis		25.07	0.43			
502650	**Low discrimination**					**502650**
Staphylococcus haemolyticus		89.27	0.94			
Staphylococcus petrasii subsp. jettensis		10.09	0.59			
Staphylococcus auricularis		0.25	0.40			
502651	**Low discrimination**					**502651**
Staphylococcus haemolyticus		80.39	0.63	MNS 3		
Staphylococcus petrasii subsp. petrasii		12.99	0.39			
Staphylococcus petrasii subsp. jettensis		2.97	0.18			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
502652	**Very good identification**						**502652**
					PYR		
Staphylococcus haemolyticus		96.32	0.89		98		
Staphylococcus petrasii subsp. jettensis		3.42	0.43		99		
Staphylococcus warneri		0.08	0.33		8		
502653	**Low discrimination**						**502653**
					PYR		
Staphylococcus haemolyticus		80.55	0.58	MNS 3	98		
Staphylococcus petrasii subsp. petrasii		12.46	0.34		99		
Staphylococcus capitis subsp. ureolyticus		5.11	0.18		1		
502660	**Low discrimination**						**502660**
Staphylococcus petrasii subsp. jettensis		83.82	0.59	XYL 1			
Staphylococcus haemolyticus		12.76	0.58				
Staphylococcus auricularis		2.04	0.40				
502662	**Low discrimination**						**502662**
Staphylococcus petrasii subsp. jettensis		66.65	0.43	XYL 1			
Staphylococcus haemolyticus		32.33	0.53				
Staphylococcus auricularis		1.02	0.21				
502670	**Low discrimination**						**502670**
Staphylococcus haemolyticus		89.85	0.53	XYL 1			
Staphylococcus petrasii subsp. jettensis		10.15	0.18				
502672	**Good identification**						**502672**
Staphylococcus haemolyticus		96.58	0.48	XYL 1			
Staphylococcus petrasii subsp. jettensis		3.42	0.02				
502700	**Low discrimination**						**502700**
Staphylococcus haemolyticus		55.38	0.59	MLT 99			
Staphylococcus auricularis		36.94	0.54				
Staphylococcus petrasii subsp. jettensis		3.26	0.18				
502701	**Low discrimination**						**502701**
Rothia mucilaginosa		54.15	0.32	ARG 1			
Staphylococcus auricularis		23.25	0.35	ESL 90			
Staphylococcus haemolyticus		9.70	0.28				
502702	**Low discrimination**						**502702**
					OXI	NOV	
Staphylococcus haemolyticus		83.28	0.54	MLT 99	1	2	
Staphylococcus auricularis		10.97	0.35		1	1	
Macrococcus caseolyticus		3.52	0.14		99	99	
502710	**Very good identification**						**502710**
Staphylococcus haemolyticus		98.87	0.54	MLT 99			
Staphylococcus auricularis		1.13	0.13				
502712	**Good identification**						**502712**
Staphylococcus haemolyticus		99.39	0.49	MLT 99			
Staphylococcus piscifermentans		0.61	0.12				
502740	**Good identification**						**502740**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Staphylococcus haemolyticus		91.26	1.00		1	2	
Staphylococcus petrasii subsp. jettensis		5.37	0.59		1	1	
Macrococcus caseolyticus		1.44	0.52		99	99	
502741	**Low discrimination**						**502741**
Staphylococcus haemolyticus		57.41	0.69	MNS 3			
Staphylococcus petrasii subsp. petrasii		28.99	0.55				
Rothia mucilaginosa		7.55	0.39				
502742	**Good identification**						**502742**
					OXI	NOV	
Staphylococcus haemolyticus		93.64	0.95		1	2	
Macrococcus caseolyticus		3.95	0.55		99	99	
Staphylococcus petrasii subsp. jettensis		1.73	0.43		1	1	
502743	**Low discrimination**						**502743**
Staphylococcus haemolyticus		63.10	0.64	MNS 3			
Staphylococcus petrasii subsp. petrasii		30.50	0.50				
Staphylococcus lugdunensis		2.64	0.29				
502750	**Excellent identification**						**502750**
					PYR		
Staphylococcus haemolyticus		99.75	0.95		98		
Staphylococcus petrasii subsp. jettensis		0.10	0.18		99		
Staphylococcus warneri		0.04	0.32		8		
502751	**Good identification**						**502751**
Staphylococcus haemolyticus		90.47	0.64	MNS 3			
Staphylococcus petrasii subsp. petrasii		8.64	0.36				
Staphylococcus caprae		0.47	0.04				
502752	**Very good identification**						**502752**
					OXI	NOV	
Staphylococcus haemolyticus		99.70	0.90		1	2	
Staphylococcus caprae		0.07	0.13		1	1	
Macrococcus caseolyticus		0.07	0.14		99	99	
502753	**Low discrimination**						**502753**
Staphylococcus haemolyticus		74.40	0.59	MNS 3			
Staphylococcus caprae		13.16	0.30				
Staphylococcus petrasii subsp. petrasii		6.80	0.30				
502760	**Good identification**						**502760**
					OXI	NOV	
Staphylococcus haemolyticus		91.56	0.59	XYL 1	1	2	
Staphylococcus petrasii subsp. jettensis		5.39	0.18		1	1	
Macrococcus caseolyticus		1.45	0.11		99	99	
502761	**Low discrimination**						**502761**
Staphylococcus haemolyticus		59.07	0.28	XYL 1			
Staphylococcus petrasii subsp. petrasii		29.82	0.14	MNS 3			
Rothia mucilaginosa		7.77	-.02				
502762	**Good identification**						**502762**
					OXI	NOV	
Staphylococcus haemolyticus		94.01	0.54	XYL 1	1	2	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus caseolyticus		3.97	0.14		99	99	
Staphylococcus petrasii subsp. jettensis		1.74	0.02		1	1	
502770	**Very good identification**						**502770**
Staphylococcus haemolyticus		100.0	0.54	XYL 1			
502772	**Good identification**						**502772**
Staphylococcus haemolyticus		100.0	0.49	XYL 1			
503001	**Good identification**						**503001**
Staphylococcus capitis subsp. capitis		95.44	0.30	ESL 1			
Staphylococcus schleiferi subsp. schleiferi		4.56	0.04	MAN 93			
503011	**Very good identification**						**503011**
Staphylococcus capitis subsp. capitis		99.50	0.53	ESL 1			
Staphylococcus carnosus subsp. carnosus		0.50	0.07				
503201	**Good identification**						**503201**
Staphylococcus capitis subsp. capitis		100.0	0.36	ESL 1			
503211	**Very good identification**						**503211**
Staphylococcus capitis subsp. capitis		100.0	0.59	ESL 1			
503240	**Low discrimination**						**503240**
Staphylococcus haemolyticus		48.40	0.35	ESL 1			
Staphylococcus petrasii subsp. jettensis		42.68	0.18	TRE 93			
Staphylococcus auricularis		8.92	0.19				
503242	**Low discrimination**						**503242**
					PYR		
Staphylococcus haemolyticus		75.50	0.30	ESL 1	98		
Staphylococcus petrasii subsp. jettensis		20.89	0.02	TRE 93	99		
Staphylococcus epidermidis		3.62	0.01		1		
503250	**Good identification**						**503250**
Staphylococcus haemolyticus		100.0	0.30	ESL 1			
503340	**Low discrimination**						**503340**
					PYR		
Staphylococcus haemolyticus		80.68	0.36	ESL 1	98		
Rothia mucilaginosa		13.08	0.08	TRE 93	75		
Staphylococcus epidermidis		3.45	0.06		1		
503342	**Low discrimination**						**503342**
					PYR		
Staphylococcus haemolyticus		80.60	0.31	ESL 1	98		
Staphylococcus epidermidis		19.41	0.16	TRE 93	1		
503350	**Good identification**						**503350**
Staphylococcus haemolyticus		100.0	0.31	ESL 1			
503402	**Low discrimination**						**503402**
Staphylococcus piscifermentans		72.11	0.32	VPT 1			
Staphylococcus auricularis		27.89	0.14	URE 90			
503440	**Low discrimination**						**503440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		65.59	0.41	VPT 12		
Staphylococcus petrasii subsp. jettensis		25.19	0.18	ESL 1		
Staphylococcus piscifermentans		5.39	0.28			
503442	**Low discrimination**					**503442**
Staphylococcus piscifermentans		36.77	0.32	VPT 1		
Staphylococcus auricularis		33.18	0.22	URE 90		
Staphylococcus petrasii subsp. jettensis		20.24	0.02			
503500	**Low discrimination**					**503500**
Staphylococcus piscifermentans		60.10	0.32	VPT 1		
Staphylococcus auricularis		23.24	0.14	URE 90		
Rothia mucilaginosa		16.66	0.00			
503502	**Good identification**					**503502**
Staphylococcus piscifermentans		100.0	0.36	VPT 1		
503512	**Good identification**					**503512**
Staphylococcus piscifermentans		100.0	0.28	VPT 1		
503552	**Low discrimination**					**503552**
Staphylococcus piscifermentans		78.43	0.28	VPT 1		
Staphylococcus haemolyticus		21.57	0.08	URE 90		
503600	**Low discrimination**					**503600**
Staphylococcus auricularis		39.75	0.33	VPT 12		
Staphylococcus petrasii subsp. jettensis		38.58	0.18	ESL 1		
Staphylococcus piscifermentans		12.37	0.32			
503602	**Low discrimination**					**503602**
Staphylococcus piscifermentans		56.09	0.36	VPT 1		
Staphylococcus petrasii subsp. jettensis		20.58	0.02	URE 90		
Staphylococcus auricularis		13.35	0.13			
503612	**Low discrimination**					**503612**
Staphylococcus piscifermentans		80.39	0.28	VPT 1		
Staphylococcus haemolyticus		19.61	0.07	URE 90		
503640	**Low discrimination**					**503640**
Staphylococcus petrasii subsp. jettensis		83.45	0.59	ESL 1		
Staphylococcus haemolyticus		12.70	0.58			
Staphylococcus auricularis		2.03	0.40			
503642	**Low discrimination**					**503642**
Staphylococcus petrasii subsp. jettensis		65.45	0.43	ESL 1		
Staphylococcus haemolyticus		31.75	0.53			
Staphylococcus piscifermentans		1.80	0.36			
503650	**Low discrimination**					**503650**
Staphylococcus haemolyticus		88.61	0.53	ESL 1		
Staphylococcus petrasii subsp. jettensis		10.01	0.18			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Staphylococcus piscifermentans	1.38	0.25			
503652	**Acceptable identification**					**503652**
	Staphylococcus haemolyticus	92.86	0.48	ESL 1		
	Staphylococcus piscifermentans	3.85	0.28			
	Staphylococcus petrasii subsp. jettensis	3.29	0.02			
503700	**Acceptable identification**					**503700**
	Rothia mucilaginosa	91.98	0.42	ARG 1		
	Staphylococcus piscifermentans	5.03	0.36			
	Staphylococcus haemolyticus	1.79	0.18			
503701	**Good identification**					**503701**
					OXI	
	Rothia mucilaginosa	99.89	0.51	ARG 1	1	
	Kocuria kristinae	0.11	-.06		99	
503702	**Low discrimination**					**503702**
	Staphylococcus piscifermentans	79.56	0.39	VPT 1		
	Staphylococcus haemolyticus	10.65	0.13	URE 90		
	Rothia mucilaginosa	9.80	0.00			
503710	**Low discrimination**					**503710**
	Staphylococcus piscifermentans	52.08	0.28	VPT 1		
	Staphylococcus haemolyticus	25.47	0.13	URE 90		
	Rothia mucilaginosa	22.45	0.00			
503712	**Low discrimination**					**503712**
	Staphylococcus piscifermentans	84.51	0.32	VPT 1		
	Staphylococcus haemolyticus	15.50	0.08	URE 90		
503713	**Good identification**					**503713**
	Staphylococcus lentus	100.0	0.30	VPT 7		
503733	**Good identification**					**503733**
	Staphylococcus lentus	100.0	0.30	VPT 7		
503740	**Low discrimination**					**503740**
	Rothia mucilaginosa	51.88	0.49	ARG 1		
	Staphylococcus haemolyticus	42.95	0.59			
	Staphylococcus petrasii subsp. jettensis	2.53	0.18			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
503741	**Very good identification**						**503741**
Rothia mucilaginosa		98.29	0.59	ARG 1			
Staphylococcus haemolyticus		0.84	0.28				
Staphylococcus petrasii subsp. petrasii		0.42	0.14				
503742	**Low discrimination**						**503742**
					OXI	NOV	
Staphylococcus haemolyticus		86.14	0.54	ESL 1	1	2	
Staphylococcus piscifermentans		6.50	0.39		1	1	
Macrococcus caseolyticus		3.64	0.14		99	99	
503750	**Very good identification**						**503750**
Staphylococcus haemolyticus		96.02	0.54	ESL 1			
Rothia mucilaginosa		2.00	0.08				
Staphylococcus piscifermentans		1.98	0.28				
503752	**Acceptable identification**						**503752**
Staphylococcus haemolyticus		94.78	0.49	ESL 1			
Staphylococcus piscifermentans		5.22	0.32				
503753	**Low discrimination**						**503753**
					OXI	NOV	
Staphylococcus lentus		86.12	0.36	VPT 7	99	99	
Staphylococcus haemolyticus		11.80	0.18	ARG 1	1	2	
Staphylococcus caprae		2.09	-.11		1	1	
503773	**Good identification**						**503773**
Staphylococcus lentus		100.0	0.36	VPT 7			
504001	**Very good identification**						**504001**
Staphylococcus schleiferi subsp. schleiferi		96.90	0.67	NIT 90			
Staphylococcus capitis subsp. capitis		2.82	0.34				
Staphylococcus carnosus subsp. carnosus		0.28	0.15				
504003	**Low discrimination**						**504003**
Staphylococcus carnosus subsp. carnosus		52.02	0.27	BGA 90			
Staphylococcus schleiferi subsp. schleiferi		45.32	0.26	NIT 99			
Staphylococcus capitis subsp. capitis		2.66	-.01				
504011	**Good identification**						**504011**
Staphylococcus capitis subsp. capitis		94.68	0.57	NIT 89			
Staphylococcus carnosus subsp. carnosus		2.84	0.27	PHS 11			
Staphylococcus schleiferi subsp. schleiferi		2.48	0.26				
504013	**Low discrimination**						**504013**
Staphylococcus carnosus subsp. carnosus		85.48	0.39	BGA 90			
Staphylococcus capitis subsp. capitis		14.52	0.22	NIT 99			
504101	**Very good identification**						**504101**
Staphylococcus schleiferi subsp. schleiferi		99.93	0.67	NIT 90			
Staphylococcus carnosus subsp. carnosus		0.07	0.02				
504113	**Low discrimination**						**504113**
					COA		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		81.01	0.27	BGA 90	1		
Staphylococcus caprae		9.71	0.00	NIT 99	1		
Staphylococcus schleiferi subsp. coagulans		9.28	0.12		99		
504153	**Good identification**						**504153**
Staphylococcus caprae		100.0	0.41	URE 90			
504201	**Low discrimination**						**504201**
Staphylococcus capitis subsp. capitis		84.24	0.40	NIT 89			
Staphylococcus schleiferi subsp. schleiferi		15.76	0.26	PHS 11			
504211	**Very good identification**						**504211**
Staphylococcus capitis subsp. capitis		100.0	0.63	NIT 89			
504213	**Good identification**						**504213**
Staphylococcus capitis subsp. capitis		96.90	0.28	NIT 89			
Staphylococcus carnosus subsp. carnosus		3.10	-.02	PHS 11			
504240	**Very good identification**						**504240**
					PYR	COA	
Staphylococcus aureus subsp. anaerobius		99.19	0.59	VPT 1	1	99	
Staphylococcus petrasii subsp. jettensis		0.47	0.10		99	1	
Staphylococcus epidermidis		0.18	0.16		1	1	
504243	**Good identification**						**504243**
Staphylococcus epidermidis		100.0	0.32	URE 92			
504340	**Low discrimination**						**504340**
					COA		
Staphylococcus epidermidis		50.91	0.31	URE 92	1		
Staphylococcus aureus subsp. anaerobius		49.09	0.18	NIT 97	99		
504341	**Good identification**						**504341**
Staphylococcus epidermidis		100.0	0.37	URE 92			
504342	**Good identification**						**504342**
Staphylococcus epidermidis		99.12	0.42	URE 92			
Staphylococcus caprae		0.88	-.12	NIT 97			
504343	**Good identification**						**504343**
Staphylococcus epidermidis		96.62	0.47	URE 92			
Staphylococcus caprae		3.38	0.06	NIT 97			
504353	**Good identification**						**504353**
Staphylococcus caprae		98.51	0.32	URE 90			
Staphylococcus epidermidis		1.50	0.06	NIT 90			
504400	**Low discrimination**						**504400**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		78.05	0.36	NIT 90	1		
Staphylococcus cohnii subsp. cohnii		9.60	0.24	MNS 99	99		
Staphylococcus carnosus subsp. utilis		7.97	0.03		1		
504401	**Very good identification**						**504401**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		99.81	0.77	NIT 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. cohnii		0.13	0.25		99		
Staphylococcus carnosus subsp. carnosus		0.04	0.07		1		
504403	**Low discrimination**						**504403**
Staphylococcus schleiferi subsp. schleiferi		85.91	0.36	NIT 90			
Staphylococcus carnosus subsp. carnosus		14.09	0.19	LAC 1			
504410	**Good identification**						**504410**
Staphylococcus cohnii subsp. cohnii		100.0	0.27	ARG 1			
504411	**Low discrimination**						**504411**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		62.75	0.36	NIT 90	1		
Staphylococcus capitis subsp. capitis		16.32	0.22	MAN 1	1		
Staphylococcus cohnii subsp. cohnii		10.64	0.27		99		
504413	**Good identification**						**504413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	BGA 90			
504421	**Good identification**						**504421**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	NIT 90			
504440	**Low discrimination**						**504440**
					NOV	PYR	
Staphylococcus cohnii subsp. cohnii		49.96	0.35	ARG 1	99	1	
Staphylococcus petrasii subsp. jettensis		24.85	0.10	PHS 9	1	99	
Staphylococcus auricularis		16.78	0.22		1	40	
504441	**Low discrimination**						**504441**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		69.61	0.36	NIT 90	1		
Staphylococcus cohnii subsp. cohnii		29.37	0.35	MLT 1	99		
Staphylococcus auricularis		1.01	0.02		1		
504450	**Low discrimination**						**504450**
					YEP		
Staphylococcus cohnii subsp. cohnii		87.12	0.37	ARG 1	12		
Staphylococcus kloosii		12.88	0.32	PHS 9	99		
504451	**Good identification**						**504451**
					NOV		
Staphylococcus cohnii subsp. cohnii		96.11	0.38	ARG 1	99		
Staphylococcus caprae		3.89	-.07	PHS 9	1		
504452	**Low discrimination**						**504452**
					NOV	YEP	
Staphylococcus kloosii		58.51	0.33	VPT 10	99	99	
Staphylococcus caprae		41.49	0.02	ARG 1	1	1	
504500	**Good identification**						**504500**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		98.65	0.36	NIT 90	1		
Staphylococcus cohnii subsp. cohnii		1.35	0.05	MNS 99	99		
504501	**Very good identification**						**504501**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		99.99	0.77	NIT 90	1		
Staphylococcus cohnii subsp. cohnii		0.02	0.05		99		
504503	**Acceptable identification**						**504503**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. schleiferi		94.10	0.36	NIT 90			
Staphylococcus carnosus subsp. carnosus		3.86	0.07	LAC 1			
Staphylococcus caprae		2.04	-.07				
504511	**Acceptable identification**						**504511**
Staphylococcus schleiferi subsp. schleiferi		92.46	0.36	NIT 90			
Staphylococcus carnosus subsp. carnosus		3.79	0.07	MAN 1			
Staphylococcus caprae		2.01	-.07				
504521	**Good identification**						**504521**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	NIT 90			
504541	**Low discrimination**						**504541**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		86.20	0.36	NIT 90	1		
Staphylococcus caprae		9.76	0.08	MLT 1	1		
Staphylococcus cohnii subsp. cohnii		4.04	0.16		99		
504543	**Good identification**						**504543**
Staphylococcus caprae		100.0	0.34	URE 90			
504551	**Good identification**						**504551**
					NOV		
Staphylococcus caprae		97.30	0.34	URE 90	1		
Staphylococcus cohnii subsp. cohnii		2.70	0.18	NIT 90	99		
504552	**Good identification**						**504552**
					NOV	YEP	
Staphylococcus caprae		97.31	0.43	URE 90	1	1	
Staphylococcus kloosii		2.69	0.39	NIT 90	99	99	
504553	**Very good identification**						**504553**
Staphylococcus caprae		100.0	0.61	URE 90			
504601	**Acceptable identification**						**504601**
					OXI	PYR	
Staphylococcus schleiferi subsp. schleiferi		94.63	0.36	NIT 90	1	37	
Staphylococcus capitis subsp. capitis		3.44	0.05	SUC 1	1	1	
Kocuria kristinae		1.93	-.02		99	93	
504611	**Good identification**						**504611**
Staphylococcus capitis subsp. capitis		100.0	0.28	NIT 89			
504640	**Low discrimination**						**504640**
					PYR	COA	
Staphylococcus petrasii subsp. jettensis		72.94	0.51	PHS 1	99	1	
Staphylococcus petrasii subsp. pragensis		24.70	0.39		99	1	
Staphylococcus aureus subsp. anaerobius		1.57	0.18		1	99	
504642	**Good identification**						**504642**
Staphylococcus petrasii subsp. jettensis		96.64	0.36	PHS 1			
Staphylococcus petrasii subsp. pragensis		1.87	-.02				
Staphylococcus haemolyticus		1.11	0.12				
504701	**Good identification**						**504701**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	NIT 90			
504752	**Good identification**						**504752**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus caprae		98.06	0.34	URE 90	1	1	
Staphylococcus haemolyticus		1.22	0.08	NIT 90	2	20	
Staphylococcus kloosii		0.73	0.18	MNS 88	99	99	
504753	**Good identification**						**504753**
					COA		
Staphylococcus caprae		96.25	0.52	URE 90	1		
Staphylococcus aureus subsp. aureus		3.75	0.18	NIT 90	99		
505401	**Good identification**						**505401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	ESL 1			
505450	**Good identification**						**505450**
Staphylococcus kloosii		100.0	0.32	VPT 10			
505452	**Good identification**						**505452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
505501	**Good identification**						**505501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	ESL 1			
505550	**Good identification**						**505550**
Staphylococcus kloosii		100.0	0.38	VPT 10			
505552	**Low discrimination**						**505552**
					NOV	YEP	
Staphylococcus kloosii		73.24	0.39	VPT 10	99	99	
Staphylococcus caprae		26.76	0.02	ARG 1	1	1	
505572	**Good identification**						**505572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
506000	**Low discrimination**						**506000**
Staphylococcus schleiferi subsp. schleiferi		88.24	0.46	MNS 99			
Staphylococcus carnosus subsp. utilis		3.00	0.03				
Staphylococcus carnosus subsp. carnosus		2.81	0.15				
506001	**Good identification**						**506001**
Staphylococcus schleiferi subsp. schleiferi		94.52	0.87				
Staphylococcus carnosus subsp. carnosus		3.01	0.56				
Staphylococcus capitis subsp. capitis		2.47	0.53				
506002	**Acceptable identification**						**506002**
Staphylococcus carnosus subsp. carnosus		92.66	0.27	BGA 90			
Staphylococcus schleiferi subsp. schleiferi		7.34	0.04	MNS 99			
506003	**Good identification**						**506003**
Staphylococcus carnosus subsp. carnosus		92.20	0.68	BGA 90			
Staphylococcus schleiferi subsp. schleiferi		7.30	0.46				
Staphylococcus capitis subsp. capitis		0.39	0.18				
506010	**Low discrimination**						**506010**
Staphylococcus capitis subsp. capitis		71.61	0.35	PHS 11			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		26.31	0.27	MNS 99		
Staphylococcus schleiferi subsp. schleiferi		2.08	0.04			
506011	**Low discrimination**					**506011**
Staphylococcus capitis subsp. capitis		71.61	0.76	PHS 11		
Staphylococcus carnosus subsp. carnosus		26.31	0.68			
Staphylococcus schleiferi subsp. schleiferi		2.08	0.46			
506012	**Good identification**					**506012**
Staphylococcus carnosus subsp. carnosus		98.63	0.39	BGA 90		
Staphylococcus capitis subsp. capitis		1.37	0.00	MNS 99		
506013	**Very good identification**					**506013**
					COA	
Staphylococcus carnosus subsp. carnosus		98.58	0.80	BGA 90	1	
Staphylococcus capitis subsp. capitis		1.37	0.41		1	
Staphylococcus schleiferi subsp. coagulans		0.03	0.12		99	
506021	**Acceptable identification**					**506021**
Staphylococcus schleiferi subsp. schleiferi		94.52	0.46	XYL 1		
Staphylococcus carnosus subsp. carnosus		3.01	0.15			
Staphylococcus capitis subsp. capitis		2.47	0.12			
506023	**Acceptable identification**					**506023**
Staphylococcus carnosus subsp. carnosus		92.66	0.27	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		7.34	0.04	XYL 1		
506031	**Low discrimination**					**506031**
Staphylococcus capitis subsp. capitis		71.61	0.35	PHS 11		
Staphylococcus carnosus subsp. carnosus		26.31	0.27	XYL 1		
Staphylococcus schleiferi subsp. schleiferi		2.08	0.04			
506033	**Good identification**					**506033**
Staphylococcus carnosus subsp. carnosus		98.63	0.39	BGA 90		
Staphylococcus capitis subsp. capitis		1.37	0.00	XYL 1		
506041	**Acceptable identification**					**506041**
Staphylococcus schleiferi subsp. schleiferi		93.17	0.46	MLT 1		
Staphylococcus carnosus subsp. carnosus		2.97	0.15			
Staphylococcus capitis subsp. capitis		2.44	0.12			
506043	**Low discrimination**					**506043**
Staphylococcus carnosus subsp. carnosus		65.97	0.27	BGA 90		
Staphylococcus epidermidis		25.06	0.22	MLT 1		
Staphylococcus schleiferi subsp. schleiferi		5.22	0.04			
506051	**Low discrimination**					**506051**
Staphylococcus capitis subsp. capitis		70.55	0.35	PHS 11		
Staphylococcus carnosus subsp. carnosus		25.92	0.27	MLT 1		
Staphylococcus schleiferi subsp. schleiferi		2.05	0.04			
506053	**Low discrimination**					**506053**
Staphylococcus carnosus subsp. carnosus		77.20	0.39	BGA 90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus caprae		20.81	0.20	MLT 1		
Staphylococcus capitis subsp. capitis		1.07	0.00			
506100	**Good identification**					**506100**
Staphylococcus schleiferi subsp. schleiferi		99.21	0.46	MNS 99		
Staphylococcus carnosus subsp. carnosus		0.79	0.02			
506101	**Very good identification**					**506101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		98.93	0.87		1	
Staphylococcus carnosus subsp. carnosus		0.79	0.43		1	
Staphylococcus schleiferi subsp. coagulans		0.26	0.37		99	
506103	**Low discrimination**					**506103**
					COA	
Staphylococcus carnosus subsp. carnosus		55.08	0.56	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		27.35	0.53		99	
Staphylococcus schleiferi subsp. schleiferi		17.45	0.46		1	
506111	**Low discrimination**					**506111**
Staphylococcus carnosus subsp. carnosus		66.51	0.56	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		21.07	0.46			
Staphylococcus capitis subsp. capitis		7.31	0.35			
506112	**Low discrimination**					**506112**
					COA	
Staphylococcus carnosus subsp. carnosus		79.26	0.27	BGA 90	1	
Staphylococcus caprae		11.66	0.02	MNS 99	1	
Staphylococcus schleiferi subsp. coagulans		9.08	0.12		99	
506113	**Low discrimination**					**506113**
					COA	
Staphylococcus carnosus subsp. carnosus		88.74	0.68	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		10.17	0.53		99	
Staphylococcus caprae		0.97	0.20		1	
506121	**Good identification**					**506121**
Staphylococcus schleiferi subsp. schleiferi		99.21	0.46	XYL 1		
Staphylococcus carnosus subsp. carnosus		0.79	0.02			
506133	**Low discrimination**					**506133**
					COA	
Staphylococcus carnosus subsp. carnosus		89.72	0.27	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		10.28	0.12	XYL 1	99	
506141	**Low discrimination**					**506141**
Staphylococcus schleiferi subsp. schleiferi		88.31	0.46	MLT 1		
Staphylococcus epidermidis		7.66	0.27			
Staphylococcus caprae		3.33	0.08			
506142	**Low discrimination**					**506142**
Staphylococcus epidermidis		60.21	0.32	URE 92		
Staphylococcus caprae		39.79	0.17	SUC 99		
506143	**Low discrimination**					**506143**
Staphylococcus caprae		68.31	0.34	URE 90		
Staphylococcus epidermidis		26.18	0.37	TRE 90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		3.04	0.15	MAN 95		
506151	**Acceptable identification**					**506151**
Staphylococcus caprae		94.47	0.34	URE 90		
Staphylococcus carnosus subsp. carnosus		4.20	0.15	TRE 90		
Staphylococcus schleiferi subsp. schleiferi		1.33	0.04	LAC 95		
506152	**Good identification**					**506152**
Staphylococcus caprae		100.0	0.43	URE 90		
506153	**Very good identification**					**506153**
					COA	
Staphylococcus caprae		98.97	0.61	URE 90	1	
Staphylococcus carnosus subsp. carnosus		0.93	0.27	TRE 90	1	
Staphylococcus schleiferi subsp. coagulans		0.11	0.12		99	
506201	**Low discrimination**					**506201**
Staphylococcus capitis subsp. capitis		82.11	0.58	PHS 11		
Staphylococcus schleiferi subsp. schleiferi		17.09	0.46	MAN 93		
Staphylococcus carnosus subsp. carnosus		0.55	0.15			
506203	**Low discrimination**					**506203**
Staphylococcus carnosus subsp. carnosus		42.62	0.27	BGA 90		
Staphylococcus capitis subsp. capitis		32.76	0.23	SUC 1		
Staphylococcus epidermidis		16.19	0.22			
506210	**Good identification**					**506210**
Staphylococcus capitis subsp. capitis		100.0	0.40	PHS 11		
506211	**Very good identification**					**506211**
Staphylococcus capitis subsp. capitis		99.79	0.81	PHS 11		
Staphylococcus carnosus subsp. carnosus		0.20	0.27			
Staphylococcus schleiferi subsp. schleiferi		0.02	0.04			
506213	**Low discrimination**					**506213**
Staphylococcus capitis subsp. capitis		71.44	0.47	PHS 11		
Staphylococcus carnosus subsp. carnosus		27.98	0.39	LAC 2		
Staphylococcus capitis subsp. ureolyticus		0.33	-.11			
506231	**Good identification**					**506231**
Staphylococcus capitis subsp. capitis		100.0	0.40	PHS 11		
506240	**Low discrimination**					**506240**
					PYR	
Staphylococcus epidermidis		62.43	0.47	URE 92	1	
Staphylococcus haemolyticus		13.08	0.35		98	
Staphylococcus petrasii subsp. jettensis		11.53	0.18		99	
506241	**Good identification**					**506241**
Staphylococcus epidermidis		95.95	0.52	URE 92		
Staphylococcus capitis subsp. capitis		3.07	0.17			
Staphylococcus schleiferi subsp. schleiferi		0.64	0.04			
506242	**Very good identification**					**506242**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		95.47	0.57	URE 92	1	
Staphylococcus haemolyticus		3.55	0.30		98	
Staphylococcus petrasii subsp. jettensis		0.98	0.02		99	
506243	**Very good identification**					**506243**
Staphylococcus epidermidis		99.87	0.63	URE 92		
Staphylococcus capitis subsp. ureolyticus		0.13	-.11			
506250	**Low discrimination**					**506250**
					PYR	
Staphylococcus haemolyticus		87.18	0.30	PHS 1	98	
Staphylococcus epidermidis		7.16	0.06	TRE 93	1	
Staphylococcus capitis subsp. capitis		5.66	-.01		1	
506251	**Acceptable identification**					**506251**
Staphylococcus capitis subsp. capitis		94.83	0.40	PHS 11		
Staphylococcus capitis subsp. ureolyticus		2.92	0.00	MLT 1		
Staphylococcus epidermidis		2.25	0.11			
506301	**Low discrimination**					**506301**
					COA	
Staphylococcus schleiferi subsp. schleiferi		81.04	0.46	SUC 1	1	
Staphylococcus epidermidis		7.03	0.27		1	
Staphylococcus schleiferi subsp. coagulans		6.53	0.27		99	
506302	**Good identification**					**506302**
					COA	
Staphylococcus epidermidis		96.73	0.32	URE 92	1	
Staphylococcus schleiferi subsp. coagulans		3.27	0.02	MLT 99	99	
506303	**Low discrimination**					**506303**
					COA	
Staphylococcus schleiferi subsp. coagulans		53.12	0.43	URE 99	99	
Staphylococcus epidermidis		29.47	0.37		1	
Staphylococcus hyicus		12.46	0.22		24	
506311	**Low discrimination**					**506311**
					COA	
Staphylococcus capitis subsp. capitis		84.71	0.40	PHS 11	1	
Staphylococcus schleiferi subsp. coagulans		9.77	0.26	GAL 1	99	
Staphylococcus carnosus subsp. carnosus		4.19	0.15		1	
506313	**Low discrimination**					**506313**
					COA	
Staphylococcus schleiferi subsp. coagulans		70.46	0.42	URE 99	99	
Staphylococcus carnosus subsp. carnosus		19.67	0.27		1	
Staphylococcus caprae		7.85	0.11		1	
506340	**Very good identification**					**506340**
					PYR	
Staphylococcus epidermidis		96.00	0.62	URE 92	1	
Staphylococcus haemolyticus		4.00	0.36		98	
506341	**Very good identification**					**506341**
Staphylococcus epidermidis		99.65	0.68	URE 92		
Staphylococcus caprae		0.16	-.01			
Staphylococcus schleiferi subsp. schleiferi		0.12	0.04			
506342	**Very good identification**					**506342**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		99.02	0.73	URE 92	1	
Staphylococcus haemolyticus		0.73	0.31		98	
Staphylococcus caprae		0.25	0.08		75	
506343	**Very good identification**					**506343**
					COA	
Staphylococcus epidermidis		99.00	0.78	URE 92	1	
Staphylococcus caprae		0.97	0.26		1	
Staphylococcus schleiferi subsp. coagulans		0.02	0.02		99	
506350	**Low discrimination**					**506350**
					PYR	
Staphylococcus haemolyticus		57.73	0.31	PHS 1	98	
Staphylococcus epidermidis		23.82	0.21	TRE 93	1	
Staphylococcus caprae		18.44	0.08		75	
506352	**Low discrimination**					**506352**
					PYR	
Staphylococcus caprae		76.45	0.34	URE 90	75	
Staphylococcus epidermidis		16.46	0.32	TRE 90	1	
Staphylococcus haemolyticus		7.09	0.26	MNS 88	98	
506353	**Low discrimination**					**506353**
					COA	
Staphylococcus caprae		89.34	0.52	URE 90	1	
Staphylococcus epidermidis		4.87	0.37	TRE 90	1	
Staphylococcus aureus subsp. aureus		3.48	0.18		99	
506361	**Good identification**					**506361**
Staphylococcus epidermidis		100.0	0.27	URE 92		
506362	**Good identification**					**506362**
Staphylococcus epidermidis		100.0	0.32	URE 92		
506363	**Good identification**					**506363**
Staphylococcus epidermidis		100.0	0.37	URE 92		
506400	**Low discrimination**					**506400**
Staphylococcus schleiferi subsp. schleiferi		89.27	0.55	MNS 99		
Staphylococcus carnosus subsp. utilis		5.32	0.18			
Staphylococcus auricularis		5.00	0.34			
506401	**Excellent identification**					**506401**
Staphylococcus schleiferi subsp. schleiferi		99.52	0.96			
Staphylococcus carnosus subsp. carnosus		0.45	0.48			
Staphylococcus capitis subsp. capitis		0.02	0.18			
506403	**Low discrimination**					**506403**
Staphylococcus carnosus subsp. carnosus		64.34	0.60	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		35.66	0.55			
506411	**Low discrimination**					**506411**
Staphylococcus carnosus subsp. carnosus		59.36	0.60	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		32.90	0.55			
Staphylococcus capitis subsp. capitis		7.69	0.41			
506412	**Good identification**					**506412**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		100.0	0.32	BGA 90		
506413	**Very good identification**					**506413**
Staphylococcus carnosus subsp. carnosus		99.74	0.73	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		0.14	0.14			
Staphylococcus capitis subsp. capitis		0.07	0.06			
506421	**Very good identification**					**506421**
Staphylococcus schleiferi subsp. schleiferi		99.55	0.55	XYL 1		
Staphylococcus carnosus subsp. carnosus		0.45	0.07			
506433	**Good identification**					**506433**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	BGA 90		
506440	**Low discrimination**					**506440**
Staphylococcus auricularis		64.74	0.41	VPT 12		
Staphylococcus petrasii subsp. jettensis		24.86	0.18	PHS 1		
Staphylococcus schleiferi subsp. schleiferi		5.00	0.14			
506441	**Very good identification**					**506441**
Staphylococcus schleiferi subsp. schleiferi		97.79	0.55	MLT 1		
Staphylococcus auricularis		1.42	0.22			
Staphylococcus carnosus subsp. carnosus		0.45	0.07			
506453	**Low discrimination**					**506453**
					COA	
Staphylococcus caprae		84.76	0.39	URE 90	1	
Staphylococcus carnosus subsp. carnosus		14.97	0.32	GAL 99	1	
Staphylococcus aureus subsp. aureus		0.27	-.17		99	
506500	**Very good identification**					**506500**
Staphylococcus schleiferi subsp. schleiferi		99.38	0.55	MNS 99		
Staphylococcus auricularis		0.62	0.14			
506501	**Excellent identification**					**506501**
Staphylococcus schleiferi subsp. schleiferi		99.89	0.96			
Staphylococcus carnosus subsp. carnosus		0.11	0.36			
506503	**Low discrimination**					**506503**
Staphylococcus schleiferi subsp. schleiferi		67.45	0.55	LAC 1		
Staphylococcus carnosus subsp. carnosus		30.42	0.48			
Staphylococcus caprae		1.47	0.13			
506511	**Low discrimination**					**506511**
Staphylococcus schleiferi subsp. schleiferi		67.79	0.55	MAN 1		
Staphylococcus carnosus subsp. carnosus		30.58	0.48			
Staphylococcus caprae		1.47	0.13			
506513	**Low discrimination**					**506513**
Staphylococcus carnosus subsp. carnosus		80.74	0.60	BGA 90		
Staphylococcus caprae		18.47	0.39			
Staphylococcus schleiferi subsp. schleiferi		0.45	0.14			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
506521	**Very good identification**						**506521**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.55	XYL 1			
506541	**Low discrimination**						**506541**
Staphylococcus schleiferi subsp. schleiferi		89.70	0.55	MLT 1			
Staphylococcus caprae		10.15	0.28				
Staphylococcus auricularis		0.15	0.02				
506542	**Low discrimination**						**506542**
					OXI	NOV	
Staphylococcus caprae		88.36	0.36	URE 90	1	1	
Macrococcus caseolyticus		9.16	0.18	MAN 95	99	99	
Staphylococcus haemolyticus		1.46	0.13	MNS 88	1	2	
506543	**Very good identification**						**506543**
Staphylococcus caprae		99.32	0.54	URE 90			
Staphylococcus schleiferi subsp. schleiferi		0.47	0.14	MAN 95			
Staphylococcus carnosus subsp. carnosus		0.21	0.07				
506550	**Good identification**						**506550**
Staphylococcus caprae		98.30	0.36	URE 90			
Staphylococcus haemolyticus		1.70	0.13	MNS 88			
506551	**Very good identification**						**506551**
Staphylococcus caprae		99.32	0.54	URE 90			
Staphylococcus schleiferi subsp. schleiferi		0.47	0.14	LAC 95			
Staphylococcus carnosus subsp. carnosus		0.21	0.07				
506552	**Very good identification**						**506552**
Staphylococcus caprae		99.95	0.63	URE 90			
Staphylococcus haemolyticus		0.05	0.08	MNS 88			
506553	**Very good identification**						**506553**
					COA		
Staphylococcus caprae		99.80	0.80	URE 90	1		
Staphylococcus aureus subsp. aureus		0.16	0.18		99		
Staphylococcus carnosus subsp. carnosus		0.05	0.19		1		
506573	**Good identification**						**506573**
Staphylococcus caprae		100.0	0.39	URE 90			
506600	**Low discrimination**						**506600**
Staphylococcus auricularis		43.22	0.33	VPT 12			
Staphylococcus petrasii subsp. jettensis		41.96	0.18	PHS 1			
Staphylococcus schleiferi subsp. schleiferi		8.44	0.14				
506601	**Very good identification**						**506601**
Staphylococcus schleiferi subsp. schleiferi		95.57	0.55	SUC 1			
Staphylococcus capitis subsp. capitis		3.12	0.23				
Staphylococcus auricularis		0.55	0.13				
506611	**Acceptable identification**						**506611**
Staphylococcus capitis subsp. capitis		93.88	0.47	PHS 11			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		3.94	0.19	TRE 2			
Staphylococcus schleiferi subsp. schleiferi		2.18	0.14				
506613	**Low discrimination**						**506613**
Staphylococcus carnosus subsp. carnosus		82.30	0.32	BGA 90			
Staphylococcus capitis subsp. capitis		10.01	0.12	SUC 1			
Staphylococcus simulans		2.63	0.04				
506640	**Low discrimination**						**506640**
Staphylococcus petrasii subsp. jettensis		83.75	0.59	PHS 1			
Staphylococcus haemolyticus		12.75	0.58				
Staphylococcus auricularis		2.03	0.40				
506642	**Low discrimination**						**506642**
Staphylococcus petrasii subsp. jettensis		65.98	0.43	PHS 1			
Staphylococcus haemolyticus		32.01	0.53				
Staphylococcus auricularis		1.01	0.21				
506650	**Low discrimination**						**506650**
Staphylococcus haemolyticus		89.85	0.53	PHS 1			
Staphylococcus petrasii subsp. jettensis		10.15	0.18				
506652	**Low discrimination**						**506652**
Staphylococcus haemolyticus		88.96	0.48	PHS 1			
Staphylococcus caprae		7.41	0.13				
Staphylococcus petrasii subsp. jettensis		3.15	0.02				
506653	**Low discrimination**						**506653**
					COA	PYR	
Staphylococcus caprae		52.17	0.30	URE 90	1	75	
Staphylococcus aureus subsp. aureus		45.19	0.24	GAL 99	99	10	
Staphylococcus haemolyticus		2.64	0.17		1	98	
506701	**Very good identification**						**506701**
					YEP		
Staphylococcus schleiferi subsp. schleiferi		95.97	0.55	SUC 1	1		
Staphylococcus hyicus		3.71	0.22		1		
Staphylococcus chromogenes		0.17	-.02		95		
506703	**Acceptable identification**						**506703**
					YEP		
Staphylococcus hyicus		91.64	0.44	VPT 1	1		
Staphylococcus chromogenes		3.35	0.18		95		
Staphylococcus schleiferi subsp. schleiferi		2.08	0.14		1		
506713	**Low discrimination**						**506713**
					COA		
Staphylococcus caprae		60.50	0.30	URE 90	1		
Staphylococcus aureus subsp. aureus		25.93	0.18	MLT 99	99		
Staphylococcus carnosus subsp. carnosus		7.22	0.19		1		
506740	**Low discrimination**						**506740**
					PYR		
Staphylococcus haemolyticus		89.11	0.59	PHS 1	98		
Staphylococcus petrasii subsp. jettensis		5.25	0.18		99		
Staphylococcus epidermidis		1.63	0.21		1		
506742	**Low discrimination**						**506742**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR		
Staphylococcus haemolyticus		71.21	0.54	PHS 1	98		
Staphylococcus caprae		16.09	0.27		75		
Staphylococcus epidermidis		7.31	0.32		1		
506743	**Low discrimination**						**506743**
					COA		
Staphylococcus caprae		79.93	0.45	URE 90	1		
Staphylococcus epidermidis		9.20	0.37	MAN 95	1		
Staphylococcus aureus subsp. aureus		6.58	0.18		99		
506750	**Low discrimination**						**506750**
Staphylococcus haemolyticus		82.21	0.54	PHS 1			
Staphylococcus caprae		17.79	0.27				
506751	**Acceptable identification**						**506751**
					COA	PYR	
Staphylococcus caprae		90.61	0.45	URE 90	1	75	
Staphylococcus aureus subsp. aureus		7.45	0.18	LAC 95	99	10	
Staphylococcus haemolyticus		1.77	0.23		1	98	
506752	**Low discrimination**						**506752**
					COA	PYR	
Staphylococcus caprae		85.58	0.54	URE 90	1	75	
Staphylococcus haemolyticus		11.71	0.49	MNS 88	1	98	
Staphylococcus aureus subsp. aureus		2.72	0.18		99	10	
506753	**Low discrimination**						**506753**
					COA	PYR	
Staphylococcus caprae		69.95	0.72	URE 90	1	75	
Staphylococcus aureus subsp. aureus		29.99	0.59		99	10	
Staphylococcus haemolyticus		0.04	0.18		1	98	
506773	**Low discrimination**						**506773**
					COA		
Staphylococcus caprae		70.00	0.30	URE 90	1		
Staphylococcus aureus subsp. aureus		30.00	0.18	XYL 1	99		
507001	**Low discrimination**						**507001**
Staphylococcus schleiferi subsp. schleiferi		72.63	0.46	ESL 1			
Staphylococcus carnosus subsp. carnosus		25.48	0.36				
Staphylococcus capitis subsp. capitis		1.90	0.12				
507003	**Good identification**						**507003**
					COA		
Staphylococcus carnosus subsp. carnosus		98.23	0.48	BGA 90	1		
Staphylococcus schleiferi subsp. coagulans		1.07	0.12	ESL 10	99		
Staphylococcus schleiferi subsp. schleiferi		0.71	0.04		1		
507011	**Low discrimination**						**507011**
Staphylococcus carnosus subsp. carnosus		79.70	0.48	BGA 90			
Staphylococcus capitis subsp. capitis		19.72	0.35	ESL 10			
Staphylococcus schleiferi subsp. schleiferi		0.57	0.04				
507013	**Very good identification**						**507013**
					COA		
Staphylococcus carnosus subsp. carnosus		99.63	0.61	BGA 90	1		
Staphylococcus schleiferi subsp. coagulans		0.25	0.11	ESL 10	99		
Staphylococcus capitis subsp. capitis		0.13	0.00		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
507101	**Low discrimination**					**507101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		75.03	0.46	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		18.40	0.37		99	
Staphylococcus carnosus subsp. carnosus		6.58	0.24		1	
507103	**Low discrimination**					**507103**
					COA	
Staphylococcus schleiferi subsp. coagulans		80.67	0.53	URE 99	99	
Staphylococcus carnosus subsp. carnosus		18.79	0.36		1	
Staphylococcus schleiferi subsp. schleiferi		0.54	0.04		1	
507111	**Low discrimination**					**507111**
					COA	
Staphylococcus carnosus subsp. carnosus		59.74	0.36	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		38.54	0.36	ESL 10	99	
Staphylococcus schleiferi subsp. schleiferi		1.72	0.04		1	
507113	**Low discrimination**					**507113**
					COA	
Staphylococcus carnosus subsp. carnosus		50.23	0.48	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		49.77	0.52	ESL 10	99	
507211	**Good identification**					**507211**
Staphylococcus capitis subsp. capitis		97.85	0.40	ESL 1		
Staphylococcus carnosus subsp. carnosus		2.15	0.07	PHS 11		
507301	**Low discrimination**					**507301**
					COA	
Staphylococcus schleiferi subsp. coagulans		88.46	0.27	URE 99	99	
Staphylococcus schleiferi subsp. schleiferi		11.54	0.04	LAC 86	1	
507303	**Good identification**					**507303**
Staphylococcus schleiferi subsp. coagulans		100.0	0.43	URE 99		
507313	**Good identification**					**507313**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.87	0.42	URE 99	99	
Staphylococcus carnosus subsp. carnosus		3.13	0.07		1	
507341	**Good identification**					**507341**
Staphylococcus epidermidis		100.0	0.27	URE 92		
507342	**Good identification**					**507342**
Staphylococcus epidermidis		100.0	0.32	URE 92		
507343	**Good identification**					**507343**
					COA	
Staphylococcus epidermidis		98.28	0.37	URE 92	1	
Staphylococcus schleiferi subsp. coagulans		1.72	0.02	ESL 1	99	
507401	**Very good identification**					**507401**
Staphylococcus schleiferi subsp. schleiferi		95.23	0.55	ESL 1		
Staphylococcus carnosus subsp. carnosus		4.77	0.28			
507402	**Low discrimination**					**507402**
Staphylococcus piscifermentans		84.14	0.28	VPT 1		
Staphylococcus carnosus subsp. carnosus		15.87	0.00	URE 90		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
507403	**Good identification**					**507403**
Staphylococcus carnosus subsp. carnosus		95.20	0.41	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		4.80	0.14	ESL 10		
507411	**Acceptable identification**					**507411**
Staphylococcus carnosus subsp. carnosus		94.15	0.41	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		4.74	0.14	ESL 10		
Staphylococcus capitis subsp. capitis		1.11	0.00			
507413	**Very good identification**					**507413**
Staphylococcus carnosus subsp. carnosus		100.0	0.53	BGA 90		
507442	**Good identification**					**507442**
Staphylococcus piscifermentans		100.0	0.28	VPT 1		
507500	**Low discrimination**					**507500**
Staphylococcus piscifermentans		51.53	0.28	VPT 1		
Staphylococcus schleiferi subsp. schleiferi		48.47	0.14	URE 90		
507501	**Very good identification**					**507501**
Staphylococcus schleiferi subsp. schleiferi		98.76	0.55	ESL 1		
Staphylococcus carnosus subsp. carnosus		1.24	0.16			
507502	**Good identification**					**507502**
Staphylococcus piscifermentans		100.0	0.32	VPT 1		
507503	**Low discrimination**					**507503**
					COA	
Staphylococcus carnosus subsp. carnosus		76.76	0.28	BGA 90	1	
Staphylococcus schleiferi subsp. schleiferi		15.47	0.14	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		7.77	0.12		99	
507511	**Low discrimination**					**507511**
Staphylococcus carnosus subsp. carnosus		83.23	0.28	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		16.78	0.14	ESL 10		
507513	**Good identification**					**507513**
					COA	
Staphylococcus carnosus subsp. carnosus		95.77	0.41	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		2.24	0.11	ESL 10	99	
Staphylococcus caprae		1.99	-.02		1	
507540	**Good identification**					**507540**
Staphylococcus piscifermentans		100.0	0.28	VPT 1		
507542	**Low discrimination**					**507542**
Staphylococcus piscifermentans		84.47	0.32	VPT 1		
Staphylococcus caprae		15.53	-.05	URE 90		
507553	**Good identification**					**507553**
Staphylococcus caprae		99.51	0.39	URE 90		
Staphylococcus carnosus subsp. carnosus		0.49	0.00	ESL 1		
507600	**Good identification**					**507600**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		100.0	0.28	VPT 1				
507602	**Good identification**							**507602**
Staphylococcus piscifermentans		100.0	0.32	VPT 1				
507642	**Low discrimination**							**507642**
Staphylococcus piscifermentans		55.03	0.32	VPT 1				
Staphylococcus petrasii subsp. jettensis		30.28	0.02	URE 90				
Staphylococcus haemolyticus		14.69	0.12					
507700	**Low discrimination**							**507700**
Staphylococcus piscifermentans		78.30	0.32	VPT 1				
Rothia mucilaginosa		21.70	0.00	URE 90				
507702	**Good identification**							**507702**
Staphylococcus piscifermentans		100.0	0.36	VPT 1				
507712	**Good identification**							**507712**
Staphylococcus piscifermentans		100.0	0.28	VPT 1				
507740	**Low discrimination**							**507740**
Staphylococcus piscifermentans		45.83	0.32	VPT 1				
Rothia mucilaginosa		29.64	0.08	URE 90				
Staphylococcus haemolyticus		24.53	0.18					
507742	**Low discrimination**							**507742**
Staphylococcus piscifermentans		83.29	0.36	VPT 1				
Staphylococcus haemolyticus		16.72	0.13	URE 90				
507753	**Low discrimination**							**507753**
					OXI	COA	NOV	
Staphylococcus caprae		64.77	0.30	URE 90	1	1	1	
Staphylococcus aureus subsp. aureus		27.76	0.18	ESL 1	1	99	4	
Staphylococcus lentus		6.75	0.23		99	1	99	
510011	**Good identification**							**510011**
Staphylococcus capitis subsp. capitis		100.0	0.35	ORN 1				
510211	**Good identification**							**510211**
Staphylococcus capitis subsp. capitis		100.0	0.40	ORN 1				
510640	**Low discrimination**							**510640**
Staphylococcus petrasii subsp. jettensis		74.10	0.51	ORN 1				
Staphylococcus petrasii subsp. pragensis		25.10	0.39					
Staphylococcus auricularis		0.47	0.21					
510642	**Good identification**							**510642**
Staphylococcus petrasii subsp. jettensis		96.64	0.36	ORN 1				
Staphylococcus petrasii subsp. pragensis		1.87	-.02					
Staphylococcus haemolyticus		1.11	0.12					
510643	**Low discrimination**							**510643**
					OXI			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus lugdunensis		87.35	0.29	ARG 5	1	
Kocuria kristinae		7.24	0.05	NIT 99	99	
Staphylococcus petrasii subsp. jettensis		5.41	-.05		1	
510743	**Good identification**					**510743**
					OXI	
Staphylococcus lugdunensis		97.31	0.29	ARG 5	1	
Kocuria kristinae		2.69	-.04	NIT 99	99	
512001	**Good identification**					**512001**
Staphylococcus capitis subsp. capitis		95.44	0.30	ORN 1		
Staphylococcus schleiferi subsp. schleiferi		4.56	0.04	MAN 93		
512011	**Very good identification**					**512011**
Staphylococcus capitis subsp. capitis		100.0	0.53	ORN 1		
512201	**Good identification**					**512201**
Staphylococcus capitis subsp. capitis		100.0	0.36	ORN 1		
512211	**Very good identification**					**512211**
Staphylococcus capitis subsp. capitis		100.0	0.59	ORN 1		
512240	**Low discrimination**					**512240**
Staphylococcus haemolyticus		45.85	0.35	ORN 1		
Staphylococcus petrasii subsp. jettensis		40.43	0.18	TRE 93		
Staphylococcus auricularis		8.45	0.19			
512242	**Low discrimination**					**512242**
					PYR	
Staphylococcus haemolyticus		51.99	0.30	ORN 1	98	
Staphylococcus epidermidis		33.63	0.24	TRE 93	1	
Staphylococcus petrasii subsp. jettensis		14.38	0.02		99	
512243	**Low discrimination**					**512243**
					PYR	
Staphylococcus lugdunensis		60.24	0.29	ARG 5	96	
Staphylococcus epidermidis		37.77	0.29	TRE 99	1	
Staphylococcus capitis subsp. ureolyticus		1.99	-.11		1	
512250	**Good identification**					**512250**
Staphylococcus haemolyticus		100.0	0.30	ORN 1		
512340	**Low discrimination**					**512340**
					PYR	
Staphylococcus haemolyticus		63.40	0.36	ORN 1	98	
Staphylococcus epidermidis		36.60	0.29	TRE 93	1	
512341	**Low discrimination**					**512341**
					PYR	
Staphylococcus epidermidis		88.66	0.35	URE 92	1	
Staphylococcus lugdunensis		8.78	0.09	ORN 12	96	
Staphylococcus haemolyticus		2.56	0.05		98	
512342	**Low discrimination**					**512342**
					PYR	
Staphylococcus epidermidis		76.47	0.39	URE 92	1	
Staphylococcus haemolyticus		23.53	0.31	ORN 12	98	
512343	**Low discrimination**					**512343**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus epidermidis		78.04	0.45	URE	92	
Staphylococcus lugdunensis		21.97	0.29	ORN	12	
512350	**Good identification**					**512350**
Staphylococcus haemolyticus		100.0	0.31	ORN	1	
512440	**Low discrimination**					**512440**
Staphylococcus auricularis		69.33	0.41	VPT	12	
Staphylococcus petrasii subsp. jettensis		26.62	0.18	ORN	1	
Staphylococcus haemolyticus		4.05	0.17			
512443	**Good identification**					**512443**
Staphylococcus lugdunensis		97.44	0.29	ARG	5	
Staphylococcus auricularis		2.56	0.02	SUC	99	
512543	**Good identification**					**512543**
Staphylococcus lugdunensis		100.0	0.29	ARG	5	
512600	**Low discrimination**					**512600**
Staphylococcus auricularis		47.20	0.33	VPT	12	
Staphylococcus petrasii subsp. jettensis		45.82	0.18	ORN	1	
Staphylococcus haemolyticus		6.97	0.17			
512603	**Good identification**					**512603**
Staphylococcus lugdunensis		100.0	0.29	ARG	5	
512640	**Low discrimination**					**512640**
Staphylococcus petrasii subsp. jettensis		83.72	0.59	ORN	1	
Staphylococcus haemolyticus		12.74	0.58			
Staphylococcus auricularis		2.03	0.40			
512641	**Low discrimination**					**512641**
Staphylococcus lugdunensis		86.15	0.50	ARG	5	
Staphylococcus petrasii subsp. jettensis		6.41	0.18	LAC	90	
Staphylococcus haemolyticus		2.99	0.27			
512642	**Low discrimination**					**512642**
Staphylococcus petrasii subsp. jettensis		63.68	0.43	ORN	1	
Staphylococcus haemolyticus		30.89	0.53			
Staphylococcus lugdunensis		4.46	0.29			
512643	**Very good identification**					**512643**
Staphylococcus lugdunensis		99.44	0.70	ARG	5	
Staphylococcus haemolyticus		0.22	0.22			
Staphylococcus petrasii subsp. petrasii		0.18	0.12			
512650	**Low discrimination**					**512650**
Staphylococcus haemolyticus		89.85	0.53	ORN	1	
Staphylococcus petrasii subsp. jettensis		10.15	0.18			
512652	**Good identification**					**512652**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		96.58	0.48	ORN 1			
Staphylococcus petrasii subsp. jettensis		3.42	0.02				
512653	**Low discrimination**						**512653**
Staphylococcus lugdunensis		88.81	0.29	ARG 5			
Staphylococcus haemolyticus		11.19	0.17	MAN 1			
512663	**Good identification**						**512663**
Staphylococcus lugdunensis		100.0	0.29	ARG 5			
512703	**Good identification**						**512703**
Staphylococcus lugdunensis		100.0	0.29	ARG 5			
512740	**Good identification**						**512740**
					OXI	NOV	
Staphylococcus haemolyticus		90.90	0.59	ORN 1	1	2	
Staphylococcus petrasii subsp. jettensis		5.35	0.18		1	1	
Macrococcus caseolyticus		1.44	0.11		99	99	
512741	**Acceptable identification**						**512741**
Staphylococcus lugdunensis		93.79	0.50	ARG 5			
Staphylococcus haemolyticus		3.67	0.28	LAC 90			
Staphylococcus petrasii subsp. petrasii		1.85	0.14				
512742	**Low discrimination**						**512742**
					OXI	NOV	
Staphylococcus haemolyticus		83.92	0.54	ORN 1	1	2	
Staphylococcus lugdunensis		10.73	0.29		1	1	
Macrococcus caseolyticus		3.54	0.14		99	99	
512743	**Very good identification**						**512743**
Staphylococcus lugdunensis		99.60	0.70	ARG 5			
Staphylococcus haemolyticus		0.24	0.23				
Staphylococcus petrasii subsp. petrasii		0.12	0.09				
512750	**Very good identification**						**512750**
Staphylococcus haemolyticus		100.0	0.54	ORN 1			
512752	**Good identification**						**512752**
Staphylococcus haemolyticus		100.0	0.49	ORN 1			
512753	**Low discrimination**						**512753**
Staphylococcus lugdunensis		85.68	0.29	ARG 5			
Staphylococcus haemolyticus		12.17	0.18	MAN 1			
Staphylococcus caprae		2.15	-.11				
512763	**Good identification**						**512763**
Staphylococcus lugdunensis		100.0	0.29	ARG 5			
513643	**Good identification**						**513643**
Staphylococcus lugdunensis		100.0	0.29	ARG 5			
513743	**Good identification**						**513743**
Staphylococcus lugdunensis		100.0	0.29	ARG 5			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
514343	**Good identification**					**514343**
Staphylococcus epidermidis		100.0	0.29	URE 92		
514401	**Good identification**					**514401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	ORN 1		
514501	**Good identification**					**514501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	ORN 1		
516001	**Acceptable identification**					**516001**
Staphylococcus schleiferi subsp. schleiferi		94.52	0.46	ORN 1		
Staphylococcus carnosus subsp. carnosus		3.01	0.15			
Staphylococcus capitis subsp. capitis		2.47	0.12			
516003	**Acceptable identification**					**516003**
Staphylococcus carnosus subsp. carnosus		92.66	0.27	ORN 1		
Staphylococcus schleiferi subsp. schleiferi		7.34	0.04	BGA 90		
516011	**Low discrimination**					**516011**
Staphylococcus capitis subsp. capitis		71.61	0.35	ORN 1		
Staphylococcus carnosus subsp. carnosus		26.31	0.27	PHS 11		
Staphylococcus schleiferi subsp. schleiferi		2.08	0.04			
516013	**Good identification**					**516013**
Staphylococcus carnosus subsp. carnosus		98.63	0.39	ORN 1		
Staphylococcus capitis subsp. capitis		1.37	0.00	BGA 90		
516101	**Good identification**					**516101**
Staphylococcus schleiferi subsp. schleiferi		99.21	0.46	ORN 1		
Staphylococcus carnosus subsp. carnosus		0.79	0.02			
516113	**Low discrimination**					**516113**
					COA	
Staphylococcus carnosus subsp. carnosus		89.72	0.27	ORN 1	1	
Staphylococcus schleiferi subsp. coagulans		10.28	0.12	BGA 90	99	
516211	**Good identification**					**516211**
Staphylococcus capitis subsp. capitis		100.0	0.40	ORN 1		
516240	**Good identification**					**516240**
Staphylococcus epidermidis		100.0	0.29	URE 92		
516241	**Good identification**					**516241**
Staphylococcus epidermidis		100.0	0.35	URE 92		
516242	**Good identification**					**516242**
Staphylococcus epidermidis		100.0	0.39	URE 92		
516243	**Good identification**					**516243**
Staphylococcus epidermidis		100.0	0.45	URE 92		
516340	**Good identification**					**516340**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		100.0	0.45	URE 92		
516341	**Good identification**					**516341**
Staphylococcus epidermidis		100.0	0.50	URE 92		
516342	**Very good identification**					**516342**
Staphylococcus epidermidis		100.0	0.55	URE 92		
516343	**Very good identification**					**516343**
Staphylococcus epidermidis		100.0	0.60	URE 92		
516401	**Very good identification**					**516401**
Staphylococcus schleiferi subsp. schleiferi		99.55	0.55	ORN 1		
Staphylococcus carnosus subsp. carnosus		0.45	0.07			
516413	**Good identification**					**516413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ORN 1		
516501	**Very good identification**					**516501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.55	ORN 1		
516553	**Good identification**					**516553**
Staphylococcus caprae		100.0	0.39	URE 90		
516643	**Good identification**					**516643**
					PYR	
Staphylococcus lugdunensis		96.54	0.29	ARG 5	96	
Staphylococcus epidermidis		3.46	0.04	PHS 1	1	
516743	**Low discrimination**					**516743**
					PYR	
Staphylococcus lugdunensis		74.95	0.29	ARG 5	96	
Staphylococcus epidermidis		15.24	0.19	PHS 1	1	
Staphylococcus caprae		9.81	0.04		75	
516753	**Low discrimination**					**516753**
					COA	
Staphylococcus caprae		70.00	0.30	URE 90	1	
Staphylococcus aureus subsp. aureus		30.00	0.18	ORN 1	99	
520001	**Low discrimination**					**520001**
Staphylococcus schleiferi subsp. schleiferi		86.45	0.30	NIT 90		
Staphylococcus capitis subsp. capitis		13.56	0.12	PHS 99		
520011	**Good identification**					**520011**
Staphylococcus capitis subsp. capitis		96.77	0.35	BGA 1		
Staphylococcus carnosus subsp. carnosus		3.23	0.05	NIT 89		
520101	**Good identification**					**520101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
520211	**Good identification**					**520211**
Staphylococcus capitis subsp. capitis		100.0	0.40	BGA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
520400	**Low discrimination**						**520400**
Staphylococcus auricularis		87.44	0.39	VPT 12			
Staphylococcus carnosus subsp. utilis		9.88	0.03	BGA 14			
Staphylococcus schleiferi subsp. schleiferi		1.47	-.02	NIT 90			
520401	**Acceptable identification**						**520401**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		92.94	0.39	NIT 90	1		
Staphylococcus auricularis		6.22	0.19	PHS 99	1		
Staphylococcus cohnii subsp. cohnii		0.84	0.05		99		
520440	**Low discrimination**						**520440**
Staphylococcus auricularis		86.14	0.46	VPT 12			
Staphylococcus petrasii subsp. jettensis		7.92	0.10	BGA 14			
Staphylococcus petrasii subsp. pragensis		2.68	-.02	NIT 90			
520441	**Low discrimination**						**520441**
					OXI	NOV	
Staphylococcus auricularis		66.25	0.27	VPT 12	1	1	
Kocuria kristinae		17.28	0.08	BGA 14	99	1	
Staphylococcus cohnii subsp. cohnii		12.19	0.15	NIT 90	1	99	
520442	**Low discrimination**						**520442**
					NOV	PYR	
Staphylococcus auricularis		83.35	0.27	VPT 12	1	40	
Staphylococcus petrasii subsp. jettensis		12.17	-.05	BGA 14	1	99	
Staphylococcus saprophyticus subsp. saprophyticus		4.48	-.12	NIT 90	99	7	
520501	**Good identification**						**520501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90			
520540	**Acceptable identification**						**520540**
Staphylococcus auricularis		92.35	0.27	VPT 12			
Staphylococcus haemolyticus		7.65	0.05	BGA 14			
520600	**Low discrimination**						**520600**
Staphylococcus auricularis		73.58	0.38	VPT 12			
Staphylococcus petrasii subsp. jettensis		17.10	0.10	BGA 14			
Staphylococcus petrasii subsp. pragensis		5.79	-.02	NIT 90			
520601	**Low discrimination**						**520601**
					OXI		
Kocuria kristinae		86.18	0.30	ARG 1	99		
Staphylococcus auricularis		11.88	0.18	MLT 90	1		
Staphylococcus schleiferi subsp. schleiferi		1.94	-.02		1		
520640	**Low discrimination**						**520640**
Staphylococcus petrasii subsp. jettensis		64.16	0.51	BGA 1			
Staphylococcus petrasii subsp. pragensis		21.73	0.39				
Staphylococcus auricularis		6.51	0.46				
520641	**Low discrimination**						**520641**
					OXI		
Kocuria kristinae		85.53	0.49	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		7.00	0.27		1		
Staphylococcus auricularis		3.06	0.26		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
520642	**Low discrimination**						**520642**
Staphylococcus petrasii subsp. jettensis		68.51	0.36	BGA 1			
Staphylococcus haemolyticus		19.39	0.40				
Staphylococcus auricularis		4.38	0.26				
520643	**Low discrimination**						**520643**
					OXI		
Kocuria kristinae		81.02	0.37	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		14.27	0.21		1		
Staphylococcus petrasii subsp. jettensis		1.83	-.05		1		
520650	**Low discrimination**						**520650**
					NOV	PYR	
Staphylococcus haemolyticus		46.91	0.41	NIT 99	2	98	
Staphylococcus saprophyticus subsp. saprophyticus		25.29	0.22		99	7	
Staphylococcus warneri		14.71	0.38		1	8	
520652	**Low discrimination**						**520652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		81.33	0.37	URE 99	99	7	
Staphylococcus haemolyticus		14.98	0.36	ARG 8	2	98	
Staphylococcus warneri		2.78	0.28		1	8	
520740	**Low discrimination**						**520740**
Staphylococcus haemolyticus		74.15	0.47	NIT 99			
Staphylococcus auricularis		8.35	0.26				
Staphylococcus petrasii subsp. jettensis		7.48	0.10				
520741	**Low discrimination**						**520741**
					OXI		
Kocuria kristinae		82.99	0.39	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		13.58	0.23		1		
Staphylococcus haemolyticus		2.45	0.16		1		
520742	**Low discrimination**						**520742**
					PYR		
Staphylococcus haemolyticus		89.74	0.41	NIT 99	98		
Staphylococcus warneri		3.79	0.20		8		
Staphylococcus petrasii subsp. jettensis		2.84	-.05		99		
520743	**Low discrimination**						**520743**
					OXI		
Kocuria kristinae		70.49	0.27	ARG 1	99		
Staphylococcus petrasii subsp. petrasii		24.84	0.18		1		
Staphylococcus haemolyticus		4.67	0.10		1		
520750	**Low discrimination**						**520750**
					NOV	PYR	
Staphylococcus haemolyticus		88.54	0.42	NIT 99	2	98	
Staphylococcus warneri		7.78	0.27		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		3.68	0.00		99	7	
520752	**Low discrimination**						**520752**
					NOV	PYR	
Staphylococcus haemolyticus		67.99	0.37	NIT 99	2	98	
Staphylococcus saprophyticus subsp. saprophyticus		28.47	0.15		99	7	
Staphylococcus warneri		3.54	0.17		1	8	
521641	**Good identification**						**521641**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Kocuria kristinae		100.0	0.37	ARG 1		
521741	**Low discrimination**					**521741**
					OXI	
Kocuria kristinae		87.76	0.27	ARG 1	99	
Rothia mucilaginosa		12.24	0.05		1	
522000	**Low discrimination**					**522000**
Staphylococcus auricularis		72.54	0.37	VPT 12		
Staphylococcus carnosus subsp. utilis		10.48	0.03	BGA 14		
Kocuria varians		7.00	0.17	TRE 92		
522001	**Low discrimination**					**522001**
Staphylococcus schleiferi subsp. schleiferi		74.09	0.49	PHS 99		
Staphylococcus carnosus subsp. carnosus		14.18	0.34			
Staphylococcus capitis subsp. capitis		10.44	0.30			
522003	**Good identification**					**522003**
Staphylococcus carnosus subsp. carnosus		98.33	0.47	PHS 99		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
Staphylococcus capitis subsp. capitis		0.37	-.05			
522011	**Low discrimination**					**522011**
Staphylococcus capitis subsp. capitis		70.72	0.53	BGA 1		
Staphylococcus carnosus subsp. carnosus		28.90	0.47			
Staphylococcus schleiferi subsp. schleiferi		0.38	0.08			
522013	**Very good identification**					**522013**
Staphylococcus carnosus subsp. carnosus		98.77	0.59	PHS 99		
Staphylococcus capitis subsp. capitis		1.23	0.19			
522040	**Acceptable identification**					**522040**
					PYR	
Staphylococcus auricularis		92.02	0.44	VPT 12	40	
Staphylococcus haemolyticus		7.15	0.22	BGA 14	98	
Kocuria varians		0.84	0.03	TRE 92	10	
522101	**Good identification**					**522101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.15	0.49	PHS 99	1	
Staphylococcus carnosus subsp. carnosus		4.55	0.22		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.02		99	
522103	**Low discrimination**					**522103**
					COA	
Staphylococcus carnosus subsp. carnosus		86.56	0.34	PHS 99	1	
Staphylococcus schleiferi subsp. coagulans		8.87	0.18		99	
Staphylococcus schleiferi subsp. schleiferi		4.57	0.08		1	
522111	**Low discrimination**					**522111**
Staphylococcus carnosus subsp. carnosus		85.69	0.34	PHS 99		
Staphylococcus capitis subsp. capitis		8.47	0.12			
Staphylococcus schleiferi subsp. schleiferi		4.52	0.08			
522113	**Good identification**					**522113**
					COA	
Staphylococcus carnosus subsp. carnosus		97.69	0.47	PHS 99	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99	
522200	**Low discrimination**					**522200**
Staphylococcus auricularis		83.59	0.36	VPT 12		
Staphylococcus haemolyticus		16.42	0.22	BGA 14		
522201	**Acceptable identification**					**522201**
Staphylococcus capitis subsp. capitis		90.90	0.36	BGA 1		
Staphylococcus auricularis		5.60	0.16	MAN 93		
Staphylococcus schleiferi subsp. schleiferi		3.51	0.08			
522211	**Very good identification**					**522211**
Staphylococcus capitis subsp. capitis		99.57	0.59	BGA 1		
Staphylococcus carnosus subsp. carnosus		0.22	0.05			
Staphylococcus capitis subsp. ureolyticus		0.21	-.05			
522240	**Low discrimination**					**522240**
Staphylococcus haemolyticus		85.90	0.63	TRE 93		
Staphylococcus auricularis		10.31	0.43			
Staphylococcus petrasii subsp. jettensis		3.06	0.18			
522241	**Low discrimination**					**522241**
					PYR	
Staphylococcus haemolyticus		45.39	0.32	TRE 93	98	
Staphylococcus epidermidis		23.20	0.25	MNS 3	1	
Staphylococcus auricularis		19.57	0.24		40	
522242	**Good identification**					**522242**
					PYR	
Staphylococcus haemolyticus		92.35	0.58	TRE 93	98	
Staphylococcus epidermidis		4.43	0.29		1	
Staphylococcus auricularis		2.19	0.24		40	
522243	**Low discrimination**					**522243**
					PYR	
Staphylococcus epidermidis		55.49	0.35	URE 92	1	
Staphylococcus capitis subsp. ureolyticus		21.47	0.13		1	
Staphylococcus haemolyticus		19.29	0.27		98	
522250	**Very good identification**					**522250**
Staphylococcus haemolyticus		99.79	0.59	TRE 93		
Staphylococcus auricularis		0.21	0.02			
522252	**Very good identification**					**522252**
					PYR	
Staphylococcus haemolyticus		98.60	0.54	TRE 93	98	
Staphylococcus capitis subsp. ureolyticus		1.40	0.00		1	
522253	**Good identification**					**522253**
					PYR	
Staphylococcus capitis subsp. ureolyticus		97.63	0.41	URE 99	1	
Staphylococcus haemolyticus		2.15	0.22	BGA 12	98	
Staphylococcus simulans		0.22	0.00		70	
522340	**Good identification**					**522340**
					PYR	
Staphylococcus haemolyticus		94.82	0.65	TRE 93	98	
Staphylococcus epidermidis		4.06	0.34		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus auricularis		1.12	0.24		40	
522341	**Low discrimination**					**522341**
					PYR	
Staphylococcus epidermidis		69.82	0.40	URE 92	1	
Staphylococcus haemolyticus		27.19	0.33		98	
Staphylococcus petrasii subsp. petrasii		1.84	0.02		99	
522342	**Low discrimination**					**522342**
					PYR	
Staphylococcus haemolyticus		80.44	0.59	TRE 93	98	
Staphylococcus epidermidis		19.37	0.45		1	
Staphylococcus auricularis		0.19	0.04		40	
522343	**Acceptable identification**					**522343**
					PYR	
Staphylococcus epidermidis		93.14	0.50	URE 92	1	
Staphylococcus haemolyticus		6.44	0.28		98	
Staphylococcus petrasii subsp. petrasii		0.42	-.04		99	
522350	**Very good identification**					**522350**
Staphylococcus haemolyticus		100.0	0.60	TRE 93		
522351	**Good identification**					**522351**
Staphylococcus haemolyticus		100.0	0.29	TRE 93		
522352	**Very good identification**					**522352**
					PYR	
Staphylococcus haemolyticus		99.59	0.55	TRE 93	98	
Staphylococcus epidermidis		0.41	0.04		1	
522400	**Good identification**					**522400**
Staphylococcus auricularis		92.18	0.58	VPT 12		
Staphylococcus carnosus subsp. utilis		6.08	0.18	BGA 14		
Staphylococcus schleiferi subsp. schleiferi		1.55	0.18			
522401	**Good identification**					**522401**
Staphylococcus schleiferi subsp. schleiferi		91.31	0.59	PHS 99		
Staphylococcus auricularis		6.11	0.39			
Staphylococcus carnosus subsp. carnosus		2.50	0.27			
522402	**Good identification**					**522402**
Staphylococcus auricularis		98.38	0.39	VPT 12		
Staphylococcus carnosus subsp. carnosus		1.62	-.02	BGA 14		
522403	**Low discrimination**					**522403**
Staphylococcus carnosus subsp. carnosus		86.18	0.39	PHS 99		
Staphylococcus schleiferi subsp. schleiferi		7.96	0.18			
Staphylococcus auricularis		5.86	0.19			
522411	**Low discrimination**					**522411**
Staphylococcus carnosus subsp. carnosus		82.72	0.39	PHS 99		
Staphylococcus capitis subsp. capitis		9.64	0.19			
Staphylococcus schleiferi subsp. schleiferi		7.64	0.18			
522413	**Good identification**					**522413**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		99.60	0.51	PHS 99		
Staphylococcus simulans		0.40	0.05			
522440	**Low discrimination**					**522440**
Staphylococcus auricularis		89.80	0.66	VPT 12		
Staphylococcus haemolyticus		8.06	0.46	BGA 14		
Staphylococcus petrasii subsp. jettensis		2.14	0.18			
522441	**Acceptable identification**					**522441**
Staphylococcus auricularis		90.01	0.46	VPT 12		
Staphylococcus schleiferi subsp. schleiferi		5.82	0.18	BGA 14		
Staphylococcus haemolyticus		2.25	0.14	MNS 10		
522442	**Low discrimination**					**522442**
Staphylococcus auricularis		67.01	0.46	VPT 12		
Staphylococcus haemolyticus		30.45	0.40	BGA 14		
Staphylococcus petrasii subsp. jettensis		2.54	0.02	LAC 10		
522443	**Low discrimination**					**522443**
Staphylococcus auricularis		77.31	0.27	VPT 12		
Staphylococcus haemolyticus		9.78	0.09	BGA 14		
Staphylococcus petrasii subsp. petrasii		7.99	0.00	MNS 10		
522450	**Low discrimination**					**522450**
Staphylococcus haemolyticus		83.92	0.41	SUC 99		
Staphylococcus auricularis		16.08	0.25			
522452	**Good identification**					**522452**
Staphylococcus haemolyticus		96.35	0.36	SUC 99		
Staphylococcus auricularis		3.65	0.05			
522500	**Low discrimination**					**522500**
Staphylococcus auricularis		85.29	0.39	VPT 12		
Staphylococcus schleiferi subsp. schleiferi		12.88	0.18	BGA 14		
Staphylococcus haemolyticus		1.83	0.05	GAL 10		
522501	**Very good identification**					**522501**
Staphylococcus schleiferi subsp. schleiferi		98.59	0.59	PHS 99		
Staphylococcus auricularis		0.73	0.19			
Staphylococcus carnosus subsp. carnosus		0.67	0.14			
522503	**Low discrimination**					**522503**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	PHS 99		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
522511	**Low discrimination**					**522511**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	PHS 99		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
522513	**Good identification**					**522513**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	PHS 99		
522540	**Low discrimination**					**522540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI	NOV	
Staphylococcus auricularis		50.06	0.46	VPT 12	1	1	
Staphylococcus haemolyticus		45.61	0.47	BGA 14	1	2	
Macrococcus caseolyticus		4.33	0.14	GAL 10	99	99	
522541	**Low discrimination**						**522541**
Staphylococcus auricularis		50.92	0.27	VPT 12			
Staphylococcus schleiferi subsp. schleiferi		29.65	0.18	BGA 14			
Staphylococcus haemolyticus		12.91	0.16	GAL 10			
522542	**Low discrimination**						**522542**
					OXI	NOV	
Staphylococcus haemolyticus		68.02	0.41	SUC 99	1	2	
Macrococcus caseolyticus		17.23	0.18		99	99	
Staphylococcus auricularis		14.75	0.27		1	1	
522550	**Good identification**						**522550**
Staphylococcus haemolyticus		98.15	0.42	SUC 99			
Staphylococcus auricularis		1.85	0.05				
522552	**Good identification**						**522552**
Staphylococcus haemolyticus		100.0	0.37	SUC 99			
522600	**Low discrimination**						**522600**
Staphylococcus auricularis		77.69	0.58	VPT 12			
Staphylococcus haemolyticus		17.63	0.46	BGA 14			
Staphylococcus petrasii subsp. jettensis		4.68	0.18				
522601	**Low discrimination**						**522601**
Staphylococcus auricularis		72.99	0.38	VPT 12			
Staphylococcus schleiferi subsp. schleiferi		11.94	0.18	BGA 14			
Staphylococcus haemolyticus		4.61	0.14	MNS 10			
522602	**Low discrimination**						**522602**
Staphylococcus haemolyticus		50.05	0.40	MLT 99			
Staphylococcus auricularis		43.57	0.38				
Staphylococcus petrasii subsp. jettensis		4.17	0.02				
522610	**Acceptable identification**						**522610**
Staphylococcus haemolyticus		92.95	0.41	MLT 99			
Staphylococcus auricularis		7.05	0.17				
522612	**Low discrimination**						**522612**
Staphylococcus haemolyticus		76.90	0.36	MLT 99			
Staphylococcus simulans		23.10	0.22				
522613	**Low discrimination**						**522613**
Staphylococcus simulans		88.02	0.33	VPT 4			
Staphylococcus carnosus subsp. carnosus		9.28	0.10	URE 98			
Staphylococcus haemolyticus		2.71	0.05				
522640	**Low discrimination**						**522640**
Staphylococcus haemolyticus		72.59	0.87				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus petrasii subsp. jettensis		19.27	0.59			
Staphylococcus auricularis		7.54	0.65			
522641	**Low discrimination**					**522641**
Staphylococcus haemolyticus		41.59	0.56	MNS 3		
Staphylococcus petrasii subsp. petrasii		35.52	0.47			
Staphylococcus auricularis		15.52	0.46			
522642	**Good identification**					**522642**
Staphylococcus haemolyticus		90.30	0.81			
Staphylococcus petrasii subsp. jettensis		7.52	0.43			
Staphylococcus auricularis		1.85	0.46			
522643	**Low discrimination**					**522643**
Staphylococcus haemolyticus		49.36	0.50	MNS 3		
Staphylococcus petrasii subsp. petrasii		40.35	0.41			
Staphylococcus auricularis		3.64	0.26			
522650	**Very good identification**					**522650**
Staphylococcus haemolyticus		99.18	0.82			
Staphylococcus petrasii subsp. jettensis		0.45	0.18			
Staphylococcus auricularis		0.18	0.24			
522651	**Low discrimination**					**522651**
Staphylococcus haemolyticus		85.17	0.51	MNS 3		
Staphylococcus petrasii subsp. petrasii		13.76	0.27			
Staphylococcus auricularis		0.55	0.05			
522652	**Very good identification**					**522652**
Staphylococcus haemolyticus		99.50	0.77			
Staphylococcus petrasii subsp. jettensis		0.14	0.02			
Staphylococcus simulans		0.14	0.15			
522653	**Low discrimination**					**522653**
Staphylococcus haemolyticus		74.80	0.46	MNS 3		
Staphylococcus petrasii subsp. petrasii		11.57	0.21			
Staphylococcus simulans		11.04	0.26			
522660	**Low discrimination**					**522660**
Staphylococcus haemolyticus		73.03	0.46	XYL 1		
Staphylococcus petrasii subsp. jettensis		19.39	0.18			
Staphylococcus auricularis		7.59	0.24			
522662	**Acceptable identification**					**522662**
Staphylococcus haemolyticus		90.60	0.40	XYL 1		
Staphylococcus petrasii subsp. jettensis		7.55	0.02			
Staphylococcus auricularis		1.86	0.05			
522670	**Good identification**					**522670**
Staphylococcus haemolyticus		100.0	0.41	XYL 1		
522672	**Good identification**					**522672**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		100.0	0.36	XYL 1			
522700	**Low discrimination**						**522700**
Staphylococcus haemolyticus		67.03	0.47	MLT 99			
Staphylococcus auricularis		29.11	0.38				
Rothia mucilaginosa		3.86	0.10				
522702	**Acceptable identification**						**522702**
Staphylococcus haemolyticus		92.10	0.41	MLT 99			
Staphylococcus auricularis		7.90	0.18				
522710	**Good identification**						**522710**
Staphylococcus haemolyticus		100.0	0.42	MLT 99			
522712	**Good identification**						**522712**
Staphylococcus haemolyticus		97.74	0.37	MLT 99			
Staphylococcus simulans		2.26	0.01				
522740	**Very good identification**						**522740**
Staphylococcus haemolyticus		98.36	0.88				
Staphylococcus auricularis		1.01	0.46				
Staphylococcus petrasii subsp. jettensis		0.23	0.18				
522741	**Low discrimination**						**522741**
Staphylococcus haemolyticus		58.77	0.57	MNS 3			
Staphylococcus petrasii subsp. petrasii		29.67	0.43				
Rothia mucilaginosa		7.73	0.27				
522742	**Very good identification**						**522742**
					OXI	NOV	
Staphylococcus haemolyticus		99.35	0.83		1	2	
Staphylococcus auricularis		0.20	0.26		1	1	
Macrococcus caseolyticus		0.17	0.14		99	99	
522743	**Low discrimination**						**522743**
Staphylococcus haemolyticus		64.16	0.51	MNS 3			
Staphylococcus petrasii subsp. petrasii		31.01	0.37				
Staphylococcus lugdunensis		2.68	0.17				
522750	**Very good identification**						**522750**
					PYR		
Staphylococcus haemolyticus		99.89	0.83		98		
Staphylococcus warneri		0.04	0.20		8		
Staphylococcus petrasii subsp. petrasii		0.03	0.04		99		
522751	**Acceptable identification**						**522751**
Staphylococcus haemolyticus		91.28	0.52	MNS 3			
Staphylococcus petrasii subsp. petrasii		8.72	0.23				
522752	**Very good identification**						**522752**
					PYR		
Staphylococcus haemolyticus		99.95	0.78		98		
Staphylococcus petrasii subsp. petrasii		0.03	-.02		99		
Staphylococcus warneri		0.02	0.10		8		
522753	**Low discrimination**						**522753**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
					COA	PYR	
Staphylococcus haemolyticus		86.11	0.47	MNS 3	1	98	
Staphylococcus petrasii subsp. petrasii		7.87	0.18		1	99	
Staphylococcus aureus subsp. aureus		2.90	-.02		99	10	
522760	**Good identification**						**522760**
Staphylococcus haemolyticus		98.99	0.47	XYL 1			
Staphylococcus auricularis		1.01	0.05				
522762	**Good identification**						**522762**
Staphylococcus haemolyticus		100.0	0.41	XYL 1			
522770	**Good identification**						**522770**
Staphylococcus haemolyticus		100.0	0.42	XYL 1			
522772	**Good identification**						**522772**
Staphylococcus haemolyticus		100.0	0.37	XYL 1			
523003	**Good identification**						**523003**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10			
523011	**Low discrimination**						**523011**
Staphylococcus carnosus subsp. carnosus		81.80	0.27	ESL 10			
Staphylococcus capitis subsp. capitis		18.20	0.12	PHS 99			
523013	**Good identification**						**523013**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ESL 10			
523113	**Low discrimination**						**523113**
					COA		
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ESL 10	1		
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	PHS 99	99		
523413	**Good identification**						**523413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ESL 10			
523640	**Low discrimination**						**523640**
Staphylococcus haemolyticus		72.30	0.46	ESL 1			
Staphylococcus petrasii subsp. jettensis		19.20	0.18				
Staphylococcus auricularis		7.51	0.24				
523642	**Acceptable identification**						**523642**
Staphylococcus haemolyticus		90.60	0.40	ESL 1			
Staphylococcus petrasii subsp. jettensis		7.55	0.02				
Staphylococcus auricularis		1.86	0.05				
523650	**Good identification**						**523650**
Staphylococcus haemolyticus		100.0	0.41	ESL 1			
523652	**Good identification**						**523652**
Staphylococcus haemolyticus		100.0	0.36	ESL 1			
523700	**Good identification**						**523700**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Rothia mucilaginosa		98.09	0.29	ARG 1			
Staphylococcus haemolyticus		1.91	0.05				
523701	**Good identification**						**523701**
Rothia mucilaginosa		100.0	0.39	ARG 1			
523713	**Good identification**						**523713**
Staphylococcus lentus		100.0	0.30	VPT 7			
523733	**Good identification**						**523733**
Staphylococcus lentus		100.0	0.30	VPT 7			
523740	**Low discrimination**						**523740**
Rothia mucilaginosa		54.46	0.37	ARG 1			
Staphylococcus haemolyticus		45.08	0.47				
Staphylococcus auricularis		0.46	0.05				
523741	**Good identification**						**523741**
					OXI		
Rothia mucilaginosa		98.20	0.47	ARG 1	1		
Staphylococcus haemolyticus		0.84	0.16		1		
Kocuria kristinae		0.54	0.04		99		
523742	**Good identification**						**523742**
Staphylococcus haemolyticus		97.88	0.41	ESL 1			
Rothia mucilaginosa		2.12	-.04				
523750	**Good identification**						**523750**
Staphylococcus haemolyticus		97.96	0.42	ESL 1			
Rothia mucilaginosa		2.04	-.04				
523752	**Good identification**						**523752**
Staphylococcus haemolyticus		100.0	0.37	ESL 1			
523753	**Good identification**						**523753**
					OXI	NOV	
Staphylococcus lentus		96.69	0.36	VPT 7	99	99	
Staphylococcus haemolyticus		3.31	0.06	ARG 1	1	2	
523773	**Good identification**						**523773**
Staphylococcus lentus		100.0	0.36	VPT 7			
524000	**Good identification**						**524000**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90			
524001	**Very good identification**						**524001**
Staphylococcus schleiferi subsp. schleiferi		98.29	0.71	NIT 90			
Staphylococcus carnosus subsp. carnosus		1.71	0.34				
524003	**Low discrimination**						**524003**
Staphylococcus carnosus subsp. carnosus		87.32	0.47	NIT 99			
Staphylococcus schleiferi subsp. schleiferi		12.68	0.30				
524011	**Low discrimination**						**524011**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		84.56	0.47	NIT 99		
Staphylococcus schleiferi subsp. schleiferi		12.28	0.30			
Staphylococcus capitis subsp. capitis		3.16	0.16			
524013	**Very good identification**					**524013**
Staphylococcus carnosus subsp. carnosus		100.0	0.59	NIT 99		
524021	**Good identification**					**524021**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
524041	**Good identification**					**524041**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
524100	**Good identification**					**524100**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
524101	**Very good identification**					**524101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		99.54	0.71	NIT 90	1	
Staphylococcus carnosus subsp. carnosus		0.43	0.22		1	
Staphylococcus schleiferi subsp. coagulans		0.03	0.02		99	
524103	**Low discrimination**					**524103**
					COA	
Staphylococcus carnosus subsp. carnosus		59.41	0.34	NIT 99	1	
Staphylococcus schleiferi subsp. schleiferi		34.50	0.30		1	
Staphylococcus schleiferi subsp. coagulans		6.09	0.18		99	
524111	**Low discrimination**					**524111**
					COA	
Staphylococcus carnosus subsp. carnosus		62.65	0.34	NIT 99	1	
Staphylococcus schleiferi subsp. schleiferi		36.38	0.30		1	
Staphylococcus schleiferi subsp. coagulans		0.97	0.01		99	
524113	**Good identification**					**524113**
					COA	
Staphylococcus carnosus subsp. carnosus		97.69	0.47	NIT 99	1	
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99	
524121	**Good identification**					**524121**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
524141	**Good identification**					**524141**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
524201	**Good identification**					**524201**
Staphylococcus schleiferi subsp. schleiferi		96.53	0.30	NIT 90		
Staphylococcus capitis subsp. capitis		3.47	-.02	SUC 1		
524301	**Good identification**					**524301**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	NIT 90		
524342	**Good identification**					**524342**
Staphylococcus epidermidis		100.0	0.29	URE 92		
524343	**Good identification**					**524343**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		100.0	0.35	URE 92		
524400	**Good identification**					**524400**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90		
524401	**Very good identification**					**524401**
Staphylococcus schleiferi subsp. schleiferi		99.75	0.80	NIT 90		
Staphylococcus carnosus subsp. carnosus		0.25	0.27			
524403	**Low discrimination**					**524403**
Staphylococcus schleiferi subsp. schleiferi		50.40	0.39	NIT 90		
Staphylococcus carnosus subsp. carnosus		49.60	0.39	LAC 1		
524411	**Low discrimination**					**524411**
Staphylococcus schleiferi subsp. schleiferi		50.40	0.39	NIT 90		
Staphylococcus carnosus subsp. carnosus		49.60	0.39	MAN 1		
524413	**Good identification**					**524413**
Staphylococcus carnosus subsp. carnosus		99.74	0.51	NIT 99		
Staphylococcus schleiferi subsp. schleiferi		0.26	-.02			
524421	**Good identification**					**524421**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90		
524441	**Good identification**					**524441**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90		
524450	**Good identification**					**524450**
Staphylococcus kloosii		100.0	0.32	VPT 10		
524452	**Good identification**					**524452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
524500	**Good identification**					**524500**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90		
524501	**Very good identification**					**524501**
Staphylococcus schleiferi subsp. schleiferi		99.94	0.80	NIT 90		
Staphylococcus carnosus subsp. carnosus		0.06	0.14			
524503	**Low discrimination**					**524503**
Staphylococcus schleiferi subsp. schleiferi		80.26	0.39	NIT 90		
Staphylococcus carnosus subsp. carnosus		19.74	0.27	LAC 1		
524511	**Low discrimination**					**524511**
Staphylococcus schleiferi subsp. schleiferi		80.26	0.39	NIT 90		
Staphylococcus carnosus subsp. carnosus		19.74	0.27	MAN 1		
524513	**Good identification**					**524513**
Staphylococcus carnosus subsp. carnosus		98.98	0.39	NIT 99		
Staphylococcus schleiferi subsp. schleiferi		1.02	-.02			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
524521	**Good identification**						**524521**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90			
524541	**Good identification**						**524541**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90			
524550	**Good identification**						**524550**
Staphylococcus kloosii		100.0	0.38	VPT 10			
524552	**Low discrimination**						**524552**
					NOV	YEP	
Staphylococcus kloosii		73.24	0.39	VPT 10	99	99	
Staphylococcus caprae		26.76	0.02	ARG 1	1	1	
524572	**Good identification**						**524572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
524601	**Good identification**						**524601**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90			
524701	**Good identification**						**524701**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	NIT 90			
525001	**Low discrimination**						**525001**
Staphylococcus schleiferi subsp. schleiferi		83.94	0.30	ESL 1			
Staphylococcus carnosus subsp. carnosus		16.06	0.15	NIT 90			
525003	**Good identification**						**525003**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10			
525011	**Good identification**						**525011**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10			
525013	**Good identification**						**525013**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ESL 10			
525101	**Acceptable identification**						**525101**
					COA		
Staphylococcus schleiferi subsp. schleiferi		92.99	0.30	ESL 1	1		
Staphylococcus carnosus subsp. carnosus		4.45	0.02	NIT 90	1		
Staphylococcus schleiferi subsp. coagulans		2.57	0.01		99		
525113	**Low discrimination**						**525113**
					COA		
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ESL 10	1		
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	NIT 99	99		
525401	**Good identification**						**525401**
Staphylococcus schleiferi subsp. schleiferi		97.34	0.39	ESL 1			
Staphylococcus carnosus subsp. carnosus		2.66	0.07	NIT 90			
525413	**Good identification**						**525413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ESL 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
525450	**Good identification**					**525450**
Staphylococcus kloosii		100.0	0.32	VPT 10		
525452	**Good identification**					**525452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
525501	**Good identification**					**525501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ESL 1		
525550	**Good identification**					**525550**
Staphylococcus kloosii		100.0	0.38	VPT 10		
525552	**Good identification**					**525552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
525572	**Good identification**					**525572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
526000	**Low discrimination**					**526000**
Staphylococcus schleiferi subsp. schleiferi		83.94	0.49	MNS 99		
Staphylococcus carnosus subsp. carnosus		16.06	0.34			
526001	**Low discrimination**					**526001**
Staphylococcus schleiferi subsp. schleiferi		83.92	0.90			
Staphylococcus carnosus subsp. carnosus		16.06	0.75			
Staphylococcus capitis subsp. capitis		0.02	0.12			
526002	**Good identification**					**526002**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	MNS 99		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
526003	**Very good identification**					**526003**
					COA	
Staphylococcus carnosus subsp. carnosus		98.67	0.88		1	
Staphylococcus schleiferi subsp. schleiferi		1.30	0.49		1	
Staphylococcus schleiferi subsp. coagulans		0.03	0.18		99	
526010	**Good identification**					**526010**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	MNS 99		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
526011	**Very good identification**					**526011**
Staphylococcus carnosus subsp. carnosus		98.40	0.88			
Staphylococcus schleiferi subsp. schleiferi		1.30	0.49			
Staphylococcus capitis subsp. capitis		0.30	0.35			
526012	**Very good identification**					**526012**
Staphylococcus carnosus subsp. carnosus		100.0	0.59	MNS 99		
526013	**Excellent identification**					**526013**
					COA	
Staphylococcus carnosus subsp. carnosus		99.99	1.00		1	
Staphylococcus schleiferi subsp. coagulans		0.01	0.17		99	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus schleiferi subsp. schleiferi		0.00	0.08		1	
526021	**Low discrimination**					**526021**
Staphylococcus schleiferi subsp. schleiferi		83.94	0.49	XYL	1	
Staphylococcus carnosus subsp. carnosus		16.06	0.34			
526023	**Good identification**					**526023**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	XYL	1	
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
526031	**Good identification**					**526031**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	XYL	1	
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
526033	**Very good identification**					**526033**
Staphylococcus carnosus subsp. carnosus		100.0	0.59	XYL	1	
526041	**Low discrimination**					**526041**
Staphylococcus schleiferi subsp. schleiferi		83.94	0.49	MLT	1	
Staphylococcus carnosus subsp. carnosus		16.06	0.34			
526043	**Good identification**					**526043**
Staphylococcus carnosus subsp. carnosus		97.68	0.47	MLT	1	
Staphylococcus schleiferi subsp. schleiferi		1.29	0.08			
Staphylococcus epidermidis		1.03	0.09			
526051	**Good identification**					**526051**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	MLT	1	
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
526053	**Very good identification**					**526053**
Staphylococcus carnosus subsp. carnosus		100.0	0.59	MLT	1	
526100	**Good identification**					**526100**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.15	0.49	MNS 99	1	
Staphylococcus carnosus subsp. carnosus		4.55	0.22		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.02		99	
526101	**Very good identification**					**526101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.15	0.90		1	
Staphylococcus carnosus subsp. carnosus		4.55	0.63		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.43		99	
526102	**Low discrimination**					**526102**
					COA	
Staphylococcus carnosus subsp. carnosus		86.56	0.34	MNS 99	1	
Staphylococcus schleiferi subsp. coagulans		8.87	0.18		99	
Staphylococcus schleiferi subsp. schleiferi		4.57	0.08		1	
526103	**Low discrimination**					**526103**
					COA	
Staphylococcus carnosus subsp. carnosus		86.56	0.75		1	
Staphylococcus schleiferi subsp. coagulans		8.87	0.59		99	
Staphylococcus schleiferi subsp. schleiferi		4.57	0.49		1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
526110	**Acceptable identification**					**526110**
					COA	
Staphylococcus carnosus subsp. carnosus		93.62	0.34	MNS 99	1	
Staphylococcus schleiferi subsp. schleiferi		4.94	0.08		1	
Staphylococcus schleiferi subsp. coagulans		1.44	0.01		99	
526111	**Good identification**					**526111**
					COA	
Staphylococcus carnosus subsp. carnosus		93.62	0.75		1	
Staphylococcus schleiferi subsp. schleiferi		4.94	0.49		1	
Staphylococcus schleiferi subsp. coagulans		1.44	0.42		99	
526112	**Good identification**					**526112**
					COA	
Staphylococcus carnosus subsp. carnosus		97.69	0.47	MNS 99	1	
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99	
526113	**Very good identification**					**526113**
					COA	
Staphylococcus carnosus subsp. carnosus		97.68	0.88		1	
Staphylococcus schleiferi subsp. coagulans		2.31	0.58		99	
Staphylococcus schleiferi subsp. schleiferi		0.01	0.08		1	
526121	**Good identification**					**526121**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.15	0.49	XYL 1	1	
Staphylococcus carnosus subsp. carnosus		4.55	0.22		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.02		99	
526123	**Low discrimination**					**526123**
					COA	
Staphylococcus carnosus subsp. carnosus		86.56	0.34	XYL 1	1	
Staphylococcus schleiferi subsp. coagulans		8.87	0.18		99	
Staphylococcus schleiferi subsp. schleiferi		4.57	0.08		1	
526131	**Acceptable identification**					**526131**
					COA	
Staphylococcus carnosus subsp. carnosus		93.62	0.34	XYL 1	1	
Staphylococcus schleiferi subsp. schleiferi		4.94	0.08		1	
Staphylococcus schleiferi subsp. coagulans		1.44	0.01		99	
526133	**Good identification**					**526133**
					COA	
Staphylococcus carnosus subsp. carnosus		97.69	0.47	XYL 1	1	
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99	
526141	**Acceptable identification**					**526141**
Staphylococcus schleiferi subsp. schleiferi		93.85	0.49	MLT 1		
Staphylococcus carnosus subsp. carnosus		4.49	0.22			
Staphylococcus epidermidis		1.36	0.14			
526143	**Low discrimination**					**526143**
					COA	
Staphylococcus carnosus subsp. carnosus		70.44	0.34	MLT 1	1	
Staphylococcus epidermidis		16.85	0.25		1	
Staphylococcus schleiferi subsp. coagulans		7.22	0.18		99	
526151	**Acceptable identification**					**526151**
Staphylococcus carnosus subsp. carnosus		91.46	0.34	MLT 1		
Staphylococcus schleiferi subsp. schleiferi		4.83	0.08			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus caprae		2.31	-.07			
526153	**Low discrimination**					**526153**
					COA	
Staphylococcus carnosus subsp. carnosus		87.45	0.47	MLT 1	1	
Staphylococcus caprae		10.48	0.20		1	
Staphylococcus schleiferi subsp. coagulans		2.07	0.17		99	
526201	**Low discrimination**					**526201**
Staphylococcus schleiferi subsp. schleiferi		81.72	0.49	SUC 1		
Staphylococcus carnosus subsp. carnosus		15.64	0.34			
Staphylococcus capitis subsp. capitis		2.64	0.17			
526203	**Good identification**					**526203**
Staphylococcus carnosus subsp. carnosus		96.91	0.47	SUC 1		
Staphylococcus schleiferi subsp. schleiferi		1.28	0.08			
Staphylococcus epidermidis		1.02	0.09			
526211	**Low discrimination**					**526211**
Staphylococcus carnosus subsp. carnosus		63.50	0.47	SUC 1		
Staphylococcus capitis subsp. capitis		35.66	0.40			
Staphylococcus schleiferi subsp. schleiferi		0.84	0.08			
526213	**Very good identification**					**526213**
					COA	
Staphylococcus carnosus subsp. carnosus		99.53	0.59	SUC 1	1	
Staphylococcus capitis subsp. capitis		0.29	0.05		1	
Staphylococcus schleiferi subsp. coagulans		0.19	0.07		99	
526240	**Low discrimination**					**526240**
					PYR	
Staphylococcus epidermidis		81.00	0.34	URE 92	1	
Staphylococcus haemolyticus		16.97	0.22		98	
Staphylococcus auricularis		2.04	0.02		40	
526241	**Good identification**					**526241**
Staphylococcus epidermidis		96.15	0.40	URE 92		
Staphylococcus schleiferi subsp. schleiferi		3.85	0.08			
526242	**Good identification**					**526242**
					PYR	
Staphylococcus epidermidis		96.41	0.45	URE 92	1	
Staphylococcus haemolyticus		3.59	0.17		98	
526243	**Good identification**					**526243**
Staphylococcus epidermidis		99.04	0.50	URE 92		
Staphylococcus carnosus subsp. carnosus		0.96	0.05			
526301	**Low discrimination**					**526301**
					COA	
Staphylococcus schleiferi subsp. schleiferi		86.05	0.49	SUC 1	1	
Staphylococcus schleiferi subsp. coagulans		8.59	0.32		99	
Staphylococcus carnosus subsp. carnosus		4.12	0.22		1	
526303	**Low discrimination**					**526303**
					COA	
Staphylococcus schleiferi subsp. coagulans		70.18	0.49	URE 99	99	
Staphylococcus carnosus subsp. carnosus		21.91	0.34		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		5.24	0.25		1		
526311	**Low discrimination**						**526311**
					COA		
Staphylococcus carnosus subsp. carnosus		64.23	0.34	SUC 1	1		
Staphylococcus schleiferi subsp. coagulans		30.92	0.32		99		
Staphylococcus schleiferi subsp. schleiferi		3.39	0.08		1		
526313	**Low discrimination**						**526313**
					COA		
Staphylococcus carnosus subsp. carnosus		56.97	0.47	SUC 1	1		
Staphylococcus schleiferi subsp. coagulans		42.11	0.48		99		
Staphylococcus intermedius		0.92	0.11		99		
526340	**Good identification**						**526340**
					PYR		
Staphylococcus epidermidis		96.00	0.50	URE 92	1		
Staphylococcus haemolyticus		4.00	0.23		98		
526341	**Very good identification**						**526341**
Staphylococcus epidermidis		99.30	0.56	URE 92			
Staphylococcus schleiferi subsp. schleiferi		0.70	0.08				
526342	**Very good identification**						**526342**
					PYR		
Staphylococcus epidermidis		99.27	0.60	URE 92	1		
Staphylococcus haemolyticus		0.74	0.18		98		
526343	**Very good identification**						**526343**
					COA	PYR	
Staphylococcus epidermidis		99.71	0.66	URE 92	1	1	
Staphylococcus intermedius		0.15	0.06		99	90	
Staphylococcus schleiferi subsp. coagulans		0.14	0.08		99	20	
526400	**Very good identification**						**526400**
Staphylococcus schleiferi subsp. schleiferi		96.77	0.59	MNS 99			
Staphylococcus carnosus subsp. carnosus		2.65	0.27				
Staphylococcus auricularis		0.59	0.17				
526401	**Very good identification**						**526401**
Staphylococcus schleiferi subsp. schleiferi		97.34	1.00				
Staphylococcus carnosus subsp. carnosus		2.66	0.68				
526402	**Acceptable identification**						**526402**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	MNS 99			
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18				
526403	**Good identification**						**526403**
Staphylococcus carnosus subsp. carnosus		91.54	0.80				
Staphylococcus schleiferi subsp. schleiferi		8.46	0.59				
526410	**Acceptable identification**						**526410**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	MNS 99			
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18				
526411	**Good identification**						**526411**
Staphylococcus carnosus subsp. carnosus		91.53	0.80				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus schleiferi subsp. schleiferi		8.46	0.59			
Staphylococcus capitis subsp. capitis		0.01	0.00			
526412	**Good identification**					**526412**
Staphylococcus carnosus subsp. carnosus		100.0	0.51	MNS 99		
526413	**Very good identification**					**526413**
Staphylococcus carnosus subsp. carnosus		99.97	0.92			
Staphylococcus schleiferi subsp. schleiferi		0.02	0.18			
Staphylococcus felis		0.01	0.01			
526421	**Very good identification**					**526421**
Staphylococcus schleiferi subsp. schleiferi		97.34	0.59	XYL 1		
Staphylococcus carnosus subsp. carnosus		2.66	0.27			
526423	**Acceptable identification**					**526423**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	XYL 1		
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18			
526431	**Acceptable identification**					**526431**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	XYL 1		
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18			
526433	**Good identification**					**526433**
Staphylococcus carnosus subsp. carnosus		100.0	0.51	XYL 1		
526441	**Very good identification**					**526441**
Staphylococcus schleiferi subsp. schleiferi		97.19	0.59	MLT 1		
Staphylococcus carnosus subsp. carnosus		2.66	0.27			
Staphylococcus auricularis		0.15	0.05			
526443	**Acceptable identification**					**526443**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	MLT 1		
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18			
526451	**Acceptable identification**					**526451**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	MLT 1		
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18			
526453	**Good identification**					**526453**
Staphylococcus carnosus subsp. carnosus		99.37	0.51	MLT 1		
Staphylococcus caprae		0.63	-.02			
526500	**Very good identification**					**526500**
Staphylococcus schleiferi subsp. schleiferi		99.32	0.59	MNS 99		
Staphylococcus carnosus subsp. carnosus		0.68	0.14			
526501	**Excellent identification**					**526501**
					COA	
Staphylococcus schleiferi subsp. schleiferi		99.32	1.00		1	
Staphylococcus carnosus subsp. carnosus		0.68	0.55		1	
Staphylococcus schleiferi subsp. coagulans		0.00	0.02		99	
526502	**Low discrimination**					**526502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		73.02	0.27	MNS 99		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
526503	**Low discrimination**					**526503**
					COA	
Staphylococcus carnosus subsp. carnosus		72.86	0.68		1	
Staphylococcus schleiferi subsp. schleiferi		26.92	0.59		1	
Staphylococcus schleiferi subsp. coagulans		0.18	0.18		99	
526510	**Low discrimination**					**526510**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	MNS 99		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
526511	**Low discrimination**					**526511**
					COA	
Staphylococcus carnosus subsp. carnosus		73.00	0.68		1	
Staphylococcus schleiferi subsp. schleiferi		26.98	0.59		1	
Staphylococcus schleiferi subsp. coagulans		0.03	0.01		99	
526512	**Good identification**					**526512**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	MNS 99		
526513	**Very good identification**					**526513**
Staphylococcus carnosus subsp. carnosus		99.53	0.80			
Staphylococcus condimenti		0.21	0.14			
Staphylococcus schleiferi subsp. schleiferi		0.09	0.18			
526521	**Very good identification**					**526521**
Staphylococcus schleiferi subsp. schleiferi		99.32	0.59	XYL 1		
Staphylococcus carnosus subsp. carnosus		0.68	0.14			
526523	**Low discrimination**					**526523**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	XYL 1		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
526531	**Low discrimination**					**526531**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	XYL 1		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
526533	**Good identification**					**526533**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	XYL 1		
526541	**Very good identification**					**526541**
Staphylococcus schleiferi subsp. schleiferi		99.32	0.59	MLT 1		
Staphylococcus carnosus subsp. carnosus		0.68	0.14			
526543	**Low discrimination**					**526543**
Staphylococcus carnosus subsp. carnosus		52.65	0.27	MLT 1		
Staphylococcus caprae		27.89	0.13			
Staphylococcus schleiferi subsp. schleiferi		19.46	0.18			
526551	**Low discrimination**					**526551**
Staphylococcus carnosus subsp. carnosus		52.65	0.27	MLT 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus caprae		27.89	0.13			
Staphylococcus schleiferi subsp. schleiferi		19.46	0.18			
526553	**Low discrimination**					**526553**
					COA	
Staphylococcus caprae		70.67	0.39	URE 90	1	
Staphylococcus carnosus subsp. carnosus		28.09	0.39	BGA 1	1	
Staphylococcus aureus subsp. aureus		1.25	-.02		99	
526601	**Very good identification**					**526601**
Staphylococcus schleiferi subsp. schleiferi		97.34	0.59	SUC 1		
Staphylococcus carnosus subsp. carnosus		2.66	0.27			
526603	**Low discrimination**					**526603**
Staphylococcus carnosus subsp. carnosus		86.29	0.39	SUC 1		
Staphylococcus schleiferi subsp. schleiferi		7.97	0.18			
Staphylococcus simulans		2.76	0.11			
526611	**Low discrimination**					**526611**
Staphylococcus carnosus subsp. carnosus		89.36	0.39	SUC 1		
Staphylococcus schleiferi subsp. schleiferi		8.26	0.18			
Staphylococcus capitis subsp. capitis		2.39	0.05			
526613	**Good identification**					**526613**
Staphylococcus carnosus subsp. carnosus		95.62	0.51	SUC 1		
Staphylococcus simulans		3.05	0.23			
Staphylococcus felis		1.33	0.07			
526640	**Low discrimination**					**526640**
Staphylococcus haemolyticus		73.03	0.46	PHS 1		
Staphylococcus petrasii subsp. jettensis		19.39	0.18			
Staphylococcus auricularis		7.59	0.24			
526642	**Low discrimination**					**526642**
Staphylococcus haemolyticus		88.95	0.40	PHS 1		
Staphylococcus petrasii subsp. jettensis		7.41	0.02			
Staphylococcus auricularis		1.83	0.05			
526650	**Good identification**					**526650**
Staphylococcus haemolyticus		100.0	0.41	PHS 1		
526652	**Good identification**					**526652**
Staphylococcus haemolyticus		95.69	0.36	PHS 1		
Staphylococcus simulans		4.31	0.05			
526701	**Very good identification**					**526701**
					COA	
Staphylococcus schleiferi subsp. schleiferi		98.72	0.59	SUC 1	1	
Staphylococcus carnosus subsp. carnosus		0.68	0.14		1	
Staphylococcus intermedius		0.33	0.09		99	
526711	**Low discrimination**					**526711**
					COA	
Staphylococcus carnosus subsp. carnosus		58.93	0.27	SUC 1	1	
Staphylococcus schleiferi subsp. schleiferi		21.78	0.18		1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus intermedius		19.29	0.18		99		
526713	**Low discrimination**						**526713**
					COA		
Staphylococcus carnosus subsp. carnosus		47.87	0.39	SUC 1	1		
Staphylococcus intermedius		24.07	0.34		99		
Staphylococcus condimenti		14.64	0.18		1		
526740	**Good identification**						**526740**
					PYR		
Staphylococcus haemolyticus		97.23	0.47	PHS 1	98		
Staphylococcus epidermidis		1.77	0.09		1		
Staphylococcus auricularis		1.00	0.05		40		
526742	**Acceptable identification**						**526742**
					PYR		
Staphylococcus haemolyticus		90.69	0.41	PHS 1	98		
Staphylococcus epidermidis		9.31	0.19		1		
526743	**Low discrimination**						**526743**
					COA	PYR	
Staphylococcus intermedius		47.45	0.29	VPT 1	99	90	
Staphylococcus epidermidis		23.74	0.25	URE 90	1	1	
Staphylococcus caprae		8.33	0.04		1	75	
526750	**Good identification**						**526750**
Staphylococcus haemolyticus		100.0	0.42	PHS 1			
526752	**Low discrimination**						**526752**
					COA	PYR	
Staphylococcus haemolyticus		71.51	0.37	PHS 1	1	98	
Staphylococcus caprae		21.12	0.13		1	75	
Staphylococcus aureus subsp. aureus		7.38	-.02		99	10	
526753	**Low discrimination**						**526753**
					COA	PYR	
Staphylococcus aureus subsp. aureus		71.84	0.39	URE 99	99	10	
Staphylococcus caprae		15.24	0.30	BGA 10	1	75	
Staphylococcus intermedius		12.35	0.38		99	90	
527001	**Low discrimination**						**527001**
					COA		
Staphylococcus carnosus subsp. carnosus		67.73	0.56	ESL 10	1		
Staphylococcus schleiferi subsp. schleiferi		32.18	0.49		1		
Staphylococcus schleiferi subsp. coagulans		0.10	0.01		99		
527002	**Good identification**						**527002**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10			
527003	**Very good identification**						**527003**
					COA		
Staphylococcus carnosus subsp. carnosus		99.66	0.68	ESL 10	1		
Staphylococcus schleiferi subsp. coagulans		0.22	0.18		99		
Staphylococcus schleiferi subsp. schleiferi		0.12	0.08		1		
527010	**Good identification**						**527010**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10			
527011	**Very good identification**						**527011**
					COA		
Staphylococcus carnosus subsp. carnosus		99.85	0.68	ESL 10	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus schleiferi subsp. schleiferi		0.12	0.08		1	
Staphylococcus schleiferi subsp. coagulans		0.03	0.01		99	
527012	**Good identification**					**527012**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ESL 10		
527013	**Very good identification**					**527013**
					COA	
Staphylococcus carnosus subsp. carnosus		99.95	0.80	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		0.05	0.17		99	
527023	**Good identification**					**527023**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10		
527031	**Good identification**					**527031**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10		
527033	**Good identification**					**527033**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ESL 10		
527043	**Good identification**					**527043**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10		
527051	**Good identification**					**527051**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ESL 10		
527053	**Good identification**					**527053**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ESL 10		
527101	**Low discrimination**					**527101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		54.65	0.49	ESL 1	1	
Staphylococcus carnosus subsp. carnosus		28.76	0.43		1	
Staphylococcus schleiferi subsp. coagulans		16.59	0.42		99	
527103	**Low discrimination**					**527103**
					COA	
Staphylococcus carnosus subsp. carnosus		52.89	0.56	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		46.86	0.59		99	
Staphylococcus schleiferi subsp. schleiferi		0.25	0.08		1	
527111	**Low discrimination**					**527111**
					COA	
Staphylococcus carnosus subsp. carnosus		87.88	0.56	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		11.70	0.42		99	
Staphylococcus schleiferi subsp. schleiferi		0.42	0.08		1	
527112	**Low discrimination**					**527112**
					COA	
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	MNS 99	99	
527113	**Low discrimination**					**527113**
					COA	
Staphylococcus carnosus subsp. carnosus		83.03	0.68	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		16.98	0.58		99	
527133	**Low discrimination**					**527133**
					COA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	XYL 1	99	
527153	**Low discrimination**					**527153**
					COA	
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	MLT 1	99	
527203	**Acceptable identification**					**527203**
					COA	
Staphylococcus carnosus subsp. carnosus		93.46	0.27	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		6.54	0.07	SUC 1	99	
527211	**Good identification**					**527211**
Staphylococcus carnosus subsp. carnosus		95.14	0.27	ESL 10		
Staphylococcus capitis subsp. capitis		4.86	-.01	SUC 1		
527213	**Good identification**					**527213**
					COA	
Staphylococcus carnosus subsp. carnosus		98.41	0.39	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		1.59	0.07	SUC 1	99	
527301	**Low discrimination**					**527301**
					COA	
Staphylococcus schleiferi subsp. coagulans		86.15	0.32	URE 99	99	
Staphylococcus schleiferi subsp. schleiferi		9.08	0.08	LAC 86	1	
Staphylococcus carnosus subsp. carnosus		4.78	0.02		1	
527303	**Good identification**					**527303**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.52	0.48	URE 99	99	
Staphylococcus carnosus subsp. carnosus		3.49	0.15		1	
527311	**Low discrimination**					**527311**
					COA	
Staphylococcus schleiferi subsp. coagulans		80.63	0.31	URE 99	99	
Staphylococcus carnosus subsp. carnosus		19.37	0.15	LAC 86	1	
527313	**Low discrimination**					**527313**
					COA	
Staphylococcus schleiferi subsp. coagulans		86.47	0.48	URE 99	99	
Staphylococcus carnosus subsp. carnosus		13.53	0.27		1	
527401	**Low discrimination**					**527401**
Staphylococcus schleiferi subsp. schleiferi		76.88	0.59	ESL 1		
Staphylococcus carnosus subsp. carnosus		23.12	0.48			
527403	**Very good identification**					**527403**
Staphylococcus carnosus subsp. carnosus		99.17	0.60	ESL 10		
Staphylococcus schleiferi subsp. schleiferi		0.83	0.18			
527411	**Very good identification**					**527411**
Staphylococcus carnosus subsp. carnosus		99.17	0.60	ESL 10		
Staphylococcus schleiferi subsp. schleiferi		0.83	0.18			
527412	**Good identification**					**527412**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ESL 10		
527413	**Very good identification**					**527413**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus carnosus subsp. carnosus		100.0	0.73	ESL 10		
527433	**Good identification**					**527433**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ESL 10		
527453	**Good identification**					**527453**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ESL 10		
527501	**Good identification**					**527501**
					COA	
Staphylococcus schleiferi subsp. schleiferi		92.92	0.59	ESL 1	1	
Staphylococcus carnosus subsp. carnosus		6.99	0.36		1	
Staphylococcus schleiferi subsp. coagulans		0.10	0.01		99	
527503	**Acceptable identification**					**527503**
					COA	
Staphylococcus carnosus subsp. carnosus		94.83	0.48	ESL 10	1	
Staphylococcus schleiferi subsp. schleiferi		3.19	0.18		1	
Staphylococcus schleiferi subsp. coagulans		1.98	0.18		99	
527511	**Good identification**					**527511**
					COA	
Staphylococcus carnosus subsp. carnosus		96.46	0.48	ESL 10	1	
Staphylococcus schleiferi subsp. schleiferi		3.24	0.18		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.01		99	
527513	**Very good identification**					**527513**
					COA	
Staphylococcus carnosus subsp. carnosus		99.52	0.60	ESL 10	1	
Staphylococcus schleiferi subsp. coagulans		0.48	0.17		99	
527613	**Good identification**					**527613**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ESL 10		
532342	**Low discrimination**					**532342**
					PYR	
Staphylococcus epidermidis		76.47	0.27	URE 92	1	
Staphylococcus haemolyticus		23.53	0.18	ORN 12	98	
532343	**Low discrimination**					**532343**
					PYR	
Staphylococcus epidermidis		78.04	0.33	URE 92	1	
Staphylococcus lugdunensis		21.97	0.17	ORN 12	96	
532640	**Low discrimination**					**532640**
Staphylococcus haemolyticus		72.55	0.46	ORN 1		
Staphylococcus petrasii subsp. jettensis		19.26	0.18			
Staphylococcus auricularis		7.54	0.24			
532641	**Acceptable identification**					**532641**
Staphylococcus lugdunensis		92.83	0.38	ARG 5		
Staphylococcus haemolyticus		3.22	0.14	LAC 90		
Staphylococcus petrasii subsp. petrasii		2.75	0.05			
532642	**Low discrimination**					**532642**
Staphylococcus haemolyticus		80.13	0.40	ORN 1		
Staphylococcus lugdunensis		11.56	0.17			
Staphylococcus petrasii subsp. jettensis		6.67	0.02			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
532643	**Very good identification**					**532643**
Staphylococcus lugdunensis		99.61	0.58	ARG 5		
Staphylococcus haemolyticus		0.22	0.09			
Staphylococcus petrasii subsp. petrasii		0.18	0.00			
532650	**Good identification**					**532650**
Staphylococcus haemolyticus		100.0	0.41	ORN 1		
532652	**Good identification**					**532652**
Staphylococcus haemolyticus		100.0	0.36	ORN 1		
532740	**Good identification**					**532740**
Staphylococcus haemolyticus		98.21	0.47	ORN 1		
Staphylococcus auricularis		1.01	0.05			
Staphylococcus lugdunensis		0.79	-.03			
532741	**Acceptable identification**					**532741**
Staphylococcus lugdunensis		94.45	0.38	ARG 5		
Staphylococcus haemolyticus		3.69	0.16	LAC 90		
Staphylococcus petrasii subsp. petrasii		1.86	0.02			
532742	**Low discrimination**					**532742**
Staphylococcus haemolyticus		88.66	0.41	ORN 1		
Staphylococcus lugdunensis		11.34	0.17			
532743	**Very good identification**					**532743**
Staphylococcus lugdunensis		99.64	0.58	ARG 5		
Staphylococcus haemolyticus		0.24	0.10			
Staphylococcus petrasii subsp. petrasii		0.12	-.04			
532750	**Good identification**					**532750**
Staphylococcus haemolyticus		100.0	0.42	ORN 1		
532752	**Good identification**					**532752**
Staphylococcus haemolyticus		100.0	0.37	ORN 1		
534001	**Good identification**					**534001**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	ORN 1		
534101	**Good identification**					**534101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	ORN 1		
534401	**Good identification**					**534401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ORN 1		
534501	**Good identification**					**534501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	ORN 1		
536001	**Low discrimination**					**536001**
Staphylococcus schleiferi subsp. schleiferi		83.94	0.49	ORN 1		
Staphylococcus carnosus subsp. carnosus		16.06	0.34			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
536003	**Good identification**					**536003**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	ORN 1		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
536011	**Good identification**					**536011**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	ORN 1		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
536013	**Very good identification**					**536013**
Staphylococcus carnosus subsp. carnosus		100.0	0.59	ORN 1		
536101	**Good identification**					**536101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.15	0.49	ORN 1	1	
Staphylococcus carnosus subsp. carnosus		4.55	0.22		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.02		99	
536103	**Low discrimination**					**536103**
					COA	
Staphylococcus carnosus subsp. carnosus		86.56	0.34	ORN 1	1	
Staphylococcus schleiferi subsp. coagulans		8.87	0.18		99	
Staphylococcus schleiferi subsp. schleiferi		4.57	0.08		1	
536111	**Acceptable identification**					**536111**
					COA	
Staphylococcus carnosus subsp. carnosus		93.62	0.34	ORN 1	1	
Staphylococcus schleiferi subsp. schleiferi		4.94	0.08		1	
Staphylococcus schleiferi subsp. coagulans		1.44	0.01		99	
536113	**Good identification**					**536113**
					COA	
Staphylococcus carnosus subsp. carnosus		97.69	0.47	ORN 1	1	
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99	
536242	**Good identification**					**536242**
Staphylococcus epidermidis		100.0	0.27	URE 92		
536243	**Good identification**					**536243**
Staphylococcus epidermidis		100.0	0.33	URE 92		
536340	**Good identification**					**536340**
Staphylococcus epidermidis		100.0	0.32	URE 92		
536341	**Good identification**					**536341**
Staphylococcus epidermidis		100.0	0.38	URE 92		
536342	**Good identification**					**536342**
Staphylococcus epidermidis		100.0	0.42	URE 92		
536343	**Good identification**					**536343**
Staphylococcus epidermidis		100.0	0.48	URE 92		
536401	**Very good identification**					**536401**
Staphylococcus schleiferi subsp. schleiferi		97.34	0.59	ORN 1		
Staphylococcus carnosus subsp. carnosus		2.66	0.27			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
536403	**Acceptable identification**						**536403**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	ORN 1			
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18				
536411	**Acceptable identification**						**536411**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	ORN 1			
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18				
536413	**Good identification**						**536413**
Staphylococcus carnosus subsp. carnosus		100.0	0.51	ORN 1			
536501	**Very good identification**						**536501**
Staphylococcus schleiferi subsp. schleiferi		99.32	0.59	ORN 1			
Staphylococcus carnosus subsp. carnosus		0.68	0.14				
536503	**Low discrimination**						**536503**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	ORN 1			
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18				
536511	**Low discrimination**						**536511**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	ORN 1			
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18				
536513	**Good identification**						**536513**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ORN 1			
537003	**Good identification**						**537003**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ORN 1			
537011	**Good identification**						**537011**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	ORN 1			
537013	**Good identification**						**537013**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	ORN 1			
537113	**Low discrimination**						**537113**
					COA		
Staphylococcus carnosus subsp. carnosus		83.03	0.27	ORN 1	1		
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	ESL 10	99		
537413	**Good identification**						**537413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	ORN 1			
540011	**Good identification**						**540011**
Staphylococcus capitis subsp. capitis		100.0	0.35	GLR 1			
540211	**Good identification**						**540211**
Staphylococcus capitis subsp. capitis		100.0	0.40	GLR 1			
540240	**Low discrimination**						**540240**
					PYR	YEP	
Staphylococcus warneri		38.34	0.39	URE 98	8	43	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. jettensis		20.37	0.10	TRE 95	99	1	
Staphylococcus petrasii subsp. croceilyticus		15.81	0.05		90	99	
540242	**Low discrimination**						**540242**
					PYR		
Staphylococcus warneri		51.28	0.29	URE 98	8		
Staphylococcus haemolyticus		34.30	0.18	TRE 95	98		
Staphylococcus petrasii subsp. jettensis		14.42	-.05		99		
540250	**Low discrimination**						**540250**
					PYR		
Staphylococcus warneri		62.50	0.36	URE 98	8		
Staphylococcus haemolyticus		20.10	0.19	TRE 95	98		
Staphylococcus pasteuri		17.40	0.10		1		
540400	**Low discrimination**						**540400**
					NOV		
Staphylococcus cohnii subsp. cohnii		44.71	0.30	ARG 1	99		
Staphylococcus auricularis		32.39	0.24		1		
Staphylococcus carnosus subsp. utilis		19.27	0.03		1		
540401	**Acceptable identification**						**540401**
					NOV		
Staphylococcus cohnii subsp. cohnii		93.08	0.31	ARG 1	99		
Staphylococcus auricularis		6.92	0.04		1		
540410	**Good identification**						**540410**
					NOV		
Staphylococcus cohnii subsp. cohnii		95.59	0.32	ARG 1	99		
Staphylococcus warneri		4.41	0.10		1		
540411	**Acceptable identification**						**540411**
					NOV		
Staphylococcus cohnii subsp. cohnii		93.89	0.33	ARG 1	99		
Staphylococcus capitis subsp. capitis		6.11	0.00		1		
540440	**Low discrimination**						**540440**
					NOV		
Staphylococcus cohnii subsp. cohnii		38.58	0.41	ARG 1	99		
Staphylococcus auricularis		20.59	0.32		1		
Staphylococcus warneri		18.76	0.39		1		
540441	**Acceptable identification**						**540441**
					NOV		
Staphylococcus cohnii subsp. cohnii		94.81	0.41	ARG 1	99		
Staphylococcus auricularis		5.19	0.12		1		
540442	**Low discrimination**						**540442**
					PYR		
Staphylococcus warneri		60.71	0.29	URE 98	8		
Staphylococcus auricularis		22.22	0.12	SUC 95	40		
Staphylococcus petrasii subsp. jettensis		17.07	-.05		99		
540450	**Low discrimination**						**540450**
					NOV	YEP	
Staphylococcus cohnii subsp. cohnii		73.87	0.43	ARG 1	99	12	
Staphylococcus warneri		20.44	0.36		1	43	
Staphylococcus pasteuri		5.69	0.10		1	86	
540451	**Good identification**						**540451**
Staphylococcus cohnii subsp. cohnii		100.0	0.43	ARG 1			
540540	**Low discrimination**						**540540**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus warneri		43.48	0.29	URE 98	1		
Staphylococcus cohnii subsp. cohnii		31.46	0.21	SUC 95	99		
Staphylococcus auricularis		16.80	0.12		1		
540600	**Low discrimination**						**540600**
					PYR	YEP	
Staphylococcus warneri		31.85	0.39	URE 98	8	43	
Staphylococcus petrasii subsp. jettensis		16.92	0.10	MLT 95	99	1	
Staphylococcus pasteuri		16.85	0.19		1	86	
540602	**Low discrimination**						**540602**
					PYR	YEP	
Staphylococcus warneri		48.21	0.29	URE 98	8	43	
Staphylococcus pasteuri		31.25	0.11	MLT 95	1	86	
Staphylococcus petrasii subsp. jettensis		13.56	-.05		99	1	
540610	**Low discrimination**						**540610**
Staphylococcus pasteuri		81.78	0.35	URE 99			
Staphylococcus warneri		18.22	0.36	MLT 86			
540612	**Low discrimination**						**540612**
Staphylococcus pasteuri		84.61	0.27	URE 99			
Staphylococcus warneri		15.39	0.26	MLT 86			
540640	**Low discrimination**						**540640**
					PYR	YEP	
Staphylococcus petrasii subsp. jettensis		37.63	0.51	GLR 1	99	1	
Staphylococcus petrasii subsp. croceilyticus		29.22	0.47		90	99	
Staphylococcus warneri		13.60	0.65		8	43	
540642	**Low discrimination**						**540642**
					PYR		
Staphylococcus petrasii subsp. jettensis		45.03	0.36	GLR 1	99		
Staphylococcus warneri		30.73	0.55		8		
Staphylococcus haemolyticus		14.38	0.41		98		
540650	**Low discrimination**						**540650**
					PYR		
Staphylococcus pasteuri		52.62	0.51	URE 99	1		
Staphylococcus warneri		36.27	0.62		8		
Staphylococcus haemolyticus		8.15	0.42		98		
540651	**Low discrimination**						**540651**
					PYR		
Staphylococcus warneri		48.57	0.28	URE 98	8		
Staphylococcus pasteuri		34.88	0.10	MNS 2	1		
Staphylococcus haemolyticus		16.55	0.11		98		
540652	**Low discrimination**						**540652**
					PYR		
Staphylococcus pasteuri		55.95	0.43	URE 99	1		
Staphylococcus warneri		31.47	0.53		8		
Staphylococcus haemolyticus		11.94	0.37		98		
540700	**Low discrimination**						**540700**
					PYR		
Staphylococcus warneri		74.47	0.29	URE 98	8		
Staphylococcus haemolyticus		14.15	0.07	MLT 95	98		
Staphylococcus auricularis		11.38	0.04		40		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
540740	**Low discrimination**					**540740**
					PYR	
Staphylococcus warneri		45.54	0.55	URE 98	8	
Staphylococcus haemolyticus		45.08	0.48		98	
Staphylococcus petrasii subsp. jettensis		4.03	0.10		99	
540742	**Low discrimination**					**540742**
					PYR	
Staphylococcus haemolyticus		61.47	0.43	NIT 99	98	
Staphylococcus warneri		36.80	0.45		8	
Staphylococcus petrasii subsp. jettensis		1.73	-.05		99	
540750	**Low discrimination**					**540750**
					PYR	
Staphylococcus warneri		54.08	0.52	URE 98	8	
Staphylococcus haemolyticus		43.41	0.43		98	
Staphylococcus pasteuri		2.51	0.10		1	
540752	**Low discrimination**					**540752**
					PYR	
Staphylococcus haemolyticus		56.18	0.38	NIT 99	98	
Staphylococcus warneri		41.47	0.42		8	
Staphylococcus pasteuri		2.36	0.02		1	
542000	**Acceptable identification**					**542000**
Kocuria rosea		91.46	0.40	ARG 1		
Staphylococcus auricularis		4.85	0.22			
Staphylococcus carnosus subsp. utilis		3.69	0.03			
542001	**Low discrimination**					**542001**
Staphylococcus capitis subsp. capitis		78.05	0.30	GLR 1		
Kocuria rosea		16.39	0.14	MAN 93		
Staphylococcus schleiferi subsp. schleiferi		3.73	0.04			
542011	**Very good identification**					**542011**
Staphylococcus capitis subsp. capitis		99.83	0.53	GLR 1		
Kocuria rosea		0.18	-.06			
542020	**Good identification**					**542020**
Kocuria rosea		100.0	0.33	ARG 1		
542040	**Low discrimination**					**542040**
					PYR	
Staphylococcus auricularis		64.81	0.29	VPT 12	40	
Staphylococcus haemolyticus		29.90	0.23	GLR 3	98	
Kocuria rosea		5.29	-.01	TRE 92	1	
542201	**Good identification**					**542201**
Staphylococcus capitis subsp. capitis		97.63	0.36	GLR 1		
Kocuria rosea		1.23	-.06	MAN 93		
Staphylococcus auricularis		1.14	0.01			
542211	**Very good identification**					**542211**
Staphylococcus capitis subsp. capitis		100.0	0.59	GLR 1		
542240	**Good identification**					**542240**
					YEP	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus haemolyticus		90.35	0.65	TRE 93	20	
Staphylococcus petrasii subsp. croceilyticus		3.80	0.18		99	
Staphylococcus petrasii subsp. jettensis		2.85	0.18		1	
542241	**Acceptable identification**					**542241**
Staphylococcus haemolyticus		93.23	0.33	TRE 93		
Staphylococcus auricularis		6.77	0.09	MNS 3		
542242	**Very good identification**					**542242**
					PYR	
Staphylococcus haemolyticus		97.84	0.59	TRE 93	98	
Staphylococcus petrasii subsp. jettensis		0.97	0.02		99	
Staphylococcus warneri		0.63	0.22		8	
542243	**Low discrimination**					**542243**
					PYR	
Staphylococcus haemolyticus		85.03	0.28	TRE 93	98	
Staphylococcus epidermidis		8.76	0.06	MNS 3	1	
Staphylococcus capitis subsp. ureolyticus		6.21	-.11		1	
542250	**Very good identification**					**542250**
					PYR	
Staphylococcus haemolyticus		96.63	0.60	TRE 93	98	
Staphylococcus pasteuri		2.07	0.18		1	
Staphylococcus warneri		1.30	0.28		8	
542251	**Low discrimination**					**542251**
					PYR	
Staphylococcus haemolyticus		54.43	0.29	TRE 93	98	
Staphylococcus capitis subsp. capitis		33.10	0.18	MNS 3	1	
Staphylococcus capitis subsp. ureolyticus		12.47	0.00		1	
542252	**Very good identification**					**542252**
					PYR	
Staphylococcus haemolyticus		97.70	0.55	TRE 93	98	
Staphylococcus pasteuri		1.52	0.10		1	
Staphylococcus warneri		0.78	0.19		8	
542340	**Very good identification**					**542340**
					PYR	
Staphylococcus haemolyticus		99.35	0.66	TRE 93	98	
Staphylococcus warneri		0.30	0.21		8	
Staphylococcus auricularis		0.20	0.09		40	
542341	**Acceptable identification**					**542341**
					PYR	
Staphylococcus haemolyticus		91.58	0.35	TRE 93	98	
Staphylococcus epidermidis		8.42	0.11	MNS 3	1	
542342	**Very good identification**					**542342**
					PYR	
Staphylococcus haemolyticus		98.97	0.60	TRE 93	98	
Staphylococcus epidermidis		0.85	0.16		1	
Staphylococcus warneri		0.18	0.11		8	
542343	**Low discrimination**					**542343**
					PYR	
Staphylococcus haemolyticus		65.89	0.29	TRE 93	98	
Staphylococcus epidermidis		34.11	0.22	MNS 3	1	
542350	**Very good identification**					**542350**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		99.62	0.61	TRE 93	98		
Staphylococcus warneri		0.38	0.18		8		
542351	**Good identification**						**542351**
Staphylococcus haemolyticus		100.0	0.30	TRE 93			
542352	**Very good identification**						**542352**
Staphylococcus haemolyticus		100.0	0.56	TRE 93			
542400	**Low discrimination**						**542400**
Staphylococcus auricularis		72.06	0.44	VPT 12			
Staphylococcus carnosus subsp. utilis		25.01	0.18	GLR 3			
Kocuria rosea		1.19	-.01				
542440	**Low discrimination**						**542440**
					YEP		
Staphylococcus auricularis		52.84	0.51	VPT 12	1		
Staphylococcus haemolyticus		28.16	0.47	GLR 3	20		
Staphylococcus petrasii subsp. croceilyticus		8.82	0.18		99		
542441	**Low discrimination**						**542441**
					NOV	PYR	
Staphylococcus auricularis		76.86	0.32	VPT 12	1	40	
Staphylococcus cohnii subsp. cohnii		11.74	0.18	GLR 3	99	1	
Staphylococcus haemolyticus		11.40	0.16	MNS 10	2	98	
542442	**Low discrimination**						**542442**
Staphylococcus haemolyticus		67.00	0.41	SUC 99			
Staphylococcus auricularis		24.83	0.32				
Staphylococcus petrasii subsp. jettensis		4.95	0.02				
542450	**Low discrimination**						**542450**
					PYR		
Staphylococcus haemolyticus		73.75	0.42	SUC 99	98		
Staphylococcus pasteuri		11.77	0.18		1		
Staphylococcus warneri		7.40	0.28		8		
542452	**Low discrimination**						**542452**
					PYR		
Staphylococcus haemolyticus		85.08	0.37	SUC 99	98		
Staphylococcus pasteuri		9.86	0.10		1		
Staphylococcus warneri		5.06	0.19		8		
542540	**Low discrimination**						**542540**
					OXI	NOV	
Staphylococcus haemolyticus		77.41	0.48	SUC 99	1	2	
Staphylococcus auricularis		14.31	0.32		1	1	
Macrococcus caseolyticus		6.52	0.14		99	99	
542542	**Low discrimination**						**542542**
					OXI	NOV	
Staphylococcus haemolyticus		78.46	0.43	SUC 99	1	2	
Macrococcus caseolyticus		17.61	0.18		99	99	
Staphylococcus auricularis		2.87	0.12		1	1	
542550	**Good identification**						**542550**
					PYR		
Staphylococcus haemolyticus		97.27	0.43	SUC 99	98		
Staphylococcus warneri		2.73	0.18		8		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
542552	**Good identification**						**542552**
Staphylococcus haemolyticus		100.0	0.38	SUC 99			
542600	**Low discrimination**						**542600**
					YEP		
Staphylococcus haemolyticus		38.20	0.47	MLT 99	20		
Staphylococcus auricularis		28.35	0.43		1		
Staphylococcus petrasii subsp. croceilyticus		11.96	0.18		99		
542602	**Low discrimination**						**542602**
					PYR	YEP	
Staphylococcus haemolyticus		69.10	0.41	MLT 99	98	20	
Staphylococcus pasteuri		12.34	0.19		1	86	
Staphylococcus auricularis		10.13	0.23		40	1	
542610	**Low discrimination**						**542610**
					PYR		
Staphylococcus pasteuri		69.80	0.43	URE 99	1		
Staphylococcus haemolyticus		27.14	0.42	MLT 86	98		
Staphylococcus warneri		2.72	0.28		8		
542612	**Low discrimination**						**542612**
					PYR		
Staphylococcus pasteuri		62.85	0.35	URE 99	1		
Staphylococcus haemolyticus		33.65	0.37	MLT 86	98		
Staphylococcus warneri		2.00	0.19		8		
542640	**Low discrimination**						**542640**
					YEP		
Staphylococcus haemolyticus		62.47	0.88		20		
Staphylococcus petrasii subsp. croceilyticus		19.57	0.59		99		
Staphylococcus petrasii subsp. jettensis		14.70	0.59		1		
542641	**Low discrimination**						**542641**
					YEP		
Staphylococcus haemolyticus		77.51	0.57	MNS 3	20		
Staphylococcus petrasii subsp. croceilyticus		7.93	0.18		99		
Staphylococcus petrasii subsp. jettensis		5.96	0.18		1		
542642	**Good identification**						**542642**
					PYR	YEP	
Staphylococcus haemolyticus		90.64	0.83		98	20	
Staphylococcus petrasii subsp. jettensis		6.69	0.43		99	1	
Staphylococcus pasteuri		1.00	0.35		1	86	
542643	**Acceptable identification**						**542643**
Staphylococcus haemolyticus		93.29	0.51	MNS 3			
Staphylococcus petrasii subsp. petrasii		2.73	0.12				
Staphylococcus petrasii subsp. jettensis		2.25	0.02				
542650	**Low discrimination**						**542650**
					PYR		
Staphylococcus haemolyticus		84.13	0.83		98		
Staphylococcus pasteuri		13.43	0.59		1		
Staphylococcus warneri		1.62	0.55		8		
542651	**Acceptable identification**						**542651**
					PYR		
Staphylococcus haemolyticus		93.40	0.52	MNS 3	98		
Staphylococcus pasteuri		4.87	0.18		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		1.19	0.20		8		
542652	**Low discrimination**						**542652**
					PYR		
Staphylococcus haemolyticus		88.59	0.78		98		
Staphylococcus pasteuri		10.27	0.51		1		
Staphylococcus warneri		1.01	0.45		8		
542653	**Acceptable identification**						**542653**
					PYR	YEP	
Staphylococcus haemolyticus		93.71	0.47	MNS 3	98	20	
Staphylococcus pasteuri		3.55	0.10		1	86	
Staphylococcus simulans		2.04	0.10		70	1	
542660	**Low discrimination**						**542660**
					YEP		
Staphylococcus haemolyticus		62.62	0.47	XYL 1	20		
Staphylococcus petrasii subsp. croceilyticus		19.62	0.18		99		
Staphylococcus petrasii subsp. jettensis		14.74	0.18		1		
542662	**Acceptable identification**						**542662**
Staphylococcus haemolyticus		93.13	0.41	XYL 1			
Staphylococcus petrasii subsp. jettensis		6.88	0.02				
542670	**Low discrimination**						**542670**
					PYR		
Staphylococcus haemolyticus		84.83	0.42	XYL 1	98		
Staphylococcus pasteuri		13.54	0.18		1		
Staphylococcus warneri		1.63	0.14		8		
542672	**Low discrimination**						**542672**
					PYR		
Staphylococcus haemolyticus		89.62	0.37	XYL 1	98		
Staphylococcus pasteuri		10.39	0.10		1		
542700	**Acceptable identification**						**542700**
					PYR		
Staphylococcus haemolyticus		91.25	0.48	MLT 99	98		
Staphylococcus auricularis		6.67	0.23		40		
Staphylococcus warneri		2.08	0.21		8		
542702	**Good identification**						**542702**
					PYR		
Staphylococcus haemolyticus		96.05	0.43	MLT 99	98		
Staphylococcus auricularis		1.39	0.04		40		
Staphylococcus warneri		1.30	0.11		8		
542703	**Good identification**						**542703**
					PYR		
Staphylococcus hyicus		97.68	0.35	VPT 1	1		
Staphylococcus haemolyticus		2.32	0.11	PHS 92	98		
542710	**Good identification**						**542710**
					PYR		
Staphylococcus haemolyticus		95.14	0.43	MLT 99	98		
Staphylococcus warneri		2.67	0.18		8		
Staphylococcus pasteuri		2.19	0.02		1		
542712	**Good identification**						**542712**
Staphylococcus haemolyticus		100.0	0.38	MLT 99			
542740	**Very good identification**						**542740**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					PYR			
Staphylococcus haemolyticus		98.84	0.89		98			
Staphylococcus warneri		0.43	0.47		8			
Staphylococcus petrasii subsp. croceilyticus		0.28	0.18		90			
542741	**Very good identification**							**542741**
Staphylococcus haemolyticus		96.91	0.58	MNS 3				
Staphylococcus petrasii subsp. petrasii		1.75	0.14					
Staphylococcus auricularis		0.60	0.11					
542742	**Very good identification**							**542742**
					OXI	PYR	NOV	
Staphylococcus haemolyticus		99.49	0.84		1	98	2	
Staphylococcus warneri		0.26	0.38		1	8	1	
Macrococcus caseolyticus		0.15	0.14		99	80	99	
542743	**Good identification**							**542743**
					PYR			
Staphylococcus haemolyticus		97.89	0.53	MNS 3	98			
Staphylococcus petrasii subsp. petrasii		1.69	0.09		99			
Staphylococcus hyicus		0.42	-.06		1			
542750	**Very good identification**							**542750**
					PYR			
Staphylococcus haemolyticus		99.32	0.84		98			
Staphylococcus warneri		0.54	0.44		8			
Staphylococcus pasteuri		0.14	0.18		1			
542751	**Good identification**							**542751**
					PYR			
Staphylococcus haemolyticus		99.31	0.53	MNS 3	98			
Staphylococcus warneri		0.35	0.10		8			
Staphylococcus petrasii subsp. petrasii		0.34	-.05		99			
542752	**Very good identification**							**542752**
					PYR			
Staphylococcus haemolyticus		99.58	0.79		98			
Staphylococcus warneri		0.32	0.35		8			
Staphylococcus pasteuri		0.10	0.10		1			
542753	**Good identification**							**542753**
Staphylococcus haemolyticus		99.37	0.48	MNS 3				
Staphylococcus caprae		0.63	-.11					
542760	**Good identification**							**542760**
Staphylococcus haemolyticus		100.0	0.48	XYL 1				
542762	**Good identification**							**542762**
Staphylococcus haemolyticus		100.0	0.43	XYL 1				
542770	**Good identification**							**542770**
Staphylococcus haemolyticus		100.0	0.43	XYL 1				
542772	**Good identification**							**542772**
Staphylococcus haemolyticus		100.0	0.38	XYL 1				
543640	**Low discrimination**							**543640**
					YEP			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		62.62	0.47	ESL 1	20		
Staphylococcus petrasii subsp. croceilyticus		19.62	0.18		99		
Staphylococcus petrasii subsp. jettensis		14.74	0.18		1		
543642	**Acceptable identification**						**543642**
Staphylococcus haemolyticus		93.13	0.41	ESL 1			
Staphylococcus petrasii subsp. jettensis		6.88	0.02				
543650	**Low discrimination**						**543650**
					PYR		
Staphylococcus haemolyticus		84.83	0.42	ESL 1	98		
Staphylococcus pasteuri		13.54	0.18		1		
Staphylococcus warneri		1.63	0.14		8		
543652	**Low discrimination**						**543652**
					PYR		
Staphylococcus haemolyticus		89.62	0.37	ESL 1	98		
Staphylococcus pasteuri		10.39	0.10		1		
543740	**Good identification**						**543740**
Staphylococcus haemolyticus		95.85	0.48	ESL 1			
Rothia mucilaginosa		4.15	0.08				
543742	**Good identification**						**543742**
Staphylococcus haemolyticus		100.0	0.43	ESL 1			
543750	**Good identification**						**543750**
Staphylococcus haemolyticus		100.0	0.43	ESL 1			
543752	**Good identification**						**543752**
Staphylococcus haemolyticus		100.0	0.38	ESL 1			
544401	**Good identification**						**544401**
					NOV		
Staphylococcus schleiferi subsp. schleiferi		97.53	0.36	GLR 1	1		
Staphylococcus cohnii subsp. cohnii		2.48	0.10	NIT 90	99		
544452	**Good identification**						**544452**
Staphylococcus kloosii		100.0	0.33	VPT 10			
544501	**Good identification**						**544501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	GLR 1			
544550	**Good identification**						**544550**
Staphylococcus kloosii		100.0	0.39	VPT 10			
544552	**Low discrimination**						**544552**
					NOV	YEP	
Staphylococcus kloosii		74.02	0.39	VPT 10	99	99	
Staphylococcus caprae		25.98	0.02	ARG 1	1	1	
544572	**Good identification**						**544572**
Staphylococcus kloosii		100.0	0.27	VPT 10			
545450	**Good identification**						**545450**
Staphylococcus kloosii		100.0	0.33	VPT 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
545452	**Good identification**					**545452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
545550	**Good identification**					**545550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
545552	**Good identification**					**545552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
545572	**Good identification**					**545572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
546001	**Acceptable identification**					**546001**
Staphylococcus schleiferi subsp. schleiferi		94.52	0.46	GLR 1		
Staphylococcus carnosus subsp. carnosus		3.01	0.15			
Staphylococcus capitis subsp. capitis		2.47	0.12			
546003	**Acceptable identification**					**546003**
Staphylococcus carnosus subsp. carnosus		92.66	0.27	BGA 90		
Staphylococcus schleiferi subsp. schleiferi		7.34	0.04	GLR 1		
546011	**Low discrimination**					**546011**
Staphylococcus capitis subsp. capitis		71.61	0.35	GLR 1		
Staphylococcus carnosus subsp. carnosus		26.31	0.27	PHS 11		
Staphylococcus schleiferi subsp. schleiferi		2.08	0.04			
546013	**Good identification**					**546013**
Staphylococcus carnosus subsp. carnosus		98.63	0.39	BGA 90		
Staphylococcus capitis subsp. capitis		1.37	0.00	GLR 1		
546101	**Good identification**					**546101**
Staphylococcus schleiferi subsp. schleiferi		99.21	0.46	GLR 1		
Staphylococcus carnosus subsp. carnosus		0.79	0.02			
546113	**Low discrimination**					**546113**
					COA	
Staphylococcus carnosus subsp. carnosus		89.72	0.27	BGA 90	1	
Staphylococcus schleiferi subsp. coagulans		10.28	0.12	GLR 1	99	
546211	**Good identification**					**546211**
Staphylococcus capitis subsp. capitis		100.0	0.40	GLR 1		
546303	**Good identification**					**546303**
Staphylococcus hyicus		99.06	0.35	VPT 1		
Staphylococcus schleiferi subsp. coagulans		0.94	0.02	TRE 92		
546341	**Good identification**					**546341**
Staphylococcus epidermidis		100.0	0.27	URE 92		
546342	**Low discrimination**					**546342**
					PYR	
Staphylococcus epidermidis		82.87	0.32	URE 92	1	
Staphylococcus haemolyticus		17.13	0.19	GLR 1	98	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
546343	**Good identification**					**546343**
Staphylococcus epidermidis		98.09	0.37	URE 92		
Staphylococcus hyicus		1.91	-.06	GLR 1		
546401	**Very good identification**					**546401**
Staphylococcus schleiferi subsp. schleiferi		99.55	0.55	GLR 1		
Staphylococcus carnosus subsp. carnosus		0.45	0.07			
546413	**Good identification**					**546413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	BGA 90		
546501	**Very good identification**					**546501**
Staphylococcus schleiferi subsp. schleiferi		99.82	0.55	GLR 1		
Staphylococcus hyicus		0.18	-.06			
546553	**Good identification**					**546553**
Staphylococcus caprae		100.0	0.39	URE 90		
546603	**Good identification**					**546603**
Staphylococcus hyicus		100.0	0.35	VPT 1		
546640	**Low discrimination**					**546640**
					YEP	
Staphylococcus haemolyticus		62.62	0.47	PHS 1	20	
Staphylococcus petrasii subsp. croceilyticus		19.62	0.18		99	
Staphylococcus petrasii subsp. jettensis		14.74	0.18		1	
546642	**Acceptable identification**					**546642**
Staphylococcus haemolyticus		93.13	0.41	PHS 1		
Staphylococcus petrasii subsp. jettensis		6.88	0.02			
546650	**Low discrimination**					**546650**
					PYR	
Staphylococcus haemolyticus		84.83	0.42	PHS 1	98	
Staphylococcus pasteuri		13.54	0.18		1	
Staphylococcus warneri		1.63	0.14		8	
546652	**Low discrimination**					**546652**
					PYR	
Staphylococcus haemolyticus		89.62	0.37	PHS 1	98	
Staphylococcus pasteuri		10.39	0.10		1	
546701	**Acceptable identification**					**546701**
Staphylococcus hyicus		94.58	0.35	VPT 1		
Staphylococcus schleiferi subsp. schleiferi		5.42	0.14	LAC 92		
546703	**Very good identification**					**546703**
Staphylococcus hyicus		100.0	0.57	VPT 1		
546740	**Good identification**					**546740**
Staphylococcus haemolyticus		100.0	0.48	PHS 1		
546742	**Good identification**					**546742**
Staphylococcus haemolyticus		100.0	0.43	PHS 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
546750	**Good identification**						**546750**
Staphylococcus haemolyticus		100.0	0.43	PHS 1			
546752	**Low discrimination**						**546752**
Staphylococcus haemolyticus		79.25	0.38	PHS 1			
Staphylococcus caprae		20.75	0.13				
546753	**Low discrimination**						**546753**
					COA	PYR	
Staphylococcus caprae		69.22	0.30	URE 90	1	75	
Staphylococcus aureus subsp. aureus		29.67	0.18	GLR 1	99	10	
Staphylococcus haemolyticus		1.12	0.07		1	98	
552640	**Low discrimination**						**552640**
					YEP		
Staphylococcus haemolyticus		62.62	0.47	ORN 1	20		
Staphylococcus petrasii subsp. croceilyticus		19.62	0.18		99		
Staphylococcus petrasii subsp. jettensis		14.74	0.18		1		
552642	**Acceptable identification**						**552642**
Staphylococcus haemolyticus		93.13	0.41	ORN 1			
Staphylococcus petrasii subsp. jettensis		6.88	0.02				
552643	**Good identification**						**552643**
Staphylococcus lugdunensis		97.09	0.35	ARG 5			
Staphylococcus haemolyticus		2.91	0.10	GLR 2			
552650	**Low discrimination**						**552650**
					PYR		
Staphylococcus haemolyticus		84.83	0.42	ORN 1	98		
Staphylococcus pasteuri		13.54	0.18		1		
Staphylococcus warneri		1.63	0.14		8		
552652	**Low discrimination**						**552652**
					PYR		
Staphylococcus haemolyticus		89.62	0.37	ORN 1	98		
Staphylococcus pasteuri		10.39	0.10		1		
552740	**Good identification**						**552740**
Staphylococcus haemolyticus		100.0	0.48	ORN 1			
552742	**Good identification**						**552742**
Staphylococcus haemolyticus		100.0	0.43	ORN 1			
552743	**Good identification**						**552743**
Staphylococcus lugdunensis		96.73	0.35	ARG 5			
Staphylococcus haemolyticus		3.27	0.11	GLR 2			
552750	**Good identification**						**552750**
Staphylococcus haemolyticus		100.0	0.43	ORN 1			
552752	**Good identification**						**552752**
Staphylococcus haemolyticus		100.0	0.38	ORN 1			
560240	**Low discrimination**						**560240**
					PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		71.61	0.27	URE 98	8		
Staphylococcus haemolyticus		28.39	0.11	TRE 95	98		
560440	**Low discrimination**						**560440**
Staphylococcus warneri		58.31	0.27	URE 98			
Staphylococcus auricularis		41.69	0.15	SUC 95			
560600	**Low discrimination**						**560600**
Staphylococcus warneri		77.95	0.27	URE 98			
Staphylococcus auricularis		22.05	0.07	MLT 95			
560640	**Low discrimination**						**560640**
					PYR		
Staphylococcus warneri		64.58	0.53	URE 98	8		
Staphylococcus haemolyticus		17.90	0.34		98		
Staphylococcus petrasii subsp. jettensis		7.22	0.10		99		
560642	**Low discrimination**						**560642**
					PYR		
Staphylococcus warneri		65.49	0.43	URE 98	8		
Staphylococcus haemolyticus		30.63	0.29		98		
Staphylococcus petrasii subsp. jettensis		3.88	-.05		99		
560650	**Low discrimination**						**560650**
					PYR		
Staphylococcus warneri		77.92	0.50	URE 98	8		
Staphylococcus haemolyticus		17.52	0.29		98		
Staphylococcus pasteuri		4.57	0.10		1		
560652	**Low discrimination**						**560652**
					NOV	PYR	
Staphylococcus warneri		62.49	0.40	URE 98	1	8	
Staphylococcus haemolyticus		23.71	0.24		2	98	
Staphylococcus saprophyticus subsp. saprophyticus		9.32	0.02		99	7	
560740	**Low discrimination**						**560740**
					PYR		
Staphylococcus warneri		50.26	0.43	URE 98	8		
Staphylococcus haemolyticus		49.75	0.35		98		
560742	**Low discrimination**						**560742**
					PYR		
Staphylococcus haemolyticus		62.55	0.30	NIT 99	98		
Staphylococcus warneri		37.45	0.33		8		
560750	**Low discrimination**						**560750**
					PYR		
Staphylococcus warneri		55.47	0.40	URE 98	8		
Staphylococcus haemolyticus		44.53	0.31		98		
562240	**Good identification**						**562240**
					PYR		
Staphylococcus haemolyticus		97.65	0.52	TRE 93	98		
Staphylococcus auricularis		1.29	0.12		40		
Staphylococcus warneri		1.07	0.19		8		
562242	**Good identification**						**562242**
					PYR		
Staphylococcus haemolyticus		99.36	0.47	TRE 93	98		
Staphylococcus warneri		0.64	0.09		8		
562250	**Good identification**						**562250**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus haemolyticus		98.67	0.47	TRE 93	98	
Staphylococcus warneri		1.33	0.16		8	
562252	**Good identification**					**562252**
Staphylococcus haemolyticus		100.0	0.42	TRE 93		
562340	**Very good identification**					**562340**
					PYR	
Staphylococcus haemolyticus		99.70	0.53	TRE 93	98	
Staphylococcus warneri		0.31	0.09		8	
562342	**Good identification**					**562342**
					PYR	
Staphylococcus haemolyticus		99.15	0.48	TRE 93	98	
Staphylococcus epidermidis		0.86	0.04		1	
562350	**Good identification**					**562350**
Staphylococcus haemolyticus		100.0	0.48	TRE 93		
562352	**Good identification**					**562352**
Staphylococcus haemolyticus		100.0	0.43	TRE 93		
562400	**Good identification**					**562400**
Staphylococcus auricularis		100.0	0.27	VPT 12		
562440	**Low discrimination**					**562440**
					PYR	
Staphylococcus auricularis		53.05	0.35	VPT 12	40	
Staphylococcus haemolyticus		43.42	0.34	BGA 14	98	
Staphylococcus warneri		3.53	0.19	GLR 3	8	
562442	**Low discrimination**					**562442**
					PYR	
Staphylococcus haemolyticus		77.55	0.29	SUC 99	98	
Staphylococcus auricularis		18.71	0.15		40	
Staphylococcus warneri		3.74	0.09		8	
562450	**Acceptable identification**					**562450**
					PYR	
Staphylococcus haemolyticus		90.88	0.29	SUC 99	98	
Staphylococcus warneri		9.12	0.16		8	
562540	**Low discrimination**					**562540**
					PYR	
Staphylococcus haemolyticus		87.48	0.35	SUC 99	98	
Staphylococcus auricularis		10.53	0.15		40	
Staphylococcus warneri		1.99	0.09		8	
562542	**Good identification**					**562542**
Staphylococcus haemolyticus		100.0	0.30	SUC 99		
562550	**Good identification**					**562550**
Staphylococcus haemolyticus		100.0	0.31	SUC 99		
562600	**Low discrimination**					**562600**
					PYR	
Staphylococcus haemolyticus		63.91	0.34	MLT 99	98	
Staphylococcus auricularis		30.89	0.27		40	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		5.20	0.19		8	
562602	**Low discrimination**					**562602**
Staphylococcus haemolyticus		72.55	0.29	MLT 99		
Staphylococcus simulans		17.02	0.14			
Staphylococcus auricularis		6.93	0.07			
562610	**Low discrimination**					**562610**
					PYR	
Staphylococcus haemolyticus		83.04	0.29	MLT 99	98	
Staphylococcus pasteuri		8.63	0.02		1	
Staphylococcus warneri		8.33	0.16		8	
562613	**Good identification**					**562613**
Staphylococcus simulans		100.0	0.37	VPT 4		
562640	**Very good identification**					**562640**
					PYR	
Staphylococcus haemolyticus		95.26	0.75		98	
Staphylococcus warneri		1.49	0.45		8	
Staphylococcus petrasii subsp. croceilyticus		1.21	0.18		90	
562641	**Acceptable identification**					**562641**
Staphylococcus haemolyticus		92.44	0.44	MNS 3		
Staphylococcus auricularis		3.78	0.15			
Staphylococcus petrasii subsp. petrasii		2.83	0.05			
562642	**Very good identification**					**562642**
					PYR	
Staphylococcus haemolyticus		98.47	0.70		98	
Staphylococcus warneri		0.91	0.35		8	
Staphylococcus petrasii subsp. jettensis		0.29	0.02		99	
562643	**Low discrimination**					**562643**
Staphylococcus haemolyticus		87.37	0.39	MNS 3		
Staphylococcus simulans		10.07	0.17			
Staphylococcus petrasii subsp. petrasii		2.56	0.00			
562650	**Very good identification**					**562650**
					PYR	
Staphylococcus haemolyticus		97.49	0.71		98	
Staphylococcus warneri		1.88	0.42		8	
Staphylococcus pasteuri		0.63	0.18		1	
562651	**Good identification**					**562651**
Staphylococcus haemolyticus		100.0	0.39	MNS 3		
562652	**Very good identification**					**562652**
					PYR	
Staphylococcus haemolyticus		97.72	0.65		98	
Staphylococcus warneri		1.12	0.33		8	
Staphylococcus simulans		0.71	0.19		70	
562653	**Low discrimination**					**562653**
Staphylococcus haemolyticus		56.02	0.34	MNS 3		
Staphylococcus simulans		43.98	0.30			
562660	**Good identification**					**562660**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus haemolyticus		100.0	0.34	XYL 1		
562662	**Good identification**					**562662**
Staphylococcus haemolyticus		100.0	0.29	XYL 1		
562670	**Good identification**					**562670**
Staphylococcus haemolyticus		100.0	0.29	XYL 1		
562700	**Acceptable identification**					**562700**
					PYR	
Staphylococcus haemolyticus		93.42	0.35	MLT 99	98	
Staphylococcus auricularis		4.45	0.07		40	
Staphylococcus warneri		2.13	0.09		8	
562702	**Good identification**					**562702**
Staphylococcus haemolyticus		100.0	0.30	MLT 99		
562710	**Good identification**					**562710**
Staphylococcus haemolyticus		100.0	0.31	MLT 99		
562740	**Very good identification**					**562740**
					PYR	
Staphylococcus haemolyticus		99.45	0.76		98	
Staphylococcus warneri		0.44	0.35		8	
Staphylococcus auricularis		0.11	0.15		40	
562741	**Good identification**					**562741**
Staphylococcus haemolyticus		98.22	0.45	MNS 3		
Staphylococcus petrasii subsp. petrasii		1.78	0.02			
562742	**Very good identification**					**562742**
					PYR	
Staphylococcus haemolyticus		99.74	0.71		98	
Staphylococcus warneri		0.26	0.25		8	
562743	**Good identification**					**562743**
Staphylococcus haemolyticus		98.30	0.40	MNS 3		
Staphylococcus petrasii subsp. petrasii		1.70	-.04			
562750	**Very good identification**					**562750**
					PYR	
Staphylococcus haemolyticus		99.46	0.72		98	
Staphylococcus warneri		0.54	0.32		8	
562751	**Good identification**					**562751**
Staphylococcus haemolyticus		100.0	0.41	MNS 3		
562752	**Very good identification**					**562752**
					PYR	
Staphylococcus haemolyticus		99.68	0.66		98	
Staphylococcus warneri		0.32	0.22		8	
562753	**Acceptable identification**					**562753**
Staphylococcus haemolyticus		94.29	0.35	MNS 3		
Staphylococcus simulans		5.71	0.08			
562760	**Good identification**					**562760**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus haemolyticus		100.0	0.35	XYL 1		
562762	**Good identification**					**562762**
Staphylococcus haemolyticus		100.0	0.30	XYL 1		
562770	**Good identification**					**562770**
Staphylococcus haemolyticus		100.0	0.31	XYL 1		
563640	**Good identification**					**563640**
Staphylococcus haemolyticus		100.0	0.34	ESL 1		
563642	**Good identification**					**563642**
Staphylococcus haemolyticus		100.0	0.29	ESL 1		
563650	**Good identification**					**563650**
Staphylococcus haemolyticus		100.0	0.29	ESL 1		
563740	**Good identification**					**563740**
Staphylococcus haemolyticus		95.85	0.35	ESL 1		
Rothia mucilaginosa		4.15	-.04			
563742	**Good identification**					**563742**
Staphylococcus haemolyticus		100.0	0.30	ESL 1		
563750	**Good identification**					**563750**
Staphylococcus haemolyticus		100.0	0.31	ESL 1		
564001	**Good identification**					**564001**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	GLR 1		
564101	**Good identification**					**564101**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.30	GLR 1		
564401	**Good identification**					**564401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	GLR 1		
564450	**Good identification**					**564450**
Staphylococcus kloosii		100.0	0.33	VPT 10		
564452	**Good identification**					**564452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
564501	**Good identification**					**564501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	GLR 1		
564550	**Good identification**					**564550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
564552	**Good identification**					**564552**
Staphylococcus kloosii		100.0	0.39	VPT 10		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
564572	**Good identification**					**564572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
565450	**Good identification**					**565450**
Staphylococcus kloosii		100.0	0.33	VPT 10		
565452	**Good identification**					**565452**
Staphylococcus kloosii		100.0	0.33	VPT 10		
565550	**Good identification**					**565550**
Staphylococcus kloosii		100.0	0.39	VPT 10		
565552	**Good identification**					**565552**
Staphylococcus kloosii		100.0	0.39	VPT 10		
565572	**Good identification**					**565572**
Staphylococcus kloosii		100.0	0.27	VPT 10		
566001	**Low discrimination**					**566001**
Staphylococcus schleiferi subsp. schleiferi		83.94	0.49	GLR 1		
Staphylococcus carnosus subsp. carnosus		16.06	0.34			
566003	**Good identification**					**566003**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	GLR 1		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
566011	**Good identification**					**566011**
Staphylococcus carnosus subsp. carnosus		98.70	0.47	GLR 1		
Staphylococcus schleiferi subsp. schleiferi		1.30	0.08			
566013	**Very good identification**					**566013**
Staphylococcus carnosus subsp. carnosus		100.0	0.59	GLR 1		
566101	**Good identification**					**566101**
					COA	
Staphylococcus schleiferi subsp. schleiferi		95.15	0.49	GLR 1	1	
Staphylococcus carnosus subsp. carnosus		4.55	0.22		1	
Staphylococcus schleiferi subsp. coagulans		0.30	0.02		99	
566103	**Low discrimination**					**566103**
					COA	
Staphylococcus carnosus subsp. carnosus		86.56	0.34	GLR 1	1	
Staphylococcus schleiferi subsp. coagulans		8.87	0.18		99	
Staphylococcus schleiferi subsp. schleiferi		4.57	0.08		1	
566111	**Acceptable identification**					**566111**
					COA	
Staphylococcus carnosus subsp. carnosus		93.62	0.34	GLR 1	1	
Staphylococcus schleiferi subsp. schleiferi		4.94	0.08		1	
Staphylococcus schleiferi subsp. coagulans		1.44	0.01		99	
566113	**Good identification**					**566113**
					COA	
Staphylococcus carnosus subsp. carnosus		97.69	0.47	GLR 1	1	
Staphylococcus schleiferi subsp. coagulans		2.31	0.17		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
566401	**Very good identification**					**566401**
Staphylococcus schleiferi subsp. schleiferi		97.34	0.59	GLR 1		
Staphylococcus carnosus subsp. carnosus		2.66	0.27			
566403	**Acceptable identification**					**566403**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	GLR 1		
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18			
566411	**Acceptable identification**					**566411**
Staphylococcus carnosus subsp. carnosus		91.54	0.39	GLR 1		
Staphylococcus schleiferi subsp. schleiferi		8.46	0.18			
566413	**Good identification**					**566413**
Staphylococcus carnosus subsp. carnosus		100.0	0.51	GLR 1		
566501	**Very good identification**					**566501**
Staphylococcus schleiferi subsp. schleiferi		99.32	0.59	GLR 1		
Staphylococcus carnosus subsp. carnosus		0.68	0.14			
566503	**Low discrimination**					**566503**
Staphylococcus carnosus subsp. carnosus		70.17	0.27	GLR 1		
Staphylococcus schleiferi subsp. schleiferi		25.93	0.18			
Staphylococcus hyicus		3.89	-.04			
566511	**Low discrimination**					**566511**
Staphylococcus carnosus subsp. carnosus		73.02	0.27	GLR 1		
Staphylococcus schleiferi subsp. schleiferi		26.98	0.18			
566513	**Good identification**					**566513**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	GLR 1		
566613	**Low discrimination**					**566613**
Staphylococcus simulans		82.59	0.27	VPT 4		
Staphylococcus carnosus subsp. carnosus		17.41	0.10	URE 98		
566640	**Good identification**					**566640**
Staphylococcus haemolyticus		100.0	0.34	PHS 1		
566642	**Good identification**					**566642**
Staphylococcus haemolyticus		100.0	0.29	PHS 1		
566650	**Good identification**					**566650**
Staphylococcus haemolyticus		100.0	0.29	PHS 1		
566703	**Good identification**					**566703**
Staphylococcus hyicus		100.0	0.38	VPT 1		
566740	**Good identification**					**566740**
Staphylococcus haemolyticus		100.0	0.35	PHS 1		
566742	**Good identification**					**566742**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus haemolyticus		100.0	0.30	PHS 1		
566750	**Good identification**					**566750**
Staphylococcus haemolyticus		100.0	0.31	PHS 1		
567003	**Good identification**					**567003**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	GLR 1		
567011	**Good identification**					**567011**
Staphylococcus carnosus subsp. carnosus		100.0	0.27	GLR 1		
567013	**Good identification**					**567013**
Staphylococcus carnosus subsp. carnosus		100.0	0.39	GLR 1		
567113	**Low discrimination**					**567113**
					COA	
Staphylococcus carnosus subsp. carnosus		83.03	0.27	GLR 1	1	
Staphylococcus schleiferi subsp. coagulans		16.98	0.17	ESL 10	99	
567413	**Good identification**					**567413**
Staphylococcus carnosus subsp. carnosus		100.0	0.32	GLR 1		
572640	**Good identification**					**572640**
Staphylococcus haemolyticus		100.0	0.34	ORN 1		
572642	**Good identification**					**572642**
Staphylococcus haemolyticus		100.0	0.29	ORN 1		
572650	**Good identification**					**572650**
Staphylococcus haemolyticus		100.0	0.29	ORN 1		
572740	**Good identification**					**572740**
Staphylococcus haemolyticus		100.0	0.35	ORN 1		
572742	**Good identification**					**572742**
Staphylococcus haemolyticus		100.0	0.30	ORN 1		
572750	**Good identification**					**572750**
Staphylococcus haemolyticus		100.0	0.31	ORN 1		
600000	**Low discrimination**					**600000**
					OXI	
Dermacoccus nishinomiyaensis		58.47	0.59	ARG 1	93	
Micrococcus sp.		31.35	0.44		89	
Staphylococcus carnosus subsp. utilis		6.31	0.29		1	
600002	**Low discrimination**					**600002**
					OXI	
Kocuria varians		49.48	0.30	ARG 1	13	
Dermacoccus nishinomiyaensis		30.73	0.18		93	
Micrococcus sp.		16.48	0.03		89	
600010	**Low discrimination**					**600010**
					YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		60.57	0.43	ARG 1	99		
Macrococcus bovicus		29.33	0.35		99		
Dermacoccus nishinomiyaensis		6.14	0.18		1		
600011	**Good identification**						**600011**
					OXI	NOV	
Staphylococcus capitis subsp. capitis		97.48	0.31	URE 1	1	1	
Macrococcus equipersicus		2.52	0.02	NIT 89	99	99	
600020	**Low discrimination**						**600020**
					OXI		
Kocuria varians		54.95	0.32	ARG 1	13		
Dermacoccus nishinomiyaensis		27.40	0.18		93		
Micrococcus sp.		14.69	0.03		89		
600050	**Low discrimination**						**600050**
					OXI	NOV	
Macrococcus equipersicus		69.06	0.52	ARG 1	99	99	
Macrococcus bovicus		28.86	0.43		99	99	
Staphylococcus warneri		2.08	0.31		1	1	
600053	**Good identification**						**600053**
Staphylococcus capitis subsp. ureolyticus		100.0	0.30	NIT 95			
600100	**Low discrimination**						**600100**
					OXI		
Dermacoccus nishinomiyaensis		66.67	0.44	ARG 1	93		
Kocuria varians		30.27	0.45		13		
Micrococcus sp.		1.90	0.03		89		
600102	**Acceptable identification**						**600102**
					OXI		
Kocuria varians		92.69	0.33	ARG 1	13		
Dermacoccus nishinomiyaensis		7.31	0.03		93		
600110	**Low discrimination**						**600110**
					YEP		
Macrococcus equipersicus		64.53	0.36	ARG 1	99		
Macrococcus bovicus		31.25	0.28		99		
Dermacoccus nishinomiyaensis		2.91	0.03		1		
600120	**Acceptable identification**						**600120**
					OXI		
Kocuria varians		94.05	0.35	ARG 1	13		
Dermacoccus nishinomiyaensis		5.95	0.03		93		
600140	**Low discrimination**						**600140**
					NOV	OXI	
Kocuria varians		62.20	0.31	ARG 1	1	13	
Macrococcus equipersicus		19.90	0.18		99	99	
Staphylococcus warneri		11.61	0.23		1	1	
600150	**Low discrimination**						**600150**
					OXI	NOV	
Macrococcus equipersicus		69.44	0.45	ARG 1	99	99	
Macrococcus bovicus		29.02	0.35		99	99	
Staphylococcus warneri		1.54	0.20		1	1	
600200	**Low discrimination**						**600200**
					OXI		
Dermacoccus nishinomiyaensis		45.96	0.34	ARG 1	93		
Micrococcus sp.		38.19	0.23	SUC 6	89		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		8.08	0.34		1		
600211	**Low discrimination**						**600211**
Staphylococcus capitis subsp. capitis		84.10	0.37	URE 1			
Staphylococcus capitis subsp. ureolyticus		15.90	0.13	NIT 89			
600213	**Acceptable identification**						**600213**
Staphylococcus capitis subsp. ureolyticus		90.47	0.30	NIT 95			
Staphylococcus simulans		8.20	0.22	MLT 96			
Staphylococcus capitis subsp. capitis		1.33	0.02				
600240	**Low discrimination**						**600240**
					COA		
Staphylococcus warneri		80.65	0.60	TRE 95	1		
Staphylococcus aureus subsp. anaerobius		8.77	0.18		99		
Staphylococcus hominis subsp. hominis		4.27	0.28		1		
600242	**Low discrimination**						**600242**
					NOV		
Staphylococcus warneri		79.58	0.50	TRE 95	1		
Staphylococcus hominis subsp. hominis		14.85	0.29		1		
Staphylococcus hominis subsp. novobiosepticus		3.16	0.04		99		
600243	**Acceptable identification**						**600243**
Staphylococcus capitis subsp. ureolyticus		94.99	0.30	NIT 95			
Staphylococcus epidermidis		2.00	0.10	MAN 96			
Staphylococcus warneri		1.38	0.15				
600250	**Low discrimination**						**600250**
					OXI	NOV	
Staphylococcus warneri		71.75	0.57	TRE 95	1	1	
Macrococcus bovicus		13.08	0.31		99	99	
Macrococcus equipersicus		12.38	0.32		99	99	
600251	**Good identification**						**600251**
Staphylococcus capitis subsp. ureolyticus		98.83	0.41	NIT 95			
Staphylococcus warneri		0.95	0.22	LAC 88			
Staphylococcus capitis subsp. capitis		0.22	-.05				
600252	**Low discrimination**						**600252**
					NOV		
Staphylococcus warneri		60.76	0.47	TRE 95	1		
Staphylococcus capitis subsp. ureolyticus		28.67	0.18		1		
Staphylococcus saprophyticus subsp. saprophyticus		6.79	0.07		99		
600253	**Very good identification**						**600253**
					COA		
Staphylococcus capitis subsp. ureolyticus		98.92	0.59	NIT 95	1		
Staphylococcus delphini		0.87	0.18		99		
Staphylococcus simulans		0.17	0.15		1		
600340	**Low discrimination**						**600340**
					NOV		
Staphylococcus warneri		67.38	0.50	TRE 95	1		
Staphylococcus hominis subsp. novobiosepticus		15.75	0.20		99		
Staphylococcus hominis subsp. hominis		12.25	0.29		1		
600342	**Low discrimination**						**600342**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		42.79	0.40	TRE 95	1		
Staphylococcus hominis subsp. hominis		27.39	0.30		1		
Staphylococcus hominis subsp. novobiosepticus		16.16	0.14		99		
600350	**Low discrimination**						**600350**
					OXI	NOV	
Staphylococcus warneri		65.84	0.47	TRE 95	1	1	
Macrococcus bovicus		16.28	0.23		99	99	
Macrococcus equipersicus		15.41	0.24		99	99	
600352	**Low discrimination**						**600352**
Staphylococcus warneri		88.33	0.37	TRE 95			
Staphylococcus hominis subsp. hominis		11.67	0.13				
600400	**Low discrimination**						**600400**
Staphylococcus carnosus subsp. utilis		89.53	0.44	URE 1			
Staphylococcus auricularis		3.64	0.32				
Staphylococcus warneri		1.77	0.34				
600402	**Low discrimination**						**600402**
Staphylococcus piscifermentans		38.01	0.32	ESL 90			
Staphylococcus carnosus subsp. utilis		26.78	0.03	NIT 99			
Staphylococcus warneri		17.51	0.24				
600410	**Low discrimination**						**600410**
					OXI	NOV	
Macrococcus equipersicus		52.68	0.43	ARG 1	99	99	
Macrococcus bovicus		38.26	0.39		99	99	
Staphylococcus warneri		4.30	0.31		1	1	
600440	**Low discrimination**						**600440**
					OXI	NOV	
Staphylococcus warneri		71.71	0.60	SUC 95	1	1	
Staphylococcus auricularis		18.07	0.39		1	1	
Macrococcus equipersicus		4.77	0.26		99	99	
600442	**Low discrimination**						**600442**
Staphylococcus warneri		81.22	0.50	SUC 95			
Staphylococcus piscifermentans		9.28	0.32				
Staphylococcus auricularis		6.82	0.20				
600450	**Low discrimination**						**600450**
					OXI	NOV	
Macrococcus equipersicus		44.31	0.52	ARG 1	99	99	
Macrococcus bovicus		27.77	0.47		99	99	
Staphylococcus warneri		25.40	0.57		1	1	
600452	**Low discrimination**						**600452**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		48.28	0.27	ARG 8	99		
Staphylococcus warneri		44.12	0.47	SUC 91	1		
Staphylococcus piscifermentans		2.98	0.25		1		
600500	**Low discrimination**						**600500**
Staphylococcus piscifermentans		26.51	0.32	ESL 90			
Kocuria varians		21.26	0.22	NIT 99			
Staphylococcus carnosus subsp. utilis		18.68	0.03				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
600502	**Low discrimination**						**600502**
Staphylococcus piscifermentans		80.14	0.36	ESL 90			
Kocuria varians		12.08	0.10	NIT 99			
Staphylococcus warneri		7.77	0.13				
600510	**Low discrimination**						**600510**
					OXI	NOV	
Macrococcus equipersicus		53.54	0.36	ARG 1	99	99	
Macrococcus bovicus		38.88	0.31		99	99	
Staphylococcus piscifermentans		4.36	0.25		1	1	
600512	**Low discrimination**						**600512**
Staphylococcus piscifermentans		85.92	0.28	ESL 90			
Staphylococcus warneri		14.08	0.11	NIT 99			
600540	**Low discrimination**						**600540**
					OXI	NOV	
Staphylococcus warneri		75.10	0.50	SUC 95	1	1	
Staphylococcus piscifermentans		9.06	0.32		1	1	
Macrococcus equipersicus		6.78	0.18		99	99	
600542	**Low discrimination**						**600542**
Staphylococcus warneri		64.82	0.40	SUC 95			
Staphylococcus piscifermentans		35.18	0.36				
600550	**Low discrimination**						**600550**
					OXI	NOV	
Macrococcus equipersicus		47.73	0.45	ARG 1	99	99	
Macrococcus bovicus		29.92	0.39		99	99	
Staphylococcus warneri		20.16	0.47		1	1	
600552	**Low discrimination**						**600552**
					NOV		
Staphylococcus warneri		57.50	0.37	SUC 95	1		
Staphylococcus piscifermentans		18.47	0.28		1		
Staphylococcus saprophyticus subsp. saprophyticus		17.33	0.05		99		
600600	**Low discrimination**						**600600**
Staphylococcus warneri		85.26	0.60	MLT 95			
Staphylococcus auricularis		8.50	0.31				
Staphylococcus piscifermentans		3.25	0.32				
600602	**Low discrimination**						**600602**
Staphylococcus warneri		71.63	0.50	MLT 95			
Staphylococcus simulans		12.99	0.25				
Staphylococcus piscifermentans		12.28	0.36				
600603	**Acceptable identification**						**600603**
Staphylococcus simulans		90.68	0.36	BGA 90			
Staphylococcus hyicus		4.36	0.02	NIT 96			
Staphylococcus warneri		3.05	0.15				
600610	**Low discrimination**						**600610**
					OXI	NOV	
Staphylococcus warneri		78.12	0.57	MLT 95	1	1	
Macrococcus bovicus		9.15	0.27		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		5.47	0.19		1	1	
600612	**Low discrimination**						**600612**
					NOV		
Staphylococcus simulans		43.54	0.38	BGA 90	1		
Staphylococcus warneri		43.47	0.47	NIT 96	1		
Staphylococcus saprophyticus subsp. saprophyticus		4.86	0.07		99		
600613	**Good identification**						**600613**
Staphylococcus simulans		98.36	0.48	BGA 90			
Staphylococcus warneri		0.60	0.12	NIT 96			
Staphylococcus capitis subsp. ureolyticus		0.58	-.11				
600640	**Very good identification**						**600640**
Staphylococcus warneri		96.19	0.86				
Staphylococcus auricularis		1.18	0.39				
Staphylococcus hominis subsp. hominis		0.85	0.38				
600641	**Low discrimination**						**600641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		52.21	0.39	VPT 99	1	99	
Staphylococcus warneri		38.94	0.51	NIT 90	1	8	
Kocuria kristinae		5.91	0.20		99	93	
600642	**Good identification**						**600642**
					NOV		
Staphylococcus warneri		92.97	0.76		1		
Staphylococcus hominis subsp. hominis		2.89	0.40		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.63	0.19		99		
600643	**Low discrimination**						**600643**
					PYR		
Staphylococcus petrasii subsp. petrasii		53.03	0.34	VPT 99	99		
Staphylococcus warneri		24.49	0.42	NIT 90	8		
Staphylococcus simulans		17.22	0.29		70		
600650	**Good identification**						**600650**
					NOV		
Staphylococcus warneri		92.24	0.83		1		
Staphylococcus saprophyticus subsp. saprophyticus		3.16	0.32		99		
Staphylococcus pasteuri		2.09	0.35		1		
600651	**Low discrimination**						**600651**
					PYR		
Staphylococcus warneri		77.46	0.49	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		15.93	0.20		99		
Staphylococcus capitis subsp. ureolyticus		4.28	0.00		1		
600652	**Low discrimination**						**600652**
					NOV		
Staphylococcus warneri		60.74	0.74		1		
Staphylococcus saprophyticus subsp. saprophyticus		35.38	0.48		99		
Staphylococcus pasteuri		1.69	0.27		1		
600653	**Low discrimination**						**600653**
Staphylococcus simulans		57.79	0.41	BGA 90			
Staphylococcus capitis subsp. ureolyticus		18.10	0.18	NIT 96			
Staphylococcus warneri		14.88	0.39				
600660	**Good identification**						**600660**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Staphylococcus warneri	100.0	0.45	XYL 1			
600662	**Good identification**						**600662**
	Staphylococcus warneri	100.0	0.35	XYL 1			
600670	**Good identification**						**600670**
					NOV		
	Staphylococcus warneri	96.69	0.42	XYL 1	1		
	Staphylococcus saprophyticus subsp. saprophyticus	3.31	-.09		99		
600672	**Low discrimination**						**600672**
					NOV		
	Staphylococcus warneri	63.20	0.32	XYL 1	1		
	Staphylococcus saprophyticus subsp. saprophyticus	36.81	0.07		99		
600700	**Low discrimination**						**600700**
	Staphylococcus warneri	81.36	0.50	MLT 95			
	Staphylococcus piscifermentans	14.72	0.36				
	Staphylococcus auricularis	2.85	0.11				
600702	**Low discrimination**						**600702**
	Staphylococcus warneri	53.65	0.40	MLT 95			
	Staphylococcus piscifermentans	43.67	0.39				
	Staphylococcus simulans	2.68	0.03				
600710	**Low discrimination**						**600710**
					OXI	NOV	
	Staphylococcus warneri	73.93	0.47	MLT 95	1	1	
	Macrococcus bovicus	11.75	0.19		99	99	
	Staphylococcus piscifermentans	7.92	0.28		1	1	
600712	**Low discrimination**						**600712**
	Staphylococcus warneri	56.90	0.37	MLT 95			
	Staphylococcus piscifermentans	27.41	0.32				
	Staphylococcus simulans	15.69	0.16				
600713	**Low discrimination**						**600713**
					YEP		
	Staphylococcus simulans	71.78	0.27	BGA 90	1		
	Staphylococcus felis	13.00	0.02	NIT 96	1		
	Staphylococcus chromogenes	11.06	0.07	GAL 8	95		
600740	**Very good identification**						**600740**
	Staphylococcus warneri	95.14	0.76				
	Staphylococcus hominis subsp. hominis	2.88	0.39				
	Staphylococcus piscifermentans	0.91	0.36				
600741	**Low discrimination**						**600741**
					OXI	PYR	
	Staphylococcus petrasii subsp. petrasii	69.79	0.36	VPT 99	1	99	
	Staphylococcus warneri	24.66	0.41	NIT 90	1	8	
	Kocuria kristinae	3.95	0.10		99	93	
600742	**Low discrimination**						**600742**
	Staphylococcus warneri	85.86	0.66				
	Staphylococcus hominis subsp. hominis	9.16	0.40				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		3.68	0.39				
600743	**Low discrimination**						**600743**
					YEP	PYR	
Staphylococcus petrasii subsp. petrasii		65.94	0.30	VPT 99	1	99	
Staphylococcus warneri		14.42	0.31	NIT 90	43	8	
Staphylococcus chromogenes		13.58	0.18		95	45	
600750	**Good identification**						**600750**
					OXI	NOV	
Staphylococcus warneri		94.50	0.73		1	1	
Macrococcus bovicus		1.85	0.27		99	99	
Macrococcus equipersicus		1.16	0.24		99	99	
600751	**Low discrimination**						**600751**
					NOV	PYR	
Staphylococcus warneri		67.37	0.38	MNS 2	1	8	
Staphylococcus petrasii subsp. petrasii		29.26	0.16		1	99	
Staphylococcus saprophyticus subsp. bovis		3.38	0.02		99	99	
600752	**Low discrimination**						**600752**
					NOV		
Staphylococcus warneri		82.15	0.63		1		
Staphylococcus saprophyticus subsp. saprophyticus		13.17	0.26		99		
Staphylococcus piscifermentans		2.08	0.32		1		
600760	**Good identification**						**600760**
Staphylococcus warneri		100.0	0.35	XYL 1			
600770	**Good identification**						**600770**
Staphylococcus warneri		100.0	0.32	XYL 1			
601010	**Good identification**						**601010**
Macrococcus equipersicus		97.53	0.50	ARG 1			
Macrococcus bovicus		2.47	0.16				
601040	**Good identification**						**601040**
Macrococcus equipersicus		100.0	0.33	ARG 1			
601050	**Very good identification**						**601050**
Macrococcus equipersicus		97.86	0.59	ARG 1			
Macrococcus bovicus		2.14	0.23				
601110	**Good identification**						**601110**
Macrococcus equipersicus		97.53	0.43	ARG 1			
Macrococcus bovicus		2.47	0.08				
601150	**Good identification**						**601150**
Macrococcus equipersicus		97.86	0.51	ARG 1			
Macrococcus bovicus		2.14	0.16				
601210	**Acceptable identification**						**601210**
Macrococcus equipersicus		93.99	0.29	ARG 1			
Macrococcus bovicus		6.02	0.03	SUC 9			
601250	**Acceptable identification**						**601250**
					OXI	NOV	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		92.35	0.38	ARG 1	99	99	
Macrococcus bovicus		5.10	0.11	SUC 9	99	99	
Staphylococcus warneri		2.55	0.16		1	1	
601350	**Acceptable identification**						**601350**
Macrococcus equipersicus		94.77	0.31	ARG 1			
Macrococcus bovicus		5.23	0.03	SUC 9			
601400	**Low discrimination**						**601400**
					OXI	NOV	
Staphylococcus piscifermentans		68.85	0.48	NIT 99	1	1	
Macrococcus equipersicus		15.74	0.24		99	99	
Staphylococcus carnosus subsp. utilis		8.08	0.03		1	1	
601402	**Good identification**						**601402**
Staphylococcus piscifermentans		100.0	0.52	NIT 99			
601410	**Low discrimination**						**601410**
					OXI	NOV	
Macrococcus equipersicus		86.78	0.50	ARG 1	99	99	
Staphylococcus piscifermentans		8.56	0.41		1	1	
Macrococcus bovicus		3.30	0.19		99	99	
601412	**Acceptable identification**						**601412**
					OXI	NOV	
Staphylococcus piscifermentans		93.61	0.44	NIT 99	1	1	
Macrococcus equipersicus		6.39	0.09		99	99	
601440	**Low discrimination**						**601440**
					OXI	NOV	
Staphylococcus piscifermentans		60.15	0.48	NIT 99	1	1	
Macrococcus equipersicus		37.19	0.33		99	99	
Staphylococcus warneri		2.66	0.19		1	1	
601442	**Good identification**						**601442**
Staphylococcus piscifermentans		99.03	0.52	NIT 99			
Staphylococcus warneri		0.97	0.09				
601450	**Good identification**						**601450**
					OXI	NOV	
Macrococcus equipersicus		93.29	0.59	ARG 1	99	99	
Staphylococcus piscifermentans		3.40	0.41		1	1	
Macrococcus bovicus		3.06	0.27		99	99	
601452	**Low discrimination**						**601452**
					OXI	NOV	
Staphylococcus piscifermentans		84.42	0.44	NIT 99	1	1	
Macrococcus equipersicus		15.58	0.18		99	99	
601500	**Acceptable identification**						**601500**
					OXI	NOV	
Staphylococcus piscifermentans		91.26	0.52	NIT 99	1	1	
Macrococcus equipersicus		5.96	0.16		99	99	
Macrococcus carouselicus		2.78	0.04		99	99	
601502	**Very good identification**						**601502**
Staphylococcus piscifermentans		100.0	0.55	NIT 99			
601510	**Low discrimination**						**601510**
					OXI	NOV	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		71.49	0.43	ARG 1	99	99	
Staphylococcus piscifermentans		24.68	0.44		1	1	
Macrococcus bovicus		2.72	0.12		99	99	
601512	**Good identification**						**601512**
					OXI	NOV	
Staphylococcus piscifermentans		98.09	0.48	NIT 99	1	1	
Macrococcus equipersicus		1.91	0.01		99	99	
601540	**Low discrimination**						**601540**
					OXI	NOV	
Staphylococcus piscifermentans		84.32	0.52	NIT 99	1	1	
Macrococcus equipersicus		14.90	0.25		99	99	
Staphylococcus warneri		0.79	0.09		1	1	
601542	**Very good identification**						**601542**
Staphylococcus piscifermentans		100.0	0.55	NIT 99			
601550	**Low discrimination**						**601550**
					OXI	NOV	
Macrococcus equipersicus		86.17	0.51	ARG 1	99	99	
Staphylococcus piscifermentans		11.00	0.44		1	1	
Macrococcus bovicus		2.82	0.19		99	99	
601552	**Acceptable identification**						**601552**
					OXI	NOV	
Staphylococcus piscifermentans		94.99	0.48	NIT 99	1	1	
Macrococcus equipersicus		5.01	0.10		99	99	
601600	**Low discrimination**						**601600**
					OXI	NOV	
Staphylococcus piscifermentans		84.13	0.52	NIT 99	1	1	
Macrococcus carouselicus		12.12	0.18		99	99	
Staphylococcus warneri		2.48	0.19		1	1	
601602	**Very good identification**						**601602**
Staphylococcus piscifermentans		99.35	0.55	NIT 99			
Staphylococcus warneri		0.65	0.09				
601610	**Low discrimination**						**601610**
					OXI	NOV	
Staphylococcus piscifermentans		50.07	0.44	NIT 99	1	1	
Macrococcus equipersicus		33.47	0.29		99	99	
Macrococcus carouselicus		10.76	0.14		99	99	
601612	**Good identification**						**601612**
Staphylococcus piscifermentans		100.0	0.48	NIT 99			
601640	**Low discrimination**						**601640**
					OXI	NOV	
Staphylococcus piscifermentans		62.47	0.52	NIT 99	1	1	
Staphylococcus warneri		34.98	0.45		1	1	
Macrococcus equipersicus		2.55	0.12		99	99	
601642	**Low discrimination**						**601642**
Staphylococcus piscifermentans		88.93	0.55	NIT 99			
Staphylococcus warneri		11.07	0.35				
601650	**Low discrimination**						**601650**
					OXI	NOV	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Macrococcus equipersicus		45.92	0.38	ARG 1	99	99	
Staphylococcus piscifermentans		25.41	0.44	SUC 9	1	1	
Staphylococcus warneri		24.04	0.42		1	1	
601652	**Low discrimination**						**601652**
					NOV		
Staphylococcus piscifermentans		75.04	0.48	NIT 99	1		
Staphylococcus warneri		15.78	0.32		1		
Staphylococcus saprophyticus subsp. saprophyticus		9.19	0.07		99		
601700	**Very good identification**						**601700**
					OXI	NOV	
Staphylococcus piscifermentans		95.48	0.55	NIT 99	1	1	
Macrococcus carouselicus		3.93	0.10		99	99	
Staphylococcus warneri		0.59	0.09		1	1	
601702	**Very good identification**						**601702**
Staphylococcus piscifermentans		100.0	0.59	NIT 99			
601710	**Low discrimination**						**601710**
					OXI	NOV	
Staphylococcus piscifermentans		78.70	0.48	NIT 99	1	1	
Macrococcus equipersicus		15.03	0.22		99	99	
Macrococcus carouselicus		4.83	0.06		99	99	
601712	**Good identification**						**601712**
Staphylococcus piscifermentans		100.0	0.51	NIT 99			
601740	**Low discrimination**						**601740**
					OXI	NOV	
Staphylococcus piscifermentans		88.53	0.55	NIT 99	1	1	
Staphylococcus warneri		10.44	0.35		1	1	
Macrococcus equipersicus		1.03	0.04		99	99	
601742	**Very good identification**						**601742**
Staphylococcus piscifermentans		97.45	0.59	NIT 99			
Staphylococcus warneri		2.55	0.25				
601750	**Low discrimination**						**601750**
					OXI	NOV	
Staphylococcus piscifermentans		56.87	0.48	NIT 99	1	1	
Macrococcus equipersicus		29.37	0.31		99	99	
Staphylococcus warneri		11.33	0.32		1	1	
601752	**Good identification**						**601752**
Staphylococcus piscifermentans		95.76	0.51	NIT 99			
Staphylococcus warneri		4.24	0.22				
602000	**Low discrimination**						**602000**
					OXI		
Staphylococcus carnosus subsp. utilis		38.42	0.44	URE 1	1		
Dermacoccus nishinomiyaensis		38.18	0.54		93		
Kocuria varians		19.11	0.55		13		
602001	**Low discrimination**						**602001**
Kocuria varians		39.76	0.34	ARG 1			
Staphylococcus capitis subsp. capitis		38.47	0.27	MNS 8			
Staphylococcus carnosus subsp. utilis		9.29	0.03				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
602002	**Low discrimination**					**602002**
Kocuria varians		78.15	0.44	ARG 1		
Staphylococcus piscifermentans		7.99	0.32			
Staphylococcus carnosus subsp. utilis		5.63	0.03			
602011	**Good identification**					**602011**
Staphylococcus capitis subsp. capitis		96.68	0.50	URE 1		
Staphylococcus capitis subsp. ureolyticus		3.32	0.11			
602013	**Low discrimination**					**602013**
Staphylococcus capitis subsp. ureolyticus		83.65	0.28	SUC 96		
Staphylococcus simulans		9.58	0.22	MLT 96		
Staphylococcus capitis subsp. capitis		6.78	0.15			
602020	**Low discrimination**					**602020**
					OXI	
Kocuria varians		85.79	0.46	ARG 1	13	
Staphylococcus carnosus subsp. utilis		4.96	0.03		1	
Dermacoccus nishinomiyaensis		4.93	0.13		93	
602022	**Good identification**					**602022**
Kocuria varians		100.0	0.35	ARG 1		
602040	**Low discrimination**					**602040**
Kocuria varians		44.50	0.42	ARG 1		
Staphylococcus auricularis		30.27	0.37			
Staphylococcus hominis subsp. hominis		8.13	0.28			
602042	**Low discrimination**					**602042**
Kocuria varians		37.99	0.30	ARG 1		
Staphylococcus hominis subsp. hominis		28.90	0.29			
Staphylococcus piscifermentans		17.69	0.32			
602043	**Good identification**					**602043**
Staphylococcus capitis subsp. ureolyticus		96.09	0.28	SUC 96		
Kocuria varians		1.84	0.09	MAN 96		
Staphylococcus simulans		1.24	0.03			
602051	**Good identification**					**602051**
Staphylococcus capitis subsp. ureolyticus		98.79	0.39	SUC 96		
Staphylococcus capitis subsp. capitis		1.21	0.09	LAC 88		
602053	**Very good identification**					**602053**
					COA	
Staphylococcus capitis subsp. ureolyticus		98.70	0.57	SUC 96	1	
Staphylococcus delphini		1.09	0.18		99	
Staphylococcus simulans		0.21	0.15		1	
602060	**Good identification**					**602060**
Kocuria varians		100.0	0.32	ARG 1		
602100	**Low discrimination**					**602100**
					OXI	
Kocuria varians		77.45	0.59	ARG 1	13	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Dermacoccus nishinomiyaensis		19.65	0.39		93	
Staphylococcus piscifermentans		1.49	0.32		1	
602101	**Good identification**					**602101**
					OXI	
Kocuria varians		97.14	0.37	ARG 1	13	
Dermacoccus nishinomiyaensis		2.86	-.02	MNS 8	93	
602102	**Low discrimination**					**602102**
					OXI	
Kocuria varians		89.98	0.48	ARG 1	13	
Staphylococcus piscifermentans		9.20	0.36		1	
Dermacoccus nishinomiyaensis		0.82	-.02		93	
602112	**Low discrimination**					**602112**
Staphylococcus piscifermentans		81.27	0.28	ESL 90		
Kocuria varians		18.73	0.07	TRE 99		
602120	**Good identification**					**602120**
					OXI	
Kocuria varians		99.28	0.50	ARG 1	13	
Dermacoccus nishinomiyaensis		0.72	-.02		93	
602121	**Good identification**					**602121**
Kocuria varians		100.0	0.28	ARG 1		
602122	**Good identification**					**602122**
Kocuria varians		100.0	0.38	ARG 1		
602140	**Low discrimination**					**602140**
Kocuria varians		76.14	0.45	ARG 1		
Staphylococcus hominis subsp. hominis		10.05	0.29			
Staphylococcus piscifermentans		6.67	0.32			
602142	**Low discrimination**					**602142**
Kocuria varians		47.06	0.34	ARG 1		
Staphylococcus hominis subsp. hominis		25.85	0.30			
Staphylococcus piscifermentans		21.92	0.36			
602152	**Low discrimination**					**602152**
Staphylococcus piscifermentans		70.86	0.28	ESL 90		
Staphylococcus hominis subsp. hominis		29.14	0.13	TRE 99		
602160	**Good identification**					**602160**
Kocuria varians		100.0	0.36	ARG 1		
602200	**Low discrimination**					**602200**
					OXI	
Kocuria varians		30.72	0.38	ARG 1	13	
Dermacoccus nishinomiyaensis		28.74	0.29	SUC 12	93	
Staphylococcus auricularis		13.31	0.29		1	
602201	**Low discrimination**					**602201**
Staphylococcus capitis subsp. capitis		70.48	0.32	URE 1		
Staphylococcus capitis subsp. ureolyticus		17.33	0.11	MAN 93		
Kocuria varians		5.35	0.16			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
602202	**Low discrimination**					**602202**
Staphylococcus simulans		28.66	0.27	BGA 90		
Staphylococcus hominis subsp. hominis		23.35	0.29	TRE 95		
Staphylococcus piscifermentans		21.44	0.36			
602203	**Low discrimination**					**602203**
Staphylococcus capitis subsp. ureolyticus		48.95	0.28	MAN 96		
Staphylococcus simulans		33.64	0.38	MLT 96		
Staphylococcus hyicus		11.02	0.21			
602211	**Low discrimination**					**602211**
Staphylococcus capitis subsp. capitis		69.07	0.55	URE 1		
Staphylococcus capitis subsp. ureolyticus		30.67	0.39			
Staphylococcus simulans		0.26	0.09			
602212	**Low discrimination**					**602212**
Staphylococcus simulans		70.24	0.40	BGA 90		
Staphylococcus capitis subsp. ureolyticus		20.74	0.16	TRE 95		
Staphylococcus piscifermentans		5.63	0.28			
602213	**Low discrimination**					**602213**
					COA	
Staphylococcus capitis subsp. ureolyticus		88.23	0.57	MLT 96	1	
Staphylococcus simulans		10.10	0.50		1	
Staphylococcus delphini		0.98	0.18		99	
602220	**Good identification**					**602220**
Kocuria varians		100.0	0.28	ARG 1		
602240	**Low discrimination**					**602240**
					NOV	
Staphylococcus hominis subsp. hominis		83.94	0.69	ARG 12	1	
Staphylococcus warneri		6.86	0.52		1	
Staphylococcus hominis subsp. novobiosepticus		3.54	0.30		99	
602241	**Low discrimination**					**602241**
Staphylococcus capitis subsp. ureolyticus		77.81	0.39	MAN 96		
Staphylococcus hominis subsp. hominis		6.84	0.34	LAC 88		
Staphylococcus epidermidis		6.46	0.30			
602242	**Good identification**					**602242**
Staphylococcus hominis subsp. hominis		90.33	0.71	ARG 12		
Staphylococcus epidermidis		2.53	0.35			
Staphylococcus warneri		2.10	0.43			
602243	**Good identification**					**602243**
Staphylococcus capitis subsp. ureolyticus		92.24	0.57	MAN 96		
Staphylococcus epidermidis		3.30	0.41			
Staphylococcus hominis subsp. hominis		1.30	0.36			
602250	**Low discrimination**					**602250**
Staphylococcus hominis subsp. hominis		54.23	0.52	ARG 12		
Staphylococcus warneri		21.47	0.49	MAN 13		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. ureolyticus		20.42	0.26			
602251	**Very good identification**					**602251**
					COA	
Staphylococcus capitis subsp. ureolyticus		99.46	0.67	LAC 88	1	
Staphylococcus delphini		0.34	0.18		99	
Staphylococcus capitis subsp. capitis		0.09	0.14		1	
602252	**Low discrimination**					**602252**
Staphylococcus capitis subsp. ureolyticus		62.63	0.44	MNS 99		
Staphylococcus hominis subsp. hominis		26.63	0.54			
Staphylococcus simulans		3.97	0.33			
602253	**Very good identification**					**602253**
					COA	
Staphylococcus capitis subsp. ureolyticus		95.42	0.85		1	
Staphylococcus delphini		4.36	0.59		99	
Staphylococcus simulans		0.21	0.43		1	
602260	**Low discrimination**					**602260**
Staphylococcus hominis subsp. hominis		74.41	0.28	ARG 12		
Kocuria varians		19.51	0.15	XYL 1		
Staphylococcus warneri		6.08	0.11			
602262	**Acceptable identification**					**602262**
Staphylococcus hominis subsp. hominis		94.08	0.29	ARG 12		
Kocuria varians		5.93	0.03	XYL 1		
602273	**Good identification**					**602273**
					COA	
Staphylococcus capitis subsp. ureolyticus		95.44	0.44	XYL 1	1	
Staphylococcus delphini		4.36	0.18		99	
Staphylococcus simulans		0.21	0.02		1	
602300	**Low discrimination**					**602300**
Kocuria varians		61.50	0.41	ARG 1		
Staphylococcus hominis subsp. hominis		13.07	0.29	SUC 12		
Staphylococcus piscifermentans		13.01	0.36			
602302	**Low discrimination**					**602302**
Staphylococcus piscifermentans		32.65	0.39	ESL 90		
Kocuria varians		29.03	0.30	TRE 99		
Staphylococcus hominis subsp. hominis		25.68	0.30			
602303	**Low discrimination**					**602303**
					YEP	
Staphylococcus hyicus		71.40	0.43	PHS 92	1	
Staphylococcus chromogenes		24.53	0.37	TRE 92	95	
Staphylococcus simulans		1.65	0.16		1	
602310	**Low discrimination**					**602310**
Staphylococcus piscifermentans		64.06	0.28	ESL 90		
Staphylococcus hominis subsp. hominis		22.44	0.12	TRE 99		
Staphylococcus warneri		13.50	0.13			
602312	**Low discrimination**					**602312**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		50.07	0.32	ESL 90			
Staphylococcus simulans		36.20	0.18	TRE 99			
Staphylococcus hominis subsp. hominis		13.73	0.13				
602320	**Good identification**						**602320**
Kocuria varians		100.0	0.32	ARG 1			
602340	**Low discrimination**						**602340**
					NOV		
Staphylococcus hominis subsp. hominis		81.70	0.70	ARG 12	1		
Staphylococcus hominis subsp. novobiosepticus		9.55	0.39		99		
Staphylococcus epidermidis		4.43	0.40		1		
602341	**Low discrimination**						**602341**
					NOV		
Staphylococcus epidermidis		60.88	0.46	VPT 96	1		
Staphylococcus hominis subsp. novobiosepticus		17.67	0.27		99		
Staphylococcus hominis subsp. hominis		12.34	0.35		1		
602342	**Low discrimination**						**602342**
					NOV		
Staphylococcus hominis subsp. hominis		81.66	0.71	ARG 12	1		
Staphylococcus epidermidis		11.94	0.51		1		
Staphylococcus hominis subsp. novobiosepticus		4.38	0.34		99		
602343	**Low discrimination**						**602343**
					YEP		
Staphylococcus epidermidis		69.85	0.56	VPT 96	1		
Staphylococcus chromogenes		12.75	0.37		95		
Staphylococcus hominis subsp. hominis		5.25	0.36		49		
602350	**Low discrimination**						**602350**
Staphylococcus hominis subsp. hominis		83.50	0.53	ARG 12			
Staphylococcus warneri		9.64	0.39	MAN 13			
Staphylococcus piscifermentans		2.41	0.28				
602352	**Low discrimination**						**602352**
					COA		
Staphylococcus hominis subsp. hominis		81.38	0.54	ARG 12	1		
Staphylococcus delphini		8.07	0.18	MAN 13	99		
Staphylococcus piscifermentans		3.00	0.32		1		
602353	**Low discrimination**						**602353**
					COA	YEP	
Staphylococcus delphini		80.51	0.59	PHS 99	99	1	
Staphylococcus capitis subsp. ureolyticus		17.81	0.44		1	77	
Staphylococcus chromogenes		0.77	0.26		1	95	
602360	**Low discrimination**						**602360**
					NOV		
Staphylococcus hominis subsp. hominis		67.57	0.29	ARG 12	1		
Kocuria varians		24.53	0.18	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		7.90	-.02		99		
602362	**Low discrimination**						**602362**
Staphylococcus hominis subsp. hominis		77.69	0.30	ARG 12			
Staphylococcus epidermidis		11.36	0.10	XYL 1			
Kocuria varians		6.78	0.07				
602400	**Low discrimination**						**602400**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. utilis		79.33	0.59	URE 1			
Staphylococcus piscifermentans		14.30	0.70				
Staphylococcus auricularis		5.53	0.51				
602402	**Good identification**						**602402**
Staphylococcus piscifermentans		91.62	0.73	ESL 90			
Staphylococcus carnosus subsp. utilis		3.42	0.18				
Staphylococcus auricularis		2.62	0.32				
602403	**Low discrimination**						**602403**
Staphylococcus simulans		68.79	0.36	BGA 90			
Staphylococcus piscifermentans		12.95	0.32	SUC 96			
Staphylococcus felis		9.43	0.09				
602410	**Low discrimination**						**602410**
					OXI	NOV	
Staphylococcus piscifermentans		83.72	0.62	ESL 90	1	1	
Staphylococcus carnosus subsp. utilis		10.95	0.18		1	1	
Macrococcus bovicus		2.93	0.19		99	99	
602412	**Low discrimination**						**602412**
Staphylococcus piscifermentans		86.73	0.66	ESL 90			
Staphylococcus simulans		12.98	0.38				
Staphylococcus warneri		0.29	0.13				
602413	**Low discrimination**						**602413**
Staphylococcus simulans		79.74	0.48	BGA 90			
Staphylococcus felis		18.28	0.26	SUC 96			
Staphylococcus piscifermentans		1.61	0.25				
602422	**Low discrimination**						**602422**
Staphylococcus piscifermentans		79.45	0.32	ESL 90			
Kocuria varians		20.55	0.12	XYL 1			
602440	**Low discrimination**						**602440**
Staphylococcus piscifermentans		45.32	0.70	ESL 90			
Staphylococcus auricularis		40.90	0.59				
Staphylococcus warneri		7.73	0.52				
602441	**Low discrimination**						**602441**
Staphylococcus auricularis		74.82	0.39	URE 1			
Staphylococcus petrasii subsp. petrasii		14.03	0.18	MNS 10			
Staphylococcus piscifermentans		7.54	0.28				
602442	**Low discrimination**						**602442**
Staphylococcus piscifermentans		85.78	0.73	ESL 90			
Staphylococcus auricularis		5.73	0.39				
Staphylococcus hominis subsp. hominis		4.48	0.40				
602443	**Low discrimination**						**602443**
Staphylococcus simulans		46.74	0.29	BGA 90			
Staphylococcus piscifermentans		19.58	0.32	SUC 96			

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		14.40	0.20				
602450	**Low discrimination**						**602450**
					NOV	PYR	
Staphylococcus piscifermentans		65.13	0.62	ESL 90	1	15	
Staphylococcus warneri		18.76	0.49		1	8	
Staphylococcus saprophyticus subsp. bovis		6.65	0.31		99	99	
602452	**Low discrimination**						**602452**
Staphylococcus piscifermentans		85.73	0.66	ESL 90			
Staphylococcus simulans		5.76	0.30				
Staphylococcus warneri		5.49	0.40				
602453	**Low discrimination**						**602453**
Staphylococcus simulans		76.23	0.41	BGA 90			
Staphylococcus capitis subsp. ureolyticus		18.90	0.16	SUC 96			
Staphylococcus piscifermentans		3.42	0.25				
602460	**Low discrimination**						**602460**
Staphylococcus piscifermentans		42.11	0.28	ESL 90			
Staphylococcus auricularis		38.00	0.18	XYL 1			
Kocuria varians		12.72	0.09				
602462	**Good identification**						**602462**
Staphylococcus piscifermentans		100.0	0.32	ESL 90			
602500	**Good identification**						**602500**
Staphylococcus piscifermentans		90.19	0.73	ESL 90			
Kocuria varians		3.57	0.36				
Staphylococcus carnosus subsp. utilis		3.37	0.18				
602501	**Low discrimination**						**602501**
Staphylococcus piscifermentans		49.43	0.32	ESL 90			
Kocuria varians		16.83	0.14	MNS 1			
Staphylococcus auricularis		15.58	0.12				
602502	**Very good identification**						**602502**
Staphylococcus piscifermentans		98.96	0.77	ESL 90			
Kocuria varians		0.74	0.25				
Staphylococcus auricularis		0.21	0.12				
602510	**Very good identification**						**602510**
					OXI	NOV	
Staphylococcus piscifermentans		98.43	0.66	ESL 90	1	1	
Micrococcus bovicus		0.98	0.12		99	99	
Staphylococcus warneri		0.31	0.13		1	1	
602512	**Very good identification**						**602512**
Staphylococcus piscifermentans		98.89	0.69	ESL 90			
Staphylococcus simulans		0.86	0.16				
Staphylococcus felis		0.26	-.04				
602513	**Low discrimination**						**602513**
Staphylococcus felis		77.08	0.37	PHS 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus condimenti		9.81	0.14				
Staphylococcus simulans		8.73	0.27				
602520	**Low discrimination**						**602520**
Kocuria varians		57.91	0.26	ARG 1			
Staphylococcus piscifermentans		42.10	0.32	TRE 7			
602522	**Low discrimination**						**602522**
Staphylococcus piscifermentans		79.45	0.36	ESL 90			
Kocuria varians		20.55	0.15	XYL 1			
602532	**Good identification**						**602532**
Staphylococcus piscifermentans		100.0	0.28	ESL 90			
602540	**Low discrimination**						**602540**
Staphylococcus piscifermentans		86.01	0.73	ESL 90			
Staphylococcus auricularis		5.75	0.39				
Staphylococcus hominis subsp. hominis		4.15	0.39				
602541	**Low discrimination**						**602541**
Staphylococcus piscifermentans		36.07	0.32	ESL 90			
Staphylococcus petrasii subsp. petrasii		29.85	0.14	MNS 1			
Staphylococcus auricularis		26.52	0.20				
602542	**Very good identification**						**602542**
Staphylococcus piscifermentans		95.03	0.77	ESL 90			
Staphylococcus hominis subsp. hominis		3.59	0.40				
Staphylococcus warneri		0.76	0.32				
602543	**Low discrimination**						**602543**
					YEP		
Staphylococcus piscifermentans		44.13	0.36	ESL 90	1		
Staphylococcus chromogenes		29.69	0.18	MNS 1	95		
Staphylococcus petrasii subsp. petrasii		13.11	0.09		1		
602550	**Low discrimination**						**602550**
					NOV	PYR	
Staphylococcus piscifermentans		72.64	0.66	ESL 90	1	15	
Staphylococcus saprophyticus subsp. bovis		19.79	0.43		99	99	
Staphylococcus warneri		4.41	0.39		1	8	
602551	**Low discrimination**						**602551**
					NOV		
Staphylococcus saprophyticus subsp. bovis		81.80	0.31	ARG 1	99		
Staphylococcus piscifermentans		12.13	0.25		1		
Staphylococcus caprae		3.47	-.09		1		
602552	**Good identification**						**602552**
					NOV	PYR	
Staphylococcus piscifermentans		92.51	0.69	ESL 90	1	15	
Staphylococcus saprophyticus subsp. bovis		4.20	0.31		99	99	
Staphylococcus warneri		1.25	0.29		1	8	
602560	**Low discrimination**						**602560**
Staphylococcus piscifermentans		76.81	0.32	ESL 90			
Kocuria varians		23.19	0.13	XYL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
602562	**Acceptable identification**					**602562**
Staphylococcus piscifermentans		94.63	0.36	ESL 90		
Kocuria varians		5.37	0.02	XYL 1		
602572	**Good identification**					**602572**
Staphylococcus piscifermentans		100.0	0.28	ESL 90		
602600	**Low discrimination**					**602600**
Staphylococcus piscifermentans		69.07	0.73	ESL 90		
Staphylococcus auricularis		16.44	0.51			
Staphylococcus warneri		7.85	0.52			
602601	**Low discrimination**					**602601**
Staphylococcus auricularis		26.15	0.31	URE 1		
Staphylococcus hyicus		21.97	0.21	MNS 10		
Staphylococcus simulans		12.87	0.23			
602602	**Low discrimination**					**602602**
Staphylococcus piscifermentans		74.69	0.77	ESL 90		
Staphylococcus simulans		19.16	0.54			
Staphylococcus hominis subsp. hominis		2.60	0.40			
602603	**Low discrimination**					**602603**
					YEP	
Staphylococcus simulans		75.88	0.64	BGA 90	1	
Staphylococcus hyicus		15.05	0.43		1	
Staphylococcus chromogenes		6.61	0.39		95	
602610	**Low discrimination**					**602610**
					YEP	
Staphylococcus piscifermentans		74.75	0.66	ESL 90	1	
Staphylococcus warneri		14.36	0.49		43	
Staphylococcus pasteuri		5.74	0.27		86	
602611	**Low discrimination**					**602611**
Staphylococcus simulans		63.41	0.36	BGA 90		
Staphylococcus capitis subsp. capitis		18.08	0.21	LAC 99		
Staphylococcus piscifermentans		5.28	0.25			
602612	**Low discrimination**					**602612**
Staphylococcus simulans		68.96	0.66	BGA 90		
Staphylococcus piscifermentans		28.81	0.69			
Staphylococcus warneri		1.23	0.40			
602613	**Very good identification**					**602613**
					YEP	
Staphylococcus simulans		96.91	0.77	BGA 90	1	
Staphylococcus felis		1.88	0.33		1	
Staphylococcus chromogenes		0.60	0.28		95	
602620	**Low discrimination**					**602620**
Staphylococcus piscifermentans		67.72	0.32	ESL 90		
Staphylococcus auricularis		16.12	0.10	XYL 1		
Kocuria varians		8.47	0.05			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
602622	**Low discrimination**					**602622**
Staphylococcus piscifermentans		79.59	0.36	ESL 90		
Staphylococcus simulans		20.41	0.13	XYL 1		
602633	**Good identification**					**602633**
Staphylococcus simulans		98.10	0.36	BGA 90		
Staphylococcus felis		1.90	-.09	XYL 1		
602640	**Low discrimination**					**602640**
Staphylococcus hominis subsp. hominis		52.15	0.79	ARG 12		
Staphylococcus warneri		25.57	0.79			
Staphylococcus piscifermentans		11.84	0.73			
602641	**Low discrimination**					**602641**
					PYR	
Staphylococcus petrasii subsp. petrasii		84.31	0.59	VPT 99	99	
Staphylococcus hominis subsp. hominis		6.11	0.45		1	
Staphylococcus auricularis		4.19	0.39		40	
602642	**Low discrimination**					**602642**
Staphylococcus hominis subsp. hominis		66.61	0.81	ARG 12		
Staphylococcus piscifermentans		19.32	0.77			
Staphylococcus warneri		9.27	0.69			
602643	**Low discrimination**					**602643**
					YEP	
Staphylococcus petrasii subsp. petrasii		43.68	0.53	VPT 99	1	
Staphylococcus simulans		37.82	0.57		1	
Staphylococcus chromogenes		7.33	0.39		95	
602650	**Low discrimination**					**602650**
Staphylococcus warneri		51.27	0.76			
Staphylococcus hominis subsp. hominis		21.58	0.62			
Staphylococcus piscifermentans		14.05	0.66			
602651	**Low discrimination**					**602651**
					PYR	
Staphylococcus petrasii subsp. petrasii		42.38	0.39	VPT 99	99	
Staphylococcus capitis subsp. ureolyticus		24.06	0.26	MAN 10	1	
Staphylococcus warneri		9.81	0.41		8	
602652	**Low discrimination**					**602652**
Staphylococcus hominis subsp. hominis		27.73	0.64	ARG 12		
Staphylococcus simulans		24.81	0.59	MAN 13		
Staphylococcus piscifermentans		23.07	0.69			
602653	**Low discrimination**					**602653**
					PYR	
Staphylococcus simulans		75.29	0.70	BGA 90	70	
Staphylococcus capitis subsp. ureolyticus		18.67	0.44		1	
Staphylococcus petrasii subsp. petrasii		2.42	0.34		99	
602660	**Low discrimination**					**602660**
Staphylococcus hominis subsp. hominis		53.35	0.38	ARG 12		
Staphylococcus warneri		26.15	0.38	XYL 1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		12.11	0.32			
602662	**Low discrimination**					**602662**
Staphylococcus hominis subsp. hominis		68.37	0.40	ARG 12		
Staphylococcus piscifermentans		19.83	0.36	XYL 1		
Staphylococcus warneri		9.52	0.28			
602670	**Low discrimination**					**602670**
Staphylococcus warneri		52.66	0.35	XYL 1		
Staphylococcus hominis subsp. hominis		22.17	0.21			
Staphylococcus piscifermentans		14.43	0.25			
602673	**Low discrimination**					**602673**
					NOV	
Staphylococcus simulans		76.55	0.29	BGA 90	1	
Staphylococcus capitis subsp. ureolyticus		18.98	0.03	XYL 1	1	
Staphylococcus xylosus		4.47	-.01		99	
602700	**Good identification**					**602700**
Staphylococcus piscifermentans		92.51	0.77	ESL 90		
Staphylococcus hominis subsp. hominis		2.97	0.39			
Staphylococcus warneri		2.21	0.42			
602701	**Low discrimination**					**602701**
					YEP	
Staphylococcus hyicus		64.34	0.43	PHS 92	1	
Staphylococcus chromogenes		28.25	0.39	LAC 92	95	
Staphylococcus piscifermentans		3.82	0.36		1	
602702	**Good identification**					**602702**
Staphylococcus piscifermentans		93.89	0.80	ESL 90		
Staphylococcus hominis subsp. hominis		2.36	0.40			
Staphylococcus simulans		1.40	0.32			
602703	**Low discrimination**					**602703**
					YEP	
Staphylococcus hyicus		70.91	0.65	PHS 92	1	
Staphylococcus chromogenes		24.37	0.59		95	
Staphylococcus simulans		2.70	0.43		1	
602710	**Good identification**					**602710**
Staphylococcus piscifermentans		94.10	0.69	ESL 90		
Staphylococcus warneri		3.81	0.39			
Staphylococcus hominis subsp. hominis		1.05	0.22			
602711	**Low discrimination**					**602711**
					YEP	
Staphylococcus chromogenes		59.61	0.28	PHS 99	95	
Staphylococcus piscifermentans		12.24	0.28	LAC 90	1	
Staphylococcus simulans		8.53	0.14		1	
602712	**Low discrimination**					**602712**
Staphylococcus piscifermentans		85.89	0.73	ESL 90		
Staphylococcus simulans		11.92	0.44			
Staphylococcus warneri		0.77	0.29			
602713	**Low discrimination**					**602713**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					YEP		
Staphylococcus simulans		38.67	0.55	BGA 90	1		
Staphylococcus felis		28.88	0.43	GAL 8	1		
Staphylococcus chromogenes		24.57	0.48		95		
602720	**Low discrimination**						**602720**
Staphylococcus piscifermentans		88.89	0.36	ESL 90			
Kocuria varians		11.12	0.09	XYL 1			
602722	**Good identification**						**602722**
Staphylococcus piscifermentans		100.0	0.39	ESL 90			
602730	**Good identification**						**602730**
Staphylococcus piscifermentans		100.0	0.28	ESL 90			
602732	**Low discrimination**						**602732**
Staphylococcus piscifermentans		87.81	0.32	ESL 90			
Staphylococcus simulans		12.19	0.03	XYL 1			
602740	**Low discrimination**						**602740**
Staphylococcus hominis subsp. hominis		66.25	0.80	ARG 12			
Staphylococcus piscifermentans		20.82	0.77				
Staphylococcus warneri		9.47	0.68				
602741	**Low discrimination**						**602741**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		76.46	0.55	VPT 99	99	1	
Staphylococcus chromogenes		10.36	0.39		45	95	
Staphylococcus hominis subsp. hominis		9.00	0.45		1	49	
602742	**Low discrimination**						**602742**
Staphylococcus hominis subsp. hominis		68.20	0.82	ARG 12			
Staphylococcus piscifermentans		27.39	0.80				
Staphylococcus warneri		2.77	0.59				
602743	**Low discrimination**						**602743**
					PYR	YEP	
Staphylococcus chromogenes		59.47	0.59	PHS 99	45	95	
Staphylococcus petrasii subsp. petrasii		26.26	0.50		99	1	
Staphylococcus hominis subsp. hominis		6.74	0.47		1	49	
602750	**Low discrimination**						**602750**
Staphylococcus hominis subsp. hominis		29.93	0.63	ARG 12			
Staphylococcus piscifermentans		26.98	0.69	MAN 13			
Staphylococcus warneri		20.73	0.65				
602751	**Low discrimination**						**602751**
					NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		41.01	0.43	ARG 1	99	80	
Staphylococcus petrasii subsp. petrasii		32.29	0.36		1	1	
Staphylococcus chromogenes		11.11	0.28		1	95	
602752	**Low discrimination**						**602752**
Staphylococcus piscifermentans		44.08	0.73	ESL 90			
Staphylococcus hominis subsp. hominis		38.27	0.65				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		7.53	0.56				
602753	**Low discrimination**						**602753**
					COA	YEP	
Staphylococcus aureus subsp. aureus		47.81	0.42	VPT 87	99	84	
Staphylococcus chromogenes		22.42	0.48	PHS 99	1	95	
Staphylococcus simulans		15.85	0.48		1	1	
602760	**Low discrimination**						**602760**
Staphylococcus hominis subsp. hominis		66.90	0.39	ARG 12			
Staphylococcus piscifermentans		21.03	0.36	XYL 1			
Staphylococcus warneri		9.56	0.27				
602762	**Low discrimination**						**602762**
Staphylococcus hominis subsp. hominis		69.34	0.40	ARG 12			
Staphylococcus piscifermentans		27.85	0.39	XYL 1			
Staphylococcus warneri		2.81	0.17				
602772	**Low discrimination**						**602772**
Staphylococcus piscifermentans		46.30	0.32	ESL 90			
Staphylococcus hominis subsp. hominis		40.20	0.23	XYL 1			
Staphylococcus warneri		7.90	0.15				
603000	**Low discrimination**						**603000**
					OXI		
Staphylococcus piscifermentans		77.37	0.48	TRE 99	1		
Staphylococcus carnosus subsp. utilis		9.08	0.03		1		
Dermacoccus nishinomiyaensis		9.03	0.13		93		
603002	**Good identification**						**603002**
Staphylococcus piscifermentans		98.91	0.52	TRE 99			
Kocuria varians		1.09	0.03				
603010	**Acceptable identification**						**603010**
					OXI	NOV	
Staphylococcus piscifermentans		90.71	0.41	TRE 99	1	1	
Macrococcus equipersicus		9.29	0.09		99	99	
603012	**Good identification**						**603012**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
603040	**Good identification**						**603040**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
603042	**Good identification**						**603042**
Staphylococcus piscifermentans		100.0	0.52	TRE 99			
603050	**Low discrimination**						**603050**
					OXI	NOV	
Staphylococcus piscifermentans		74.44	0.41	TRE 99	1	1	
Macrococcus equipersicus		20.61	0.18		99	99	
Macrococcus bovicus		4.95	0.04		99	99	
603052	**Good identification**						**603052**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
603100	**Acceptable identification**						**603100**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					OXI		
Staphylococcus piscifermentans		93.18	0.52	TRE 99	1		
Kocuria varians		5.44	0.18		13		
Dermacoccus nishinomiyaensis		1.38	-.02		93		
603102	**Very good identification**						**603102**
Staphylococcus piscifermentans		98.91	0.55	TRE 99			
Kocuria varians		1.09	0.07				
603110	**Good identification**						**603110**
					OXI	NOV	
Staphylococcus piscifermentans		97.16	0.44	TRE 99	1	1	
Macrococcus equipersicus		2.84	0.01		99	99	
603112	**Good identification**						**603112**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
603140	**Good identification**						**603140**
Staphylococcus piscifermentans		98.74	0.52	TRE 99			
Kocuria varians		1.27	0.04				
603142	**Very good identification**						**603142**
Staphylococcus piscifermentans		100.0	0.55	TRE 99			
603150	**Acceptable identification**						**603150**
					OXI	NOV	
Staphylococcus piscifermentans		92.67	0.44	TRE 99	1	1	
Macrococcus equipersicus		7.33	0.10		99	99	
603152	**Good identification**						**603152**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
603200	**Good identification**						**603200**
Staphylococcus piscifermentans		100.0	0.52	TRE 99			
603202	**Very good identification**						**603202**
Staphylococcus piscifermentans		100.0	0.55	TRE 99			
603210	**Low discrimination**						**603210**
					OXI	NOV	
Staphylococcus piscifermentans		86.27	0.44	TRE 99	1	1	
Staphylococcus vitulinus		13.73	0.09		99	99	
603212	**Good identification**						**603212**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
603240	**Low discrimination**						**603240**
					NOV		
Staphylococcus piscifermentans		81.74	0.52	TRE 99	1		
Staphylococcus hominis subsp. hominis		12.64	0.28		1		
Staphylococcus hominis subsp. novobiosepticus		4.59	0.08		99		
603242	**Low discrimination**						**603242**
					NOV		
Staphylococcus piscifermentans		87.63	0.55	TRE 99	1		
Staphylococcus hominis subsp. hominis		10.60	0.29		1		
Staphylococcus hominis subsp. novobiosepticus		1.77	0.02		99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
603250	**Acceptable identification**					**603250**
	Staphylococcus piscifermentans	94.89	0.44	TRE 99		
	Staphylococcus hominis subsp. hominis	5.11	0.11			
603252	**Low discrimination**					**603252**
	Staphylococcus piscifermentans	87.61	0.48	TRE 99		
	Staphylococcus capitis subsp. ureolyticus	8.69	0.03			
	Staphylococcus hominis subsp. hominis	3.70	0.12			
603253	**Good identification**					**603253**
					COA	
	Staphylococcus capitis subsp. ureolyticus	95.44	0.44	ESL 1	1	
	Staphylococcus delphini	4.36	0.18		99	
	Staphylococcus simulans	0.21	0.02		1	
603300	**Very good identification**					**603300**
	Staphylococcus piscifermentans	100.0	0.55	TRE 99		
603302	**Very good identification**					**603302**
	Staphylococcus piscifermentans	100.0	0.59	TRE 99		
603310	**Good identification**					**603310**
	Staphylococcus piscifermentans	100.0	0.48	TRE 99		
603312	**Good identification**					**603312**
	Staphylococcus piscifermentans	100.0	0.51	TRE 99		
603340	**Low discrimination**					**603340**
					NOV	
	Staphylococcus piscifermentans	81.70	0.55	TRE 99	1	
	Staphylococcus hominis subsp. novobiosepticus	9.18	0.17		99	
	Staphylococcus hominis subsp. hominis	9.12	0.29		1	
603342	**Low discrimination**					**603342**
					NOV	
	Staphylococcus piscifermentans	87.68	0.59	TRE 99	1	
	Staphylococcus hominis subsp. hominis	7.66	0.30		1	
	Staphylococcus hominis subsp. novobiosepticus	3.54	0.12		99	
603350	**Good identification**					**603350**
	Staphylococcus piscifermentans	96.25	0.48	TRE 99		
	Staphylococcus hominis subsp. hominis	3.75	0.12			
603352	**Good identification**					**603352**
	Staphylococcus piscifermentans	97.04	0.51	TRE 99		
	Staphylococcus hominis subsp. hominis	2.96	0.13			
603400	**Very good identification**					**603400**
	Staphylococcus piscifermentans	99.34	0.89			
	Staphylococcus carnosus subsp. utilis	0.62	0.18			
	Staphylococcus auricularis	0.04	0.10			
603401	**Good identification**					**603401**
	Staphylococcus piscifermentans	100.0	0.48	MNS 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
603402	**Very good identification**						**603402**
Staphylococcus piscifermentans		100.0	0.93				
603403	**Good identification**						**603403**
Staphylococcus piscifermentans		100.0	0.52	MNS 1			
603410	**Very good identification**						**603410**
					OXI	NOV	
Staphylococcus piscifermentans		99.85	0.82		1	1	
Macrococcus equipersicus		0.10	0.09		99	99	
Macrococcus bovicus		0.04	0.00		99	99	
603411	**Good identification**						**603411**
Staphylococcus piscifermentans		100.0	0.41	MNS 1			
603412	**Very good identification**						**603412**
Staphylococcus piscifermentans		100.0	0.85				
603413	**Acceptable identification**						**603413**
Staphylococcus piscifermentans		94.73	0.44	MNS 1			
Staphylococcus simulans		5.27	0.07				
603420	**Good identification**						**603420**
Staphylococcus piscifermentans		100.0	0.48	XYL 1			
603422	**Good identification**						**603422**
Staphylococcus piscifermentans		100.0	0.52	XYL 1			
603430	**Good identification**						**603430**
Staphylococcus piscifermentans		100.0	0.41	XYL 1			
603432	**Good identification**						**603432**
Staphylococcus piscifermentans		100.0	0.44	XYL 1			
603440	**Very good identification**						**603440**
Staphylococcus piscifermentans		99.88	0.89				
Staphylococcus auricularis		0.10	0.18				
Staphylococcus warneri		0.02	0.11				
603441	**Good identification**						**603441**
Staphylococcus piscifermentans		100.0	0.48	MNS 1			
603442	**Very good identification**						**603442**
Staphylococcus piscifermentans		100.0	0.93				
603443	**Good identification**						**603443**
Staphylococcus piscifermentans		100.0	0.52	MNS 1			
603450	**Very good identification**						**603450**
					OXI	NOV	
Staphylococcus piscifermentans		99.62	0.82		1	1	
Macrococcus equipersicus		0.28	0.18		99	99	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Micrococcus bovicus		0.10	0.07		99	99	
603451	**Good identification**						**603451**
Staphylococcus piscifermentans		100.0	0.41	MNS 1			
603452	**Very good identification**						**603452**
Staphylococcus piscifermentans		100.0	0.85				
603453	**Good identification**						**603453**
Staphylococcus piscifermentans		97.56	0.44	MNS 1			
Staphylococcus simulans		2.44	0.00				
603460	**Good identification**						**603460**
Staphylococcus piscifermentans		100.0	0.48	XYL 1			
603462	**Good identification**						**603462**
Staphylococcus piscifermentans		100.0	0.52	XYL 1			
603470	**Good identification**						**603470**
Staphylococcus piscifermentans		100.0	0.41	XYL 1			
603472	**Good identification**						**603472**
Staphylococcus piscifermentans		100.0	0.44	XYL 1			
603500	**Very good identification**						**603500**
Staphylococcus piscifermentans		100.0	0.93				
603501	**Good identification**						**603501**
Staphylococcus piscifermentans		100.0	0.52	MNS 1			
603502	**Excellent identification**						**603502**
Staphylococcus piscifermentans		100.0	0.96				
603503	**Very good identification**						**603503**
Staphylococcus piscifermentans		100.0	0.55	MNS 1			
603510	**Very good identification**						**603510**
					OXI	NOV	
Staphylococcus piscifermentans		99.97	0.85		1	1	
Macrococcus equipersicus		0.03	0.01		99	99	
603511	**Good identification**						**603511**
Staphylococcus piscifermentans		100.0	0.44	MNS 1			
603512	**Very good identification**						**603512**
Staphylococcus piscifermentans		100.0	0.89				
603513	**Good identification**						**603513**
Staphylococcus piscifermentans		97.23	0.48	MNS 1			
Staphylococcus felis		2.77	-.04				
603520	**Good identification**						**603520**

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
	Staphylococcus piscifermentans	100.0	0.52	XYL 1				
603522	**Very good identification**							**603522**
	Staphylococcus piscifermentans	100.0	0.55	XYL 1				
603530	**Good identification**							**603530**
	Staphylococcus piscifermentans	100.0	0.44	XYL 1				
603532	**Good identification**							**603532**
	Staphylococcus piscifermentans	100.0	0.48	XYL 1				
603540	**Very good identification**							**603540**
	Staphylococcus piscifermentans	100.0	0.93					
603541	**Good identification**							**603541**
	Staphylococcus piscifermentans	100.0	0.52	MNS 1				
603542	**Excellent identification**							**603542**
	Staphylococcus piscifermentans	100.0	0.96					
603543	**Very good identification**							**603543**
	Staphylococcus piscifermentans	100.0	0.55	MNS 1				
603550	**Very good identification**							**603550**
					OXI	NOV	PYR	
	Staphylococcus piscifermentans	99.86	0.85		1	1	15	
	Macrococcus equipersicus	0.08	0.10		99	99	1	
	Staphylococcus saprophyticus subsp. bovis	0.03	0.02		1	99	99	
603551	**Good identification**							**603551**
	Staphylococcus piscifermentans	100.0	0.44	MNS 1				
603552	**Very good identification**							**603552**
	Staphylococcus piscifermentans	100.0	0.89					
603553	**Good identification**							**603553**
	Staphylococcus piscifermentans	100.0	0.48	MNS 1				
603560	**Good identification**							**603560**
	Staphylococcus piscifermentans	100.0	0.52	XYL 1				
603562	**Very good identification**							**603562**
	Staphylococcus piscifermentans	100.0	0.55	XYL 1				
603570	**Good identification**							**603570**
	Staphylococcus piscifermentans	100.0	0.44	XYL 1				
603572	**Good identification**							**603572**
	Staphylococcus piscifermentans	100.0	0.48	XYL 1				
603600	**Very good identification**							**603600**
	Staphylococcus piscifermentans	99.96	0.93					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus auricularis		0.03	0.10				
Staphylococcus warneri		0.01	0.11				
603601	**Good identification**						**603601**
Staphylococcus piscifermentans		100.0	0.52	MNS 1			
603602	**Excellent identification**						**603602**
Staphylococcus piscifermentans		99.97	0.96				
Staphylococcus simulans		0.03	0.13				
603603	**Low discrimination**						**603603**
Staphylococcus piscifermentans		89.08	0.55	MNS 1			
Staphylococcus simulans		8.50	0.23				
Staphylococcus hyicus		1.69	0.02				
603610	**Very good identification**						**603610**
					OXI	NOV	
Staphylococcus piscifermentans		99.91	0.85		1	1	
Staphylococcus vitulinus		0.09	0.04		99	99	
603611	**Good identification**						**603611**
Staphylococcus piscifermentans		100.0	0.44	MNS 1			
603612	**Very good identification**						**603612**
Staphylococcus piscifermentans		99.73	0.89				
Staphylococcus simulans		0.27	0.25				
603613	**Low discrimination**						**603613**
Staphylococcus piscifermentans		52.42	0.48	MNS 1			
Staphylococcus simulans		46.68	0.36				
Staphylococcus felis		0.91	-.09				
603620	**Good identification**						**603620**
Staphylococcus piscifermentans		100.0	0.52	XYL 1			
603622	**Very good identification**						**603622**
Staphylococcus piscifermentans		100.0	0.55	XYL 1			
603630	**Low discrimination**						**603630**
					OXI	NOV	
Staphylococcus piscifermentans		80.51	0.44	XYL 1	1	1	
Staphylococcus vitulinus		19.49	0.13		99	99	
603632	**Good identification**						**603632**
Staphylococcus piscifermentans		100.0	0.48	XYL 1			
603640	**Very good identification**						**603640**
Staphylococcus piscifermentans		99.19	0.93				
Staphylococcus hominis subsp. hominis		0.49	0.38				
Staphylococcus warneri		0.24	0.38				
603641	**Low discrimination**						**603641**
					PYR		
Staphylococcus piscifermentans		87.12	0.52	MNS 1	15		
Staphylococcus petrasii subsp. petrasii		12.01	0.18		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		0.87	0.04		1		
603642	**Excellent identification**						**603642**
Staphylococcus piscifermentans		99.55	0.96				
Staphylococcus hominis subsp. hominis		0.39	0.40				
Staphylococcus warneri		0.05	0.28				
603643	**Good identification**						**603643**
Staphylococcus piscifermentans		90.21	0.55	MNS 1			
Staphylococcus petrasii subsp. petrasii		4.47	0.12				
Staphylococcus simulans		3.87	0.16				
603650	**Very good identification**						**603650**
Staphylococcus piscifermentans		99.34	0.85				
Staphylococcus warneri		0.41	0.35				
Staphylococcus hominis subsp. hominis		0.17	0.21				
603651	**Acceptable identification**						**603651**
					OXI	NOV	
Staphylococcus piscifermentans		92.05	0.44	MNS 1	1	1	
Staphylococcus sciuri		4.66	0.07		99	99	
Staphylococcus petrasii subsp. petrasii		3.29	-.02		1	1	
603652	**Very good identification**						**603652**
Staphylococcus piscifermentans		99.66	0.89				
Staphylococcus hominis subsp. hominis		0.13	0.23				
Staphylococcus simulans		0.12	0.18				
603653	**Low discrimination**						**603653**
Staphylococcus piscifermentans		66.70	0.48	MNS 1			
Staphylococcus simulans		26.68	0.29				
Staphylococcus capitis subsp. ureolyticus		6.62	0.03				
603660	**Good identification**						**603660**
Staphylococcus piscifermentans		100.0	0.52	XYL 1			
603662	**Very good identification**						**603662**
Staphylococcus piscifermentans		100.0	0.55	XYL 1			
603670	**Good identification**						**603670**
Staphylococcus piscifermentans		100.0	0.44	XYL 1			
603672	**Good identification**						**603672**
Staphylococcus piscifermentans		100.0	0.48	XYL 1			
603700	**Excellent identification**						**603700**
Staphylococcus piscifermentans		100.0	0.96				
603701	**Very good identification**						**603701**
					YEP		
Staphylococcus piscifermentans		97.35	0.55	MNS 1	1		
Staphylococcus hyicus		1.84	0.02		1		
Staphylococcus chromogenes		0.81	-.02		95		
603702	**Excellent identification**						**603702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		100.0	1.00				
603703	**Low discrimination**						**603703**
					YEP		
Staphylococcus piscifermentans		83.31	0.59	MNS 1	1		
Staphylococcus hyicus		12.08	0.24		1		
Staphylococcus chromogenes		4.15	0.18		95		
603710	**Very good identification**						**603710**
Staphylococcus piscifermentans		100.0	0.89				
603711	**Good identification**						**603711**
					OXI	NOV	
Staphylococcus piscifermentans		98.29	0.48	MNS 1	1	1	
Staphylococcus sciuri		1.71	0.01		99	99	
603712	**Very good identification**						**603712**
Staphylococcus piscifermentans		99.98	0.92				
Staphylococcus simulans		0.02	0.03				
603713	**Low discrimination**						**603713**
Staphylococcus piscifermentans		86.73	0.51	MNS 1			
Staphylococcus simulans		4.48	0.14				
Staphylococcus felis		3.34	0.02				
603720	**Very good identification**						**603720**
Staphylococcus piscifermentans		100.0	0.55	XYL 1			
603722	**Very good identification**						**603722**
Staphylococcus piscifermentans		100.0	0.59	XYL 1			
603730	**Good identification**						**603730**
Staphylococcus piscifermentans		100.0	0.48	XYL 1			
603732	**Good identification**						**603732**
Staphylococcus piscifermentans		100.0	0.51	XYL 1			
603740	**Excellent identification**						**603740**
Staphylococcus piscifermentans		99.58	0.96				
Staphylococcus hominis subsp. hominis		0.36	0.39				
Staphylococcus warneri		0.05	0.27				
603741	**Good identification**						**603741**
					YEP		
Staphylococcus piscifermentans		92.22	0.55	MNS 1	1		
Staphylococcus petrasii subsp. petrasii		5.65	0.14		1		
Staphylococcus chromogenes		0.77	-.02		95		
603742	**Excellent identification**						**603742**
Staphylococcus piscifermentans		99.71	1.00				
Staphylococcus hominis subsp. hominis		0.28	0.40				
Staphylococcus warneri		0.01	0.17				
603743	**Good identification**						**603743**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		92.81	0.59	MNS 1	1			
Staphylococcus chromogenes		4.63	0.18		95			
Staphylococcus petrasii subsp. petrasii		2.04	0.09		1			
603750	**Very good identification**							**603750**
Staphylococcus piscifermentans		99.71	0.89					
Staphylococcus hominis subsp. hominis		0.12	0.22					
Staphylococcus warneri		0.09	0.24					
603751	**Low discrimination**							**603751**
					OXI	NOV		
Staphylococcus piscifermentans		88.70	0.48	MNS 1	1	1		
Staphylococcus sciuri		5.81	0.13		99	99		
Staphylococcus lentus		2.29	0.07		99	99		
603752	**Very good identification**							**603752**
Staphylococcus piscifermentans		99.87	0.92					
Staphylococcus hominis subsp. hominis		0.10	0.23					
Staphylococcus warneri		0.02	0.15					
603753	**Low discrimination**							**603753**
					OXI	COA	NOV	
Staphylococcus piscifermentans		84.24	0.51	MNS 1	1	1	1	
Staphylococcus aureus subsp. aureus		5.89	0.01		1	99	4	
Staphylococcus lentus		5.15	0.18		99	1	99	
603760	**Very good identification**							**603760**
Staphylococcus piscifermentans		100.0	0.55	XYL 1				
603762	**Very good identification**							**603762**
Staphylococcus piscifermentans		100.0	0.59	XYL 1				
603770	**Good identification**							**603770**
Staphylococcus piscifermentans		100.0	0.48	XYL 1				
603771	**Low discrimination**							**603771**
					OXI			
Staphylococcus equorum		76.14	0.27	ARG 1	1			
Staphylococcus sciuri		8.50	0.11	GLR 90	99			
Staphylococcus gallinarum		7.34	0.04		1			
603772	**Good identification**							**603772**
Staphylococcus piscifermentans		100.0	0.51	XYL 1				
603773	**Low discrimination**							**603773**
					OXI			
Staphylococcus equorum		80.81	0.36	ARG 1	1			
Staphylococcus gallinarum		7.41	0.12	GLR 90	1			
Staphylococcus lentus		6.51	0.18		99			
604000	**Low discrimination**							**604000**
					OXI			
Dermacoccus nishinomiyaensis		89.00	0.33	ARG 1	93			
Micrococcus sp.		9.16	0.03	PHS 5	89			
Staphylococcus carnosus subsp. utilis		1.84	-.12		1			
604240	**Very good identification**							**604240**
					COA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. anaerobius		99.82	0.59	URE 1	99		
Staphylococcus warneri		0.09	0.19		1		
Staphylococcus epidermidis		0.09	0.09		1		
604253	**Very good identification**						**604253**
					COA	YEP	
Staphylococcus delphini		98.68	0.59	NIT 99	99	1	
Staphylococcus capitis subsp. ureolyticus		1.15	0.18		1	77	
Staphylococcus chromogenes		0.11	0.06		1	95	
604303	**Low discrimination**						**604303**
					YEP	COA	
Staphylococcus chromogenes		72.95	0.37	NIT 99	95	1	
Staphylococcus hyicus		24.66	0.24	TRE 92	1	24	
Staphylococcus schleiferi subsp. coagulans		0.87	0.02		1	99	
604341	**Low discrimination**						**604341**
					YEP		
Staphylococcus epidermidis		73.18	0.30	VPT 96	1		
Staphylococcus chromogenes		26.82	0.17	NIT 97	95		
604342	**Good identification**						**604342**
					YEP		
Staphylococcus epidermidis		98.08	0.35	VPT 96	1		
Staphylococcus chromogenes		1.92	-.04	NIT 97	95		
604343	**Low discrimination**						**604343**
					COA	YEP	
Staphylococcus chromogenes		43.95	0.37	NIT 99	1	95	
Staphylococcus epidermidis		42.19	0.41	TRE 92	1	1	
Staphylococcus delphini		13.13	0.18		99	1	
604353	**Very good identification**						**604353**
					COA	YEP	
Staphylococcus delphini		98.25	0.59	NIT 99	99	1	
Staphylococcus chromogenes		0.94	0.26		1	95	
Staphylococcus caprae		0.40	0.11		1	1	
604402	**Good identification**						**604402**
Staphylococcus piscifermentans		100.0	0.28	ESL 90			
604442	**Low discrimination**						**604442**
Staphylococcus piscifermentans		88.29	0.28	ESL 90			
Staphylococcus warneri		11.71	0.09	NIT 99			
604450	**Low discrimination**						**604450**
					OXI		
Staphylococcus kloosii		72.53	0.46	ARG 1	1		
Macrococcus equipersicus		9.92	0.11		99		
Macrococcus bovicus		6.22	0.05		99		
604452	**Acceptable identification**						**604452**
					NOV	YEP	
Staphylococcus kloosii		90.28	0.47	ARG 1	99	99	
Staphylococcus piscifermentans		9.72	0.21		1	1	
604470	**Good identification**						**604470**
Staphylococcus kloosii		100.0	0.34	ARG 1			
604472	**Good identification**						**604472**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.35	ARG 1			
604500	**Good identification**						**604500**
Staphylococcus piscifermentans		100.0	0.28	ESL 90			
604502	**Good identification**						**604502**
Staphylococcus piscifermentans		100.0	0.32	ESL 90			
604513	**Low discrimination**						**604513**
					YEP		
Staphylococcus felis		86.17	0.37	NIT 99	1		
Staphylococcus condimenti		10.96	0.14		1		
Staphylococcus chromogenes		1.50	0.07		95		
604540	**Low discrimination**						**604540**
Staphylococcus piscifermentans		88.84	0.28	ESL 90			
Staphylococcus warneri		11.16	0.09	NIT 99			
604542	**Low discrimination**						**604542**
Staphylococcus piscifermentans		84.47	0.32	ESL 90			
Staphylococcus caprae		15.53	-.05	NIT 99			
604550	**Low discrimination**						**604550**
					NOV	YEP	
Staphylococcus kloosii		87.12	0.52	ARG 1	99	99	
Staphylococcus piscifermentans		5.24	0.21		1	1	
Staphylococcus caprae		3.37	-.05		1	1	
604552	**Low discrimination**						**604552**
					NOV	YEP	
Staphylococcus kloosii		56.77	0.53	ARG 1	99	99	
Staphylococcus caprae		38.51	0.22		1	1	
Staphylococcus piscifermentans		4.72	0.25		1	1	
604553	**Good identification**						**604553**
					COA	YEP	
Staphylococcus caprae		95.65	0.39	VPT 99	1	1	
Staphylococcus aureus subsp. aureus		2.27	0.01	NIT 90	99	84	
Staphylococcus chromogenes		1.06	0.07		1	95	
604570	**Good identification**						**604570**
Staphylococcus kloosii		100.0	0.40	ARG 1			
604572	**Good identification**						**604572**
Staphylococcus kloosii		100.0	0.41	ARG 1			
604600	**Low discrimination**						**604600**
Staphylococcus piscifermentans		71.54	0.28	ESL 90			
Staphylococcus warneri		28.46	0.19	NIT 99			
604602	**Low discrimination**						**604602**
					YEP		
Staphylococcus piscifermentans		57.82	0.32	ESL 90	1		
Staphylococcus simulans		30.59	0.15	NIT 99	1		
Staphylococcus chromogenes		6.48	-.02		95		
604603	**Low discrimination**						**604603**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus chromogenes		64.21	0.39	NIT 99	95	
Staphylococcus hyicus		16.99	0.24	GAL 90	1	
Staphylococcus simulans		10.25	0.26		1	
604612	**Low discrimination**					**604612**
Staphylococcus simulans		80.02	0.28	BGA 90		
Staphylococcus piscifermentans		16.21	0.25	NIT 96		
Staphylococcus felis		3.78	-.09			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
604613	**Low discrimination**							**604613**
					YEP			
Staphylococcus felis		49.20	0.33	NIT 99	1			
Staphylococcus simulans		35.23	0.39		1			
Staphylococcus chromogenes		15.57	0.28		95			
604640	**Low discrimination**							**604640**
					COA			
Staphylococcus warneri		57.95	0.45	PHS 1	1			
Staphylococcus aureus subsp. anaerobius		32.82	0.18		99			
Staphylococcus piscifermentans		7.67	0.28		1			
604642	**Low discrimination**							**604642**
Staphylococcus warneri		52.67	0.35	PHS 1				
Staphylococcus piscifermentans		31.35	0.32					
Staphylococcus simulans		7.45	0.08					
604643	**Acceptable identification**							**604643**
					YEP			
Staphylococcus chromogenes		93.31	0.39	NIT 99	95			
Staphylococcus simulans		6.69	0.19	GAL 90	1			
604650	**Low discrimination**							**604650**
					NOV	YEP		
Staphylococcus warneri		84.82	0.42	PHS 1	1	43		
Staphylococcus piscifermentans		6.64	0.21		1	1		
Staphylococcus kloosii		5.63	0.26		99	99		
604652	**Low discrimination**							**604652**
					NOV			
Staphylococcus warneri		34.11	0.32	PHS 1	1			
Staphylococcus simulans		26.66	0.21		1			
Staphylococcus saprophyticus subsp. saprophyticus		19.86	0.07		99			
604653	**Low discrimination**							**604653**
					COA	YEP		
Staphylococcus simulans		30.81	0.31	BGA 90	1	1		
Staphylococcus chromogenes		30.31	0.28	NIT 96	1	95		
Staphylococcus delphini		25.12	0.18		99	1		
604700	**Low discrimination**							**604700**
					YEP			
Staphylococcus piscifermentans		83.62	0.32	ESL 90	1			
Staphylococcus chromogenes		9.38	-.02	NIT 99	95			
Staphylococcus warneri		7.00	0.09		43			
604701	**Low discrimination**							**604701**
					YEP	PYR	COA	
Staphylococcus chromogenes		77.92	0.39	NIT 99	95	45	1	
Staphylococcus hyicus		20.62	0.24	LAC 90	1	1	24	
Staphylococcus intermedius		1.47	0.05		1	90	99	
604702	**Low discrimination**							**604702**
					YEP			
Staphylococcus piscifermentans		52.63	0.36	ESL 90	1			
Staphylococcus chromogenes		35.41	0.18	NIT 99	95			
Staphylococcus hyicus		11.97	0.04		1			
604703	**Low discrimination**							**604703**
					YEP			
Staphylococcus chromogenes		71.33	0.59	NIT 99	95			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hyicus		24.11	0.45		1			
Staphylococcus felis		3.53	0.26		1			
604711	**Low discrimination**							**604711**
					COA	YEP		
Staphylococcus chromogenes		77.69	0.28	NIT 99	1	95		
Staphylococcus intermedius		14.01	0.14	LAC 90	99	1		
Staphylococcus felis		8.30	0.02		1	1		
604712	**Low discrimination**							**604712**
					YEP			
Staphylococcus piscifermentans		44.46	0.28	ESL 90	1			
Staphylococcus felis		23.14	0.02	NIT 99	1			
Staphylococcus chromogenes		19.68	0.07		95			
604713	**Low discrimination**							**604713**
					COA	YEP		
Staphylococcus felis		47.63	0.43	NIT 99	1	1		
Staphylococcus chromogenes		40.52	0.48		1	95		
Staphylococcus intermedius		4.99	0.30		99	1		
604740	**Low discrimination**							**604740**
Staphylococcus warneri		55.82	0.35	PHS 1				
Staphylococcus piscifermentans		35.07	0.32					
Staphylococcus hominis subsp. hominis		5.18	0.08					
604741	**Good identification**							**604741**
					COA	YEP		
Staphylococcus chromogenes		96.48	0.39	NIT 99	1	95		
Staphylococcus intermedius		2.72	0.09	LAC 90	99	1		
Staphylococcus petrasii subsp. petrasii		0.80	-.05		1	1		
604742	**Low discrimination**							**604742**
					YEP			
Staphylococcus piscifermentans		46.69	0.36	ESL 90	1			
Staphylococcus chromogenes		31.41	0.18	NIT 99	95			
Staphylococcus warneri		16.51	0.25		43			
604743	**Very good identification**							**604743**
					COA	PYR	YEP	
Staphylococcus chromogenes		96.66	0.59	NIT 99	1	45	95	
Staphylococcus intermedius		1.86	0.25		99	90	1	
Staphylococcus aureus subsp. aureus		0.59	0.01		99	10	84	
604750	**Low discrimination**							**604750**
					NOV	YEP		
Staphylococcus warneri		53.16	0.32	PHS 1	1	43		
Staphylococcus kloosii		21.69	0.32		99	99		
Staphylococcus piscifermentans		19.77	0.25		1	1		
604751	**Low discrimination**							**604751**
					COA	PYR	YEP	
Staphylococcus chromogenes		61.59	0.28	NIT 99	1	45	95	
Staphylococcus intermedius		16.66	0.18	LAC 90	99	90	1	
Staphylococcus aureus subsp. aureus		11.94	0.01		99	10	84	
604753	**Low discrimination**							**604753**
					COA	YEP		
Staphylococcus aureus subsp. aureus		56.61	0.42	VPT 87	99	84		
Staphylococcus chromogenes		26.54	0.48	NIT 99	1	95		
Staphylococcus caprae		8.93	0.30		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
605400	**Good identification**							**605400**
Staphylococcus piscifermentans		100.0	0.44	NIT 99				
605402	**Good identification**							**605402**
Staphylococcus piscifermentans		100.0	0.48	NIT 99				
605410	**Low discrimination**							**605410**
					OXI	NOV	YEP	
Staphylococcus piscifermentans		81.38	0.37	NIT 99	1	1	1	
Macrococcus equipersicus		12.50	0.09		99	99	99	
Staphylococcus kloosii		6.12	0.20		1	99	99	
605412	**Acceptable identification**							**605412**
					NOV	YEP		
Staphylococcus piscifermentans		94.85	0.41	NIT 99	1	1		
Staphylococcus kloosii		5.15	0.21		99	99		
605440	**Good identification**							**605440**
Staphylococcus piscifermentans		100.0	0.44	NIT 99				
605442	**Good identification**							**605442**
Staphylococcus piscifermentans		100.0	0.48	NIT 99				
605450	**Low discrimination**							**605450**
					OXI	NOV	YEP	
Staphylococcus kloosii		50.24	0.46	ARG 1	1	99	99	
Staphylococcus piscifermentans		35.16	0.37		1	1	1	
Macrococcus equipersicus		14.60	0.18		99	99	99	
605452	**Low discrimination**							**605452**
					NOV	YEP		
Staphylococcus kloosii		50.78	0.47	ARG 1	99	99		
Staphylococcus piscifermentans		49.22	0.41		1	1		
605470	**Good identification**							**605470**
Staphylococcus kloosii		100.0	0.34	ARG 1				
605472	**Good identification**							**605472**
Staphylococcus kloosii		100.0	0.35	ARG 1				
605500	**Good identification**							**605500**
Staphylococcus piscifermentans		100.0	0.48	NIT 99				
605502	**Good identification**							**605502**
Staphylococcus piscifermentans		100.0	0.52	NIT 99				
605510	**Low discrimination**							**605510**
					OXI	NOV	YEP	
Staphylococcus piscifermentans		87.63	0.41	NIT 99	1	1	1	
Staphylococcus kloosii		8.53	0.26		1	99	99	
Macrococcus equipersicus		3.85	0.01		99	99	99	
605512	**Acceptable identification**							**605512**
					NOV	YEP		
Staphylococcus piscifermentans		93.43	0.44	NIT 99	1	1		
Staphylococcus kloosii		6.57	0.27		99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
605540	**Good identification**							**605540**
Staphylococcus piscifermentans		100.0	0.48	NIT 99				
605542	**Good identification**							**605542**
Staphylococcus piscifermentans		100.0	0.52	NIT 99				
605550	**Low discrimination**							**605550**
					OXI	NOV	YEP	
Staphylococcus kloosii		62.30	0.52	ARG 1	1	99	99	
Staphylococcus piscifermentans		33.70	0.41		1	1	1	
Macrococcus equipersicus		4.00	0.10		99	99	99	
605552	**Low discrimination**							**605552**
					NOV	YEP		
Staphylococcus kloosii		57.18	0.53	ARG 1	99	99		
Staphylococcus piscifermentans		42.82	0.44		1	1		
605570	**Good identification**							**605570**
Staphylococcus kloosii		100.0	0.40	ARG 1				
605572	**Good identification**							**605572**
Staphylococcus kloosii		100.0	0.41	ARG 1				
605600	**Good identification**							**605600**
Staphylococcus piscifermentans		100.0	0.48	NIT 99				
605602	**Good identification**							**605602**
Staphylococcus piscifermentans		100.0	0.52	NIT 99				
605610	**Good identification**							**605610**
Staphylococcus piscifermentans		100.0	0.41	NIT 99				
605612	**Good identification**							**605612**
Staphylococcus piscifermentans		100.0	0.44	NIT 99				
605640	**Good identification**							**605640**
Staphylococcus piscifermentans		100.0	0.48	NIT 99				
605642	**Good identification**							**605642**
Staphylococcus piscifermentans		100.0	0.52	NIT 99				
605650	**Acceptable identification**							**605650**
					NOV	YEP		
Staphylococcus piscifermentans		91.39	0.41	NIT 99	1	1		
Staphylococcus kloosii		8.61	0.26		99	99		
605652	**Acceptable identification**							**605652**
					NOV	YEP		
Staphylococcus piscifermentans		93.63	0.44	NIT 99	1	1		
Staphylococcus kloosii		6.37	0.26		99	99		
605700	**Good identification**							**605700**
Staphylococcus piscifermentans		100.0	0.52	NIT 99				
605702	**Very good identification**							**605702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		100.0	0.55	NIT 99			
605710	**Good identification**						**605710**
Staphylococcus piscifermentans		100.0	0.44	NIT 99			
605712	**Good identification**						**605712**
Staphylococcus piscifermentans		100.0	0.48	NIT 99			
605740	**Good identification**						**605740**
Staphylococcus piscifermentans		100.0	0.52	NIT 99			
605742	**Very good identification**						**605742**
Staphylococcus piscifermentans		100.0	0.55	NIT 99			
605750	**Low discrimination**						**605750**
					NOV	YEP	
Staphylococcus piscifermentans		89.13	0.44	NIT 99	1	1	
Staphylococcus kloosii		10.87	0.32		99	99	
605752	**Acceptable identification**						**605752**
					NOV	YEP	
Staphylococcus piscifermentans		91.91	0.48	NIT 99	1	1	
Staphylococcus kloosii		8.09	0.32		99	99	
606000	**Low discrimination**						**606000**
					OXI		
Dermacoccus nishinomiyaensis		70.87	0.28	ARG 1	93		
Staphylococcus carnosus subsp. utilis		13.69	0.03	PHS 5	1		
Staphylococcus piscifermentans		8.64	0.25		1		
606002	**Low discrimination**						**606002**
Staphylococcus piscifermentans		87.09	0.28	ESL 90			
Kocuria varians		12.91	0.03	TRE 99			
606011	**Low discrimination**						**606011**
Staphylococcus capitis subsp. capitis		73.13	0.31	URE 1			
Staphylococcus carnosus subsp. carnosus		26.87	0.23	PHS 11			
606013	**Low discrimination**						**606013**
					COA		
Staphylococcus carnosus subsp. carnosus		44.98	0.36	URE 1	1		
Staphylococcus felis		31.14	0.26	BGA 90	1		
Staphylococcus delphini		16.58	0.18		99		
606042	**Low discrimination**						**606042**
Staphylococcus piscifermentans		72.04	0.28	ESL 90			
Staphylococcus epidermidis		27.97	0.10	TRE 99			
606053	**Very good identification**						**606053**
					COA	YEP	
Staphylococcus delphini		98.82	0.59	SUC 99	99	1	
Staphylococcus capitis subsp. ureolyticus		0.91	0.16		1	77	
Staphylococcus chromogenes		0.11	0.06		1	95	
606101	**Low discrimination**						**606101**
					YEP	COA	
Staphylococcus schleiferi subsp. coagulans		77.43	0.37	VPT 99	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus chromogenes		14.15	0.17	LAC 86	95	1	
Staphylococcus hyicus		3.74	0.02		1	24	
606102	**Low discrimination**						**606102**
					COA		
Staphylococcus piscifermentans		63.95	0.32	ESL 90	1		
Staphylococcus schleiferi subsp. coagulans		20.96	0.12	TRE 99	99		
Kocuria varians		9.48	0.07		1		
606103	**Low discrimination**						**606103**
					YEP	COA	
Staphylococcus schleiferi subsp. coagulans		69.11	0.53	VPT 99	1	99	
Staphylococcus chromogenes		18.51	0.37		95	1	
Staphylococcus hyicus		6.26	0.24		1	24	
606111	**Low discrimination**						**606111**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		84.75	0.36	VPT 99	99	1	
Staphylococcus carnosus subsp. carnosus		7.65	0.11	LAC 86	1	1	
Staphylococcus chromogenes		4.73	0.06		1	95	
606113	**Low discrimination**						**606113**
					COA		
Staphylococcus schleiferi subsp. coagulans		54.06	0.53	VPT 99	99		
Staphylococcus felis		29.45	0.37		1		
Staphylococcus delphini		4.68	0.18		99		
606140	**Low discrimination**						**606140**
Staphylococcus piscifermentans		53.54	0.28	ESL 90			
Staphylococcus epidermidis		37.20	0.15	TRE 99			
Kocuria varians		9.26	0.04				
606143	**Low discrimination**						**606143**
					COA	YEP	
Staphylococcus chromogenes		55.40	0.37	SUC 99	1	95	
Staphylococcus epidermidis		17.37	0.31	TRE 92	1	1	
Staphylococcus delphini		16.55	0.18		99	1	
606153	**Low discrimination**						**606153**
					COA	YEP	
Staphylococcus delphini		89.81	0.59	SUC 99	99	1	
Staphylococcus caprae		8.96	0.39		1	1	
Staphylococcus chromogenes		0.86	0.26		1	95	
606200	**Low discrimination**						**606200**
					YEP	OXI	
Staphylococcus piscifermentans		66.22	0.28	ESL 90	1	1	
Dermacoccus nishinomiyaensis		23.12	0.03	TRE 99	1	93	
Staphylococcus chromogenes		10.65	-.02		95	1	
606201	**Low discrimination**						**606201**
					YEP		
Staphylococcus chromogenes		77.86	0.39	GAL 90	95		
Staphylococcus hyicus		16.12	0.21	TRE 92	1		
Staphylococcus capitis subsp. capitis		4.91	0.14	LAC 90	1		
606202	**Low discrimination**						**606202**
					YEP		
Staphylococcus piscifermentans		31.77	0.32	ESL 90	1		
Staphylococcus chromogenes		30.67	0.17	TRE 99	95		
Staphylococcus simulans		21.23	0.17		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
606203	**Low discrimination**						**606203**
					YEP	COA	
Staphylococcus chromogenes		74.75	0.59	GAL 90	95	1	
Staphylococcus hyicus		19.78	0.43	TRE 92	1	24	
Staphylococcus delphini		2.03	0.18		1	99	
606211	**Low discrimination**						**606211**
					COA	YEP	
Staphylococcus capitis subsp. capitis		59.31	0.37	URE 1	1	1	
Staphylococcus chromogenes		19.97	0.28	PHS 11	1	95	
Staphylococcus delphini		17.30	0.18		99	1	
606212	**Low discrimination**						**606212**
					COA		
Staphylococcus simulans		44.05	0.30	BGA 90	1		
Staphylococcus delphini		42.76	0.18	TRE 95	99		
Staphylococcus piscifermentans		7.06	0.25		1		
606213	**Low discrimination**						**606213**
					COA	YEP	
Staphylococcus delphini		84.06	0.59	MLT 99	99	1	
Staphylococcus chromogenes		8.82	0.47		1	95	
Staphylococcus felis		3.24	0.33		1	1	
606240	**Low discrimination**						**606240**
					COA		
Staphylococcus epidermidis		50.28	0.40	VPT 96	1		
Staphylococcus hominis subsp. hominis		26.47	0.38		1		
Staphylococcus aureus subsp. anaerobius		17.73	0.18		99		
606241	**Low discrimination**						**606241**
					COA	YEP	
Staphylococcus epidermidis		52.45	0.46	VPT 96	1	1	
Staphylococcus chromogenes		37.06	0.39		1	95	
Staphylococcus delphini		9.06	0.18		99	1	
606242	**Low discrimination**						**606242**
					COA		
Staphylococcus epidermidis		72.21	0.51	VPT 96	1		
Staphylococcus hominis subsp. hominis		14.09	0.39		1		
Staphylococcus delphini		7.31	0.18		99		
606243	**Low discrimination**						**606243**
					COA	YEP	
Staphylococcus delphini		63.62	0.59	MAN 99	99	1	
Staphylococcus chromogenes		23.67	0.59		1	95	
Staphylococcus epidermidis		11.79	0.56		1	1	
606251	**Very good identification**						**606251**
					COA	YEP	
Staphylococcus delphini		95.90	0.59	LAC 99	99	1	
Staphylococcus capitis subsp. ureolyticus		2.89	0.26		1	77	
Staphylococcus chromogenes		1.12	0.28		1	95	
606252	**Very good identification**						**606252**
					COA		
Staphylococcus delphini		98.66	0.59	MNS 99	99		
Staphylococcus simulans		0.46	0.23		1		
Staphylococcus hominis subsp. hominis		0.29	0.22		1		
606253	**Excellent identification**						**606253**
					COA	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus delphini		99.65	1.00		99	1	
Staphylococcus capitis subsp. ureolyticus		0.22	0.44		1	77	
Staphylococcus chromogenes		0.11	0.47		1	95	
606273	**Very good identification**						**606273**
					COA	YEP	
Staphylococcus delphini		99.67	0.59	XYL 1	99	1	
Staphylococcus capitis subsp. ureolyticus		0.22	0.03		1	77	
Staphylococcus chromogenes		0.11	0.06		1	95	
606300	**Low discrimination**						**606300**
					YEP		
Staphylococcus piscifermentans		37.26	0.32	ESL 90	1		
Staphylococcus chromogenes		35.97	0.17	TRE 99	95		
Staphylococcus epidermidis		17.26	0.15		1		
606301	**Low discrimination**						**606301**
					YEP	COA	
Staphylococcus chromogenes		77.33	0.59	TRE 92	95	1	
Staphylococcus hyicus		20.46	0.43	LAC 90	1	24	
Staphylococcus schleiferi subsp. coagulans		1.35	0.27		1	99	
606302	**Low discrimination**						**606302**
					YEP		
Staphylococcus chromogenes		59.12	0.37	TRE 92	95		
Staphylococcus hyicus		19.99	0.24	MNS 99	1		
Staphylococcus piscifermentans		10.21	0.36		1		
606303	**Low discrimination**						**606303**
					YEP	COA	
Staphylococcus chromogenes		73.33	0.78	TRE 92	95	1	
Staphylococcus hyicus		24.79	0.65		1	24	
Staphylococcus schleiferi subsp. coagulans		0.87	0.43		1	99	
606311	**Low discrimination**						**606311**
					COA	YEP	
Staphylococcus chromogenes		83.37	0.47	TRE 92	1	95	
Staphylococcus delphini		8.03	0.18	LAC 90	99	1	
Staphylococcus schleiferi subsp. coagulans		4.76	0.26		99	1	
606313	**Low discrimination**						**606313**
					COA	YEP	
Staphylococcus delphini		46.45	0.59	MLT 99	99	1	
Staphylococcus chromogenes		43.87	0.67		1	95	
Staphylococcus felis		5.99	0.43		1	1	
606323	**Low discrimination**						**606323**
					YEP	COA	
Staphylococcus chromogenes		74.08	0.37	TRE 92	95	1	
Staphylococcus hyicus		25.04	0.24	XYL 1	1	24	
Staphylococcus schleiferi subsp. coagulans		0.88	0.02		1	99	
606340	**Low discrimination**						**606340**
Staphylococcus epidermidis		87.16	0.56	VPT 96			
Staphylococcus hominis subsp. hominis		8.77	0.39				
Staphylococcus piscifermentans		1.90	0.32				
606341	**Low discrimination**						**606341**
					COA	YEP	
Staphylococcus chromogenes		51.94	0.59	TRE 92	1	95	
Staphylococcus epidermidis		46.29	0.61	LAC 90	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus delphini		1.41	0.18		99	1	
606342	**Low discrimination**						**606342**
					YEP		
Staphylococcus epidermidis		88.81	0.66	VPT 96	1		
Staphylococcus chromogenes		5.31	0.37		95		
Staphylococcus hominis subsp. hominis		3.31	0.40		49		
606343	**Low discrimination**						**606343**
					COA	YEP	
Staphylococcus chromogenes		61.75	0.78	TRE 92	1	95	
Staphylococcus epidermidis		19.36	0.72		1	1	
Staphylococcus delphini		18.44	0.59		99	1	
606351	**Low discrimination**						**606351**
					COA	YEP	
Staphylococcus delphini		89.62	0.59	LAC 99	99	1	
Staphylococcus chromogenes		9.40	0.47		1	95	
Staphylococcus epidermidis		0.30	0.20		1	1	
606352	**Very good identification**						**606352**
					COA	YEP	
Staphylococcus delphini		97.21	0.59	MNS 99	99	1	
Staphylococcus chromogenes		0.93	0.26		1	95	
Staphylococcus epidermidis		0.56	0.25		1	1	
606353	**Very good identification**						**606353**
					COA	YEP	
Staphylococcus delphini		98.75	1.00		99	1	
Staphylococcus chromogenes		0.94	0.67		1	95	
Staphylococcus aureus subsp. aureus		0.23	0.42		99	84	
606363	**Low discrimination**						**606363**
					COA	YEP	
Staphylococcus chromogenes		62.03	0.37	TRE 92	1	95	
Staphylococcus epidermidis		19.45	0.31	XYL 1	1	1	
Staphylococcus delphini		18.53	0.18		99	1	
606373	**Very good identification**						**606373**
					COA	YEP	
Staphylococcus delphini		98.79	0.59	XYL 1	99	1	
Staphylococcus chromogenes		0.94	0.26		1	95	
Staphylococcus aureus subsp. aureus		0.23	0.01		99	84	
606400	**Good identification**						**606400**
Staphylococcus piscifermentans		91.75	0.66	ESL 90			
Staphylococcus carnosus subsp. utilis		7.71	0.18				
Staphylococcus auricularis		0.54	0.10				
606402	**Very good identification**						**606402**
Staphylococcus piscifermentans		97.97	0.70	ESL 90			
Staphylococcus felis		1.08	0.09				
Staphylococcus simulans		0.79	0.15				
606403	**Low discrimination**						**606403**
					YEP		
Staphylococcus felis		80.71	0.50	MAN 87	1		
Staphylococcus chromogenes		12.43	0.39		95		
Staphylococcus hyicus		3.29	0.24		1		
606410	**Very good identification**						**606410**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.58	ESL 90		
606412	**Low discrimination**					**606412**
Staphylococcus piscifermentans		80.19	0.62	ESL 90		
Staphylococcus felis		13.82	0.26			
Staphylococcus simulans		6.00	0.28			
606413	**Very good identification**					**606413**
					YEP	
Staphylococcus felis		96.87	0.67		1	
Staphylococcus simulans		1.42	0.39		1	
Staphylococcus chromogenes		0.63	0.28		95	
606422	**Good identification**					**606422**
Staphylococcus piscifermentans		100.0	0.28	ESL 90		
606440	**Very good identification**					**606440**
Staphylococcus piscifermentans		98.10	0.66	ESL 90		
Staphylococcus auricularis		1.34	0.18			
Staphylococcus hominis subsp. hominis		0.30	0.07			
606442	**Very good identification**					**606442**
Staphylococcus piscifermentans		99.23	0.70	ESL 90		
Staphylococcus simulans		0.36	0.08			
Staphylococcus hominis subsp. hominis		0.24	0.09			
606443	**Low discrimination**					**606443**
					YEP	
Staphylococcus chromogenes		82.76	0.39	GAL 90	95	
Staphylococcus simulans		5.93	0.19	SUC 99	1	
Staphylococcus felis		5.43	0.09		1	
606450	**Very good identification**					**606450**
					NOV	
Staphylococcus piscifermentans		96.31	0.58	ESL 90	1	
Staphylococcus saprophyticus subsp. bovis		3.69	0.18		99	
606452	**Very good identification**					**606452**
Staphylococcus piscifermentans		95.13	0.62	ESL 90		
Staphylococcus simulans		3.20	0.21			
Staphylococcus caprae		1.07	0.00			
606462	**Good identification**					**606462**
Staphylococcus piscifermentans		100.0	0.28	ESL 90		
606500	**Very good identification**					**606500**
					YEP	
Staphylococcus piscifermentans		99.83	0.70	ESL 90	1	
Staphylococcus chromogenes		0.17	-.02		95	
606501	**Low discrimination**					**606501**
					YEP	
Staphylococcus chromogenes		62.47	0.39	SUC 99	95	
Staphylococcus hyicus		16.53	0.24	LAC 90	1	
Staphylococcus felis		13.72	0.20		1	
606502	**Very good identification**					**606502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					YEP	
Staphylococcus piscifermentans		96.32	0.73	ESL 90	1	
Staphylococcus felis		2.37	0.20		1	
Staphylococcus chromogenes		0.98	0.18		95	
606503	**Low discrimination**					**606503**
					YEP	
Staphylococcus felis		63.67	0.61	MAN 87	1	
Staphylococcus chromogenes		26.36	0.59		95	
Staphylococcus hyicus		8.91	0.45		1	
606510	**Very good identification**					**606510**
Staphylococcus piscifermentans		99.42	0.62	ESL 90		
Staphylococcus felis		0.58	-.04			
606511	**Low discrimination**					**606511**
					YEP	
Staphylococcus felis		73.76	0.37	LAC 99	1	
Staphylococcus chromogenes		14.16	0.28		95	
Staphylococcus condimenti		9.38	0.14		1	
606512	**Low discrimination**					**606512**
Staphylococcus piscifermentans		68.87	0.66	ESL 90		
Staphylococcus felis		26.48	0.37			
Staphylococcus condimenti		3.37	0.14			
606513	**Low discrimination**					**606513**
					YEP	
Staphylococcus felis		87.09	0.78		1	
Staphylococcus condimenti		11.08	0.55		1	
Staphylococcus chromogenes		1.52	0.48		95	
606520	**Good identification**					**606520**
Staphylococcus piscifermentans		100.0	0.28	ESL 90		
606522	**Good identification**					**606522**
Staphylococcus piscifermentans		100.0	0.32	ESL 90		
606533	**Low discrimination**					**606533**
					YEP	
Staphylococcus felis		87.36	0.37	XYL 1	1	
Staphylococcus condimenti		11.11	0.14		1	
Staphylococcus chromogenes		1.52	0.07		95	
606540	**Very good identification**					**606540**
					YEP	
Staphylococcus piscifermentans		99.48	0.70	ESL 90	1	
Staphylococcus hominis subsp. hominis		0.22	0.08		49	
Staphylococcus chromogenes		0.17	-.02		95	
606541	**Low discrimination**					**606541**
					YEP	
Staphylococcus chromogenes		89.48	0.39	SUC 99	95	
Staphylococcus piscifermentans		5.37	0.28	LAC 90	1	
Staphylococcus caprae		5.15	0.06		1	
606542	**Very good identification**					**606542**
					YEP	
Staphylococcus piscifermentans		97.21	0.73	ESL 90	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus caprae		1.63	0.15		1		
Staphylococcus chromogenes		0.99	0.18		95		
606543	**Low discrimination**						**606543**
					YEP		
Staphylococcus chromogenes		85.62	0.59	SUC 99	95		
Staphylococcus caprae		10.40	0.33		1		
Staphylococcus felis		2.09	0.20		1		
606550	**Low discrimination**						**606550**
					NOV		
Staphylococcus piscifermentans		86.16	0.62	ESL 90	1		
Staphylococcus saprophyticus subsp. bovis		8.80	0.31		99		
Staphylococcus caprae		5.04	0.15		1		
606551	**Low discrimination**						**606551**
					NOV	YEP	
Staphylococcus caprae		70.72	0.33	VPT 99	1	1	
Staphylococcus chromogenes		18.24	0.28	LAC 95	1	95	
Staphylococcus saprophyticus subsp. bovis		4.21	0.18		99	80	
606552	**Low discrimination**						**606552**
					NOV		
Staphylococcus piscifermentans		56.08	0.66	ESL 90	1		
Staphylococcus caprae		41.57	0.41		1		
Staphylococcus saprophyticus subsp. bovis		0.96	0.18		99		
606553	**Low discrimination**						**606553**
					COA	YEP	
Staphylococcus caprae		67.54	0.59	VPT 99	1	1	
Staphylococcus aureus subsp. aureus		17.60	0.42		99	84	
Staphylococcus chromogenes		8.25	0.48		1	95	
606560	**Good identification**						**606560**
Staphylococcus piscifermentans		100.0	0.28	ESL 90			
606562	**Good identification**						**606562**
Staphylococcus piscifermentans		100.0	0.32	ESL 90			
606600	**Very good identification**						**606600**
					YEP		
Staphylococcus piscifermentans		96.91	0.70	ESL 90	1		
Staphylococcus chromogenes		1.81	0.20		95		
Staphylococcus hyicus		0.38	0.02		1		
606601	**Low discrimination**						**606601**
					YEP		
Staphylococcus chromogenes		81.42	0.61	GAL 90	95		
Staphylococcus hyicus		16.86	0.43	LAC 90	1		
Staphylococcus felis		0.99	0.16		1		
606602	**Low discrimination**						**606602**
					YEP		
Staphylococcus piscifermentans		77.73	0.73	ESL 90	1		
Staphylococcus simulans		9.97	0.44		1		
Staphylococcus chromogenes		8.72	0.39		95		
606603	**Low discrimination**						**606603**
					YEP		
Staphylococcus chromogenes		69.57	0.80	GAL 90	95		
Staphylococcus hyicus		18.41	0.65		1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus felis		9.27	0.57		1	
606610	**Very good identification**					**606610**
					YEP	
Staphylococcus piscifermentans		96.72	0.62	ESL 90	1	
Staphylococcus simulans		1.75	0.15		1	
Staphylococcus chromogenes		1.19	0.08		95	
606611	**Low discrimination**					**606611**
					YEP	
Staphylococcus chromogenes		73.63	0.49	GAL 90	95	
Staphylococcus felis		21.15	0.33	LAC 90	1	
Staphylococcus simulans		3.67	0.26		1	
606612	**Low discrimination**					**606612**
Staphylococcus simulans		47.80	0.56	BGA 90		
Staphylococcus piscifermentans		39.94	0.66			
Staphylococcus felis		9.31	0.33			
606613	**Low discrimination**					**606613**
					YEP	
Staphylococcus felis		66.62	0.74		1	
Staphylococcus chromogenes		21.08	0.69		95	
Staphylococcus simulans		11.56	0.67		1	
606622	**Low discrimination**					**606622**
					YEP	
Staphylococcus piscifermentans		80.62	0.32	ESL 90	1	
Staphylococcus simulans		10.34	0.03	XYL 1	1	
Staphylococcus chromogenes		9.04	-.02		95	
606623	**Low discrimination**					**606623**
					YEP	
Staphylococcus chromogenes		69.62	0.39	GAL 90	95	
Staphylococcus hyicus		18.42	0.24	XYL 1	1	
Staphylococcus felis		9.27	0.16		1	
606633	**Low discrimination**					**606633**
					YEP	
Staphylococcus felis		67.11	0.33	XYL 1	1	
Staphylococcus chromogenes		21.24	0.28		95	
Staphylococcus simulans		11.65	0.26		1	
606640	**Low discrimination**					**606640**
Staphylococcus piscifermentans		78.97	0.70	ESL 90		
Staphylococcus hominis subsp. hominis		16.14	0.48			
Staphylococcus warneri		2.58	0.38			
606641	**Very good identification**					**606641**
					YEP	
Staphylococcus chromogenes		97.77	0.61	GAL 90	95	
Staphylococcus petrasii subsp. petrasii		0.99	0.18	LAC 90	1	
Staphylococcus piscifermentans		0.53	0.28		1	
606642	**Low discrimination**					**606642**
					YEP	
Staphylococcus piscifermentans		74.65	0.73	ESL 90	1	
Staphylococcus hominis subsp. hominis		11.94	0.50		49	
Staphylococcus chromogenes		8.37	0.39		95	
606643	**Very good identification**					**606643**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					YEP		
Staphylococcus chromogenes		97.34	0.80	GAL 90	95		
Staphylococcus simulans		1.69	0.47		1		
Staphylococcus hyicus		0.26	0.24		1		
606650	**Low discrimination**						**606650**
					NOV	PYR	
Staphylococcus piscifermentans		79.60	0.62	ESL 90	1	15	
Staphylococcus saprophyticus subsp. bovis		8.13	0.31		99	99	
Staphylococcus hominis subsp. hominis		5.67	0.31		1	1	
606651	**Low discrimination**						**606651**
					COA	YEP	
Staphylococcus chromogenes		85.26	0.49	GAL 90	1	95	
Staphylococcus delphini		6.42	0.18	LAC 90	99	1	
Staphylococcus aureus subsp. aureus		3.04	0.07		99	84	
606652	**Low discrimination**						**606652**
					YEP		
Staphylococcus piscifermentans		55.81	0.66	ESL 90	1		
Staphylococcus simulans		30.01	0.49		1		
Staphylococcus chromogenes		4.12	0.28		95		
606653	**Low discrimination**						**606653**
					COA	YEP	
Staphylococcus chromogenes		39.88	0.69	GAL 90	1	95	
Staphylococcus delphini		33.05	0.59		99	1	
Staphylococcus aureus subsp. aureus		15.62	0.48		99	84	
606660	**Low discrimination**						**606660**
Staphylococcus piscifermentans		83.03	0.28	ESL 90			
Staphylococcus hominis subsp. hominis		16.97	0.07	XYL 1			
606662	**Low discrimination**						**606662**
					YEP		
Staphylococcus piscifermentans		78.61	0.32	ESL 90	1		
Staphylococcus hominis subsp. hominis		12.58	0.09	XYL 1	49		
Staphylococcus chromogenes		8.81	-.02		95		
606663	**Good identification**						**606663**
					YEP		
Staphylococcus chromogenes		98.29	0.39	GAL 90	95		
Staphylococcus simulans		1.71	0.06	XYL 1	1		
606673	**Low discrimination**						**606673**
					COA	YEP	
Staphylococcus chromogenes		40.02	0.28	GAL 90	1	95	
Staphylococcus delphini		33.16	0.18	XYL 1	99	1	
Staphylococcus aureus subsp. aureus		15.68	0.07		99	84	
606700	**Low discrimination**						**606700**
					YEP		
Staphylococcus piscifermentans		87.47	0.73	ESL 90	1		
Staphylococcus chromogenes		9.81	0.39		95		
Staphylococcus hyicus		2.60	0.24		1		
606701	**Low discrimination**						**606701**
					YEP		
Staphylococcus chromogenes		78.59	0.80	LAC 90	95		
Staphylococcus hyicus		20.79	0.65		1		
Staphylococcus felis		0.35	0.26		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
606702	**Low discrimination**						**606702**
					YEP		
Staphylococcus piscifermentans		51.46	0.77	ESL 90	1		
Staphylococcus chromogenes		34.62	0.59		95		
Staphylococcus hyicus		11.70	0.45		1		
606703	**Low discrimination**						**606703**
					YEP		
Staphylococcus chromogenes		71.93	1.00		95		
Staphylococcus hyicus		24.32	0.86		1		
Staphylococcus felis		3.56	0.68		1		
606710	**Good identification**						**606710**
					YEP		
Staphylococcus piscifermentans		92.21	0.66	ESL 90	1		
Staphylococcus chromogenes		6.81	0.28		95		
Staphylococcus felis		0.73	0.02		1		
606711	**Low discrimination**						**606711**
					COA	YEP	
Staphylococcus chromogenes		86.63	0.69	LAC 90	1	95	
Staphylococcus felis		9.26	0.43		1	1	
Staphylococcus intermedius		2.10	0.37		99	1	
606712	**Low discrimination**						**606712**
					YEP		
Staphylococcus piscifermentans		47.49	0.69	ESL 90	1		
Staphylococcus felis		24.72	0.43		1		
Staphylococcus chromogenes		21.03	0.48		95		
606713	**Low discrimination**						**606713**
					YEP		
Staphylococcus felis		50.17	0.85		1		
Staphylococcus chromogenes		42.68	0.89		95		
Staphylococcus condimenti		4.72	0.59		1		
606720	**Low discrimination**						**606720**
					YEP		
Staphylococcus piscifermentans		89.92	0.32	ESL 90	1		
Staphylococcus chromogenes		10.08	-.02	XYL 1	95		
606721	**Low discrimination**						**606721**
					YEP		
Staphylococcus chromogenes		79.08	0.39	XYL 1	95		
Staphylococcus hyicus		20.92	0.24	LAC 90	1		
606722	**Low discrimination**						**606722**
					YEP		
Staphylococcus piscifermentans		52.63	0.36	ESL 90	1		
Staphylococcus chromogenes		35.41	0.18	XYL 1	95		
Staphylococcus hyicus		11.97	0.04		1		
606723	**Low discrimination**						**606723**
					YEP		
Staphylococcus chromogenes		71.98	0.59	XYL 1	95		
Staphylococcus hyicus		24.33	0.45		1		
Staphylococcus felis		3.57	0.26		1		
606731	**Acceptable identification**						**606731**
					YEP		
Staphylococcus chromogenes		90.35	0.28	XYL 1	95		
Staphylococcus felis		9.66	0.02	LAC 90	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
606732	**Low discrimination**							**606732**
					YEP			
Staphylococcus piscifermentans		50.94	0.28	ESL 90	1			
Staphylococcus felis		26.51	0.02	XYL 1	1			
Staphylococcus chromogenes		22.55	0.07		95			
606733	**Low discrimination**							**606733**
					YEP			
Staphylococcus felis		50.20	0.43	XYL 1	1			
Staphylococcus chromogenes		42.71	0.48		95			
Staphylococcus condimenti		4.72	0.18		1			
606740	**Low discrimination**							**606740**
					YEP			
Staphylococcus piscifermentans		78.48	0.73	ESL 90	1			
Staphylococcus hominis subsp. hominis		11.58	0.49		49			
Staphylococcus chromogenes		8.80	0.39		95			
606741	**Very good identification**							**606741**
					YEP	PYR	COA	
Staphylococcus chromogenes		99.02	0.80	LAC 90	95	45	1	
Staphylococcus intermedius		0.38	0.32		1	90	99	
Staphylococcus hyicus		0.27	0.24		1	1	24	
606742	**Low discrimination**							**606742**
					YEP			
Staphylococcus piscifermentans		55.09	0.77	ESL 90	1			
Staphylococcus chromogenes		37.06	0.59		95			
Staphylococcus hominis subsp. hominis		6.36	0.50		49			
606743	**Very good identification**							**606743**
					COA	YEP		
Staphylococcus chromogenes		98.62	1.00		1	95		
Staphylococcus aureus subsp. aureus		0.60	0.42		99	84		
Staphylococcus hyicus		0.34	0.45		24	1		
606750	**Low discrimination**							**606750**
					NOV	YEP		
Staphylococcus piscifermentans		69.41	0.66	ESL 90	1	1		
Staphylococcus saprophyticus subsp. bovis		18.91	0.43		99	80		
Staphylococcus chromogenes		5.12	0.28		1	95		
606751	**Low discrimination**							**606751**
					COA	PYR	YEP	
Staphylococcus chromogenes		79.04	0.69	LAC 90	1	45	95	
Staphylococcus aureus subsp. aureus		15.33	0.42		99	10	84	
Staphylococcus intermedius		2.87	0.41		99	90	1	
606752	**Low discrimination**							**606752**
					COA	YEP		
Staphylococcus piscifermentans		36.41	0.69	ESL 90	1	1		
Staphylococcus aureus subsp. aureus		34.39	0.42		99	84		
Staphylococcus chromogenes		16.12	0.48		1	95		
606753	**Low discrimination**							**606753**
					COA	YEP		
Staphylococcus aureus subsp. aureus		64.70	0.83	VPT 87	99	84		
Staphylococcus chromogenes		30.34	0.89		1	95		
Staphylococcus delphini		2.79	0.59		99	1		
606760	**Low discrimination**							**606760**
					YEP			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		79.38	0.32	ESL 90	1		
Staphylococcus hominis subsp. hominis		11.72	0.08	XYL 1	49		
Staphylococcus chromogenes		8.90	-.02		95		
606761	**Good identification**						**606761**
Staphylococcus chromogenes		100.0	0.39	XYL 1			
606762	**Low discrimination**						**606762**
					YEP		
Staphylococcus piscifermentans		55.92	0.36	ESL 90	1		
Staphylococcus chromogenes		37.62	0.18	XYL 1	95		
Staphylococcus hominis subsp. hominis		6.46	0.09		49		
606763	**Very good identification**						**606763**
					COA	YEP	
Staphylococcus chromogenes		97.94	0.59	XYL 1	1	95	
Staphylococcus lutrae		0.87	0.12		99	1	
Staphylococcus aureus subsp. aureus		0.60	0.01		99	84	
606771	**Low discrimination**						**606771**
					COA	NOV	
Staphylococcus chromogenes		73.95	0.28	XYL 1	1	1	
Staphylococcus aureus subsp. aureus		14.34	0.01	LAC 90	99	4	
Staphylococcus gallinarum		5.83	0.04		1	99	
606772	**Low discrimination**						**606772**
					COA	YEP	
Staphylococcus piscifermentans		38.91	0.28	ESL 90	1	1	
Staphylococcus aureus subsp. aureus		36.75	0.01	XYL 1	99	84	
Staphylococcus chromogenes		17.23	0.07		1	95	
606773	**Low discrimination**						**606773**
					COA	YEP	
Staphylococcus aureus subsp. aureus		62.88	0.42	VPT 87	99	84	
Staphylococcus chromogenes		29.48	0.48	XYL 1	1	95	
Staphylococcus delphini		2.71	0.18		99	1	
607000	**Good identification**						**607000**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
607002	**Good identification**						**607002**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
607010	**Good identification**						**607010**
Staphylococcus piscifermentans		100.0	0.37	TRE 99			
607012	**Good identification**						**607012**
Staphylococcus piscifermentans		100.0	0.41	TRE 99			
607040	**Good identification**						**607040**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
607042	**Good identification**						**607042**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
607050	**Good identification**						**607050**
Staphylococcus piscifermentans		100.0	0.37	TRE 99			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
607052	**Good identification**						**607052**
Staphylococcus piscifermentans		100.0	0.41	TRE 99			
607100	**Good identification**						**607100**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
607101	**Good identification**						**607101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	VPT 99			
607102	**Good identification**						**607102**
					COA		
Staphylococcus piscifermentans		96.62	0.52	TRE 99	1		
Staphylococcus schleiferi subsp. coagulans		3.38	0.12		99		
607103	**Very good identification**						**607103**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		99.43	0.53	VPT 99	99	1	
Staphylococcus piscifermentans		0.29	0.11		1	1	
Staphylococcus chromogenes		0.28	-.04		1	95	
607110	**Good identification**						**607110**
Staphylococcus piscifermentans		100.0	0.41	TRE 99			
607111	**Good identification**						**607111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	VPT 99			
607112	**Acceptable identification**						**607112**
					COA		
Staphylococcus piscifermentans		92.99	0.44	TRE 99	1		
Staphylococcus schleiferi subsp. coagulans		7.01	0.11		99		
607113	**Good identification**						**607113**
					COA		
Staphylococcus schleiferi subsp. coagulans		98.76	0.52	VPT 99	99		
Staphylococcus carnosus subsp. carnosus		0.67	0.04		1		
Staphylococcus felis		0.57	-.04		1		
607140	**Good identification**						**607140**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
607142	**Good identification**						**607142**
Staphylococcus piscifermentans		100.0	0.52	TRE 99			
607150	**Good identification**						**607150**
Staphylococcus piscifermentans		100.0	0.41	TRE 99			
607152	**Good identification**						**607152**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
607200	**Good identification**						**607200**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
607202	**Good identification**						**607202**
Staphylococcus piscifermentans		100.0	0.52	TRE 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
607210	**Good identification**						**607210**
Staphylococcus piscifermentans		100.0	0.41	TRE 99			
607212	**Good identification**						**607212**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
607240	**Good identification**						**607240**
Staphylococcus piscifermentans		100.0	0.48	TRE 99			
607242	**Good identification**						**607242**
Staphylococcus piscifermentans		97.21	0.52	TRE 99			
Staphylococcus epidermidis		2.80	0.10				
607250	**Good identification**						**607250**
Staphylococcus piscifermentans		100.0	0.41	TRE 99			
607252	**Low discrimination**						**607252**
					COA		
Staphylococcus piscifermentans		59.79	0.44	TRE 99	1		
Staphylococcus delphini		40.21	0.18		99		
607253	**Very good identification**						**607253**
					COA	YEP	
Staphylococcus delphini		99.67	0.59	ESL 1	99	1	
Staphylococcus capitis subsp. ureolyticus		0.22	0.03		1	77	
Staphylococcus chromogenes		0.11	0.06		1	95	
607300	**Good identification**						**607300**
Staphylococcus piscifermentans		100.0	0.52	TRE 99			
607301	**Low discrimination**						**607301**
					YEP	COA	
Staphylococcus schleiferi subsp. coagulans		54.99	0.27	VPT 99	1	99	
Staphylococcus chromogenes		33.13	0.17	LAC 86	95	1	
Staphylococcus hyicus		8.76	0.02		1	24	
607302	**Very good identification**						**607302**
					COA	YEP	
Staphylococcus piscifermentans		98.63	0.55	TRE 99	1	1	
Staphylococcus schleiferi subsp. coagulans		0.73	0.02		99	1	
Staphylococcus chromogenes		0.64	-.04		1	95	
607303	**Low discrimination**						**607303**
					YEP	COA	
Staphylococcus schleiferi subsp. coagulans		45.56	0.43	VPT 99	1	99	
Staphylococcus chromogenes		40.21	0.37		95	1	
Staphylococcus hyicus		13.59	0.24		1	24	
607310	**Good identification**						**607310**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
607312	**Good identification**						**607312**
					COA		
Staphylococcus piscifermentans		98.44	0.48	TRE 99	1		
Staphylococcus schleiferi subsp. coagulans		1.56	0.01		99		
607313	**Low discrimination**						**607313**
					COA	YEP	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. coagulans		62.81	0.42	VPT 99	99	1	
Staphylococcus delphini		17.94	0.18		99	1	
Staphylococcus chromogenes		16.94	0.26		1	95	
607340	**Good identification**						**607340**
Staphylococcus piscifermentans		95.11	0.52	TRE 99			
Staphylococcus epidermidis		4.89	0.15				
607342	**Low discrimination**						**607342**
					YEP		
Staphylococcus piscifermentans		89.68	0.55	TRE 99	1		
Staphylococcus epidermidis		9.74	0.25		1		
Staphylococcus chromogenes		0.58	-.04		95		
607343	**Low discrimination**						**607343**
					COA	YEP	
Staphylococcus chromogenes		61.00	0.37	ESL 1	1	95	
Staphylococcus epidermidis		19.13	0.31	TRE 92	1	1	
Staphylococcus delphini		18.22	0.18		99	1	
607350	**Good identification**						**607350**
Staphylococcus piscifermentans		100.0	0.44	TRE 99			
607352	**Low discrimination**						**607352**
					COA		
Staphylococcus piscifermentans		69.04	0.48	TRE 99	1		
Staphylococcus delphini		30.96	0.18		99		
607353	**Very good identification**						**607353**
					COA	YEP	
Staphylococcus delphini		98.75	0.59	ESL 1	99	1	
Staphylococcus chromogenes		0.94	0.26		1	95	
Staphylococcus aureus subsp. aureus		0.23	0.01		99	84	
607400	**Very good identification**						**607400**
Staphylococcus piscifermentans		100.0	0.86				
607401	**Good identification**						**607401**
Staphylococcus piscifermentans		100.0	0.44	MNS 1			
607402	**Very good identification**						**607402**
Staphylococcus piscifermentans		100.0	0.89				
607403	**Low discrimination**						**607403**
					YEP		
Staphylococcus piscifermentans		86.65	0.48	MNS 1	1		
Staphylococcus felis		10.52	0.09		1		
Staphylococcus chromogenes		1.62	-.02		95		
607410	**Very good identification**						**607410**
Staphylococcus piscifermentans		100.0	0.78				
607411	**Good identification**						**607411**
Staphylococcus piscifermentans		95.34	0.37	MNS 1			
Staphylococcus carnosus subsp. carnosus		4.66	-.04				
607412	**Very good identification**						**607412**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.82			
607420	**Good identification**					**607420**
Staphylococcus piscifermentans		100.0	0.44	XYL 1		
607422	**Good identification**					**607422**
Staphylococcus piscifermentans		100.0	0.48	XYL 1		
607430	**Good identification**					**607430**
Staphylococcus piscifermentans		100.0	0.37	XYL 1		
607432	**Good identification**					**607432**
Staphylococcus piscifermentans		100.0	0.41	XYL 1		
607440	**Very good identification**					**607440**
Staphylococcus piscifermentans		100.0	0.86			
607441	**Good identification**					**607441**
Staphylococcus piscifermentans		100.0	0.44	MNS 1		
607442	**Very good identification**					**607442**
Staphylococcus piscifermentans		100.0	0.89			
607443	**Good identification**					**607443**
					YEP	
Staphylococcus piscifermentans		98.17	0.48	MNS 1	1	
Staphylococcus chromogenes		1.83	-.02		95	
607450	**Very good identification**					**607450**
Staphylococcus piscifermentans		100.0	0.78			
607451	**Good identification**					**607451**
Staphylococcus piscifermentans		100.0	0.37	MNS 1		
607452	**Very good identification**					**607452**
Staphylococcus piscifermentans		100.0	0.82			
607453	**Good identification**					**607453**
Staphylococcus piscifermentans		100.0	0.41	MNS 1		
607460	**Good identification**					**607460**
Staphylococcus piscifermentans		100.0	0.44	XYL 1		
607462	**Good identification**					**607462**
Staphylococcus piscifermentans		100.0	0.48	XYL 1		
607470	**Good identification**					**607470**
Staphylococcus piscifermentans		100.0	0.37	XYL 1		
607472	**Good identification**					**607472**
Staphylococcus piscifermentans		100.0	0.41	XYL 1		
607500	**Very good identification**					**607500**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.89			
607501	**Good identification**					**607501**
					YEP	
Staphylococcus piscifermentans		98.17	0.48	MNS 1	1	
Staphylococcus chromogenes		1.83	-.02		95	
607502	**Very good identification**					**607502**
Staphylococcus piscifermentans		100.0	0.93			
607503	**Low discrimination**					**607503**
					YEP	
Staphylococcus piscifermentans		68.69	0.52	MNS 1	1	
Staphylococcus felis		18.61	0.20		1	
Staphylococcus chromogenes		7.70	0.18		95	
607510	**Very good identification**					**607510**
Staphylococcus piscifermentans		100.0	0.82			
607511	**Acceptable identification**					**607511**
Staphylococcus piscifermentans		93.98	0.41	MNS 1		
Staphylococcus felis		6.02	-.04			
607512	**Very good identification**					**607512**
Staphylococcus piscifermentans		99.96	0.85			
Staphylococcus felis		0.04	-.04			
607513	**Low discrimination**					**607513**
Staphylococcus felis		71.22	0.37	ESL 1		
Staphylococcus piscifermentans		16.84	0.44			
Staphylococcus condimenti		9.06	0.14			
607520	**Good identification**					**607520**
Staphylococcus piscifermentans		100.0	0.48	XYL 1		
607522	**Good identification**					**607522**
Staphylococcus piscifermentans		100.0	0.52	XYL 1		
607530	**Good identification**					**607530**
Staphylococcus piscifermentans		100.0	0.41	XYL 1		
607532	**Good identification**					**607532**
Staphylococcus piscifermentans		100.0	0.44	XYL 1		
607540	**Very good identification**					**607540**
Staphylococcus piscifermentans		100.0	0.89			
607541	**Good identification**					**607541**
					YEP	
Staphylococcus piscifermentans		98.17	0.48	MNS 1	1	
Staphylococcus chromogenes		1.83	-.02		95	
607542	**Very good identification**					**607542**
Staphylococcus piscifermentans		100.0	0.93			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
607543	**Low discrimination**					**607543**
					YEP	
Staphylococcus piscifermentans		88.83	0.52	MNS 1	1	
Staphylococcus chromogenes		9.96	0.18		95	
Staphylococcus caprae		1.21	-.09		1	
607550	**Very good identification**					**607550**
Staphylococcus piscifermentans		100.0	0.82			
607551	**Good identification**					**607551**
Staphylococcus piscifermentans		95.45	0.41	MNS 1		
Staphylococcus caprae		4.55	-.09			
607552	**Very good identification**					**607552**
Staphylococcus piscifermentans		99.92	0.85			
Staphylococcus caprae		0.08	0.00			
607553	**Low discrimination**					**607553**
					COA	
Staphylococcus piscifermentans		53.25	0.44	MNS 1	1	
Staphylococcus caprae		32.16	0.18		1	
Staphylococcus aureus subsp. aureus		8.38	0.01		99	
607560	**Good identification**					**607560**
Staphylococcus piscifermentans		100.0	0.48	XYL 1		
607562	**Good identification**					**607562**
Staphylococcus piscifermentans		100.0	0.52	XYL 1		
607570	**Good identification**					**607570**
Staphylococcus piscifermentans		100.0	0.41	XYL 1		
607572	**Good identification**					**607572**
Staphylococcus piscifermentans		100.0	0.44	XYL 1		
607600	**Very good identification**					**607600**
Staphylococcus piscifermentans		100.0	0.89			
607601	**Low discrimination**					**607601**
					YEP	
Staphylococcus piscifermentans		80.12	0.48	MNS 1	1	
Staphylococcus chromogenes		16.47	0.20		95	
Staphylococcus hyicus		3.41	0.02		1	
607602	**Very good identification**					**607602**
					YEP	
Staphylococcus piscifermentans		99.97	0.93		1	
Staphylococcus simulans		0.01	0.03		1	
Staphylococcus chromogenes		0.01	-.02		95	
607603	**Low discrimination**					**607603**
					YEP	
Staphylococcus chromogenes		44.50	0.39	ESL 1	95	
Staphylococcus piscifermentans		36.08	0.52	GAL 90	1	
Staphylococcus hyicus		11.77	0.24		1	
607610	**Very good identification**					**607610**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.82			
607611	**Low discrimination**					**607611**
					YEP	
Staphylococcus piscifermentans		82.75	0.41	MNS 1	1	
Staphylococcus chromogenes		11.19	0.08		95	
Staphylococcus felis		3.22	-.09		1	
607612	**Very good identification**					**607612**
Staphylococcus piscifermentans		99.84	0.85			
Staphylococcus simulans		0.13	0.15			
Staphylococcus felis		0.03	-.09			
607613	**Low discrimination**					**607613**
					YEP	
Staphylococcus felis		53.20	0.33	ESL 1	1	
Staphylococcus piscifermentans		20.74	0.44		1	
Staphylococcus chromogenes		16.83	0.28		95	
607620	**Good identification**					**607620**
Staphylococcus piscifermentans		100.0	0.48	XYL 1		
607622	**Good identification**					**607622**
Staphylococcus piscifermentans		100.0	0.52	XYL 1		
607630	**Good identification**					**607630**
Staphylococcus piscifermentans		100.0	0.41	XYL 1		
607632	**Good identification**					**607632**
Staphylococcus piscifermentans		100.0	0.44	XYL 1		
607640	**Very good identification**					**607640**
Staphylococcus piscifermentans		99.98	0.89			
Staphylococcus hominis subsp. hominis		0.02	0.07			
607641	**Low discrimination**					**607641**
					YEP	
Staphylococcus piscifermentans		82.95	0.48	MNS 1	1	
Staphylococcus chromogenes		17.05	0.20		95	
607642	**Very good identification**					**607642**
					YEP	
Staphylococcus piscifermentans		99.97	0.93		1	
Staphylococcus hominis subsp. hominis		0.02	0.09		49	
Staphylococcus chromogenes		0.01	-.02		95	
607643	**Low discrimination**					**607643**
					YEP	
Staphylococcus chromogenes		54.70	0.39	ESL 1	95	
Staphylococcus piscifermentans		44.35	0.52	GAL 90	1	
Staphylococcus simulans		0.95	0.06		1	
607650	**Very good identification**					**607650**
Staphylococcus piscifermentans		100.0	0.82			
607651	**Low discrimination**					**607651**
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Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		79.08	0.41	MNS 1	1	1	1	
Staphylococcus chromogenes		10.70	0.08		1	95	1	
Staphylococcus sciuri		10.23	0.12		99	57	99	
607652	**Very good identification**							**607652**
Staphylococcus piscifermentans		99.94	0.85					
Staphylococcus simulans		0.06	0.08					
607653	**Low discrimination**							**607653**
					COA	YEP		
Staphylococcus piscifermentans		33.02	0.44	MNS 1	1	1		
Staphylococcus chromogenes		26.80	0.28		1	95		
Staphylococcus delphini		22.21	0.18		99	1		
607660	**Good identification**							**607660**
Staphylococcus piscifermentans		100.0	0.48	XYL 1				
607662	**Good identification**							**607662**
Staphylococcus piscifermentans		100.0	0.52	XYL 1				
607670	**Good identification**							**607670**
Staphylococcus piscifermentans		100.0	0.41	XYL 1				
607672	**Good identification**							**607672**
Staphylococcus piscifermentans		100.0	0.44	XYL 1				
607700	**Very good identification**							**607700**
					YEP			
Staphylococcus piscifermentans		99.99	0.93		1			
Staphylococcus chromogenes		0.01	-.02		95			
607701	**Low discrimination**							**607701**
					YEP			
Staphylococcus chromogenes		48.19	0.39	ESL 1	95			
Staphylococcus piscifermentans		39.07	0.52	LAC 90	1			
Staphylococcus hyicus		12.75	0.24		1			
607702	**Excellent identification**							**607702**
					YEP			
Staphylococcus piscifermentans		99.90	0.96		1			
Staphylococcus chromogenes		0.08	0.18		95			
Staphylococcus hyicus		0.03	0.04		1			
607703	**Low discrimination**							**607703**
					YEP			
Staphylococcus chromogenes		65.56	0.59	ESL 1	95			
Staphylococcus hyicus		22.16	0.45		1			
Staphylococcus piscifermentans		8.86	0.55		1			
607710	**Very good identification**							**607710**
Staphylococcus piscifermentans		100.0	0.85					
607711	**Low discrimination**							**607711**
					YEP			
Staphylococcus piscifermentans		51.47	0.44	MNS 1	1			
Staphylococcus chromogenes		41.78	0.28		95			
Staphylococcus felis		4.47	0.02		1			
607712	**Very good identification**							**607712**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					YEP			
Staphylococcus piscifermentans		99.89	0.89		1			
Staphylococcus felis		0.06	0.02		1			
Staphylococcus chromogenes		0.05	0.07		95			
607713	**Low discrimination**							**607713**
					YEP			
Staphylococcus felis		46.10	0.43	ESL 1	1			
Staphylococcus chromogenes		39.22	0.48		95			
Staphylococcus piscifermentans		8.05	0.48		1			
607720	**Good identification**							**607720**
Staphylococcus piscifermentans		100.0	0.52	XYL 1				
607722	**Very good identification**							**607722**
Staphylococcus piscifermentans		100.0	0.55	XYL 1				
607730	**Good identification**							**607730**
Staphylococcus piscifermentans		100.0	0.44	XYL 1				
607732	**Good identification**							**607732**
Staphylococcus piscifermentans		100.0	0.48	XYL 1				
607740	**Very good identification**							**607740**
					YEP			
Staphylococcus piscifermentans		99.97	0.93		1			
Staphylococcus hominis subsp. hominis		0.02	0.08		49			
Staphylococcus chromogenes		0.01	-.02		95			
607741	**Low discrimination**							**607741**
					YEP			
Staphylococcus chromogenes		55.23	0.39	ESL 1	95			
Staphylococcus piscifermentans		44.78	0.52	LAC 90	1			
607742	**Excellent identification**							**607742**
					YEP			
Staphylococcus piscifermentans		99.91	0.96		1			
Staphylococcus chromogenes		0.08	0.18		95			
Staphylococcus hominis subsp. hominis		0.01	0.09		49			
607743	**Low discrimination**							**607743**
					COA	YEP		
Staphylococcus chromogenes		87.17	0.59	ESL 1	1	95		
Staphylococcus piscifermentans		11.78	0.55		1	1		
Staphylococcus aureus subsp. aureus		0.53	0.01		99	84		
607750	**Very good identification**							**607750**
					NOV			
Staphylococcus piscifermentans		99.97	0.85		1			
Staphylococcus saprophyticus subsp. bovis		0.03	0.02		99			
607751	**Low discrimination**							**607751**
					OXI	YEP	NOV	
Staphylococcus piscifermentans		44.85	0.44	MNS 1	1	1	1	
Staphylococcus chromogenes		36.40	0.28		1	95	1	
Staphylococcus sciuri		7.51	0.18		99	57	99	
607752	**Very good identification**							**607752**
					COA	YEP		
Staphylococcus piscifermentans		99.82	0.89		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. aureus		0.11	0.01		99	84	
Staphylococcus chromogenes		0.05	0.07		1	95	
607753	**Low discrimination**						**607753**
					COA	YEP	
Staphylococcus aureus subsp. aureus		60.41	0.42	VPT 87	99	84	
Staphylococcus chromogenes		28.32	0.48	ESL 1	1	95	
Staphylococcus piscifermentans		5.82	0.48		1	1	
607760	**Good identification**						**607760**
Staphylococcus piscifermentans		100.0	0.52	XYL 1			
607762	**Very good identification**						**607762**
Staphylococcus piscifermentans		100.0	0.55	XYL 1			
607763	**Low discrimination**						**607763**
					NOV	YEP	
Staphylococcus gallinarum		82.61	0.32	ARG 1	99	78	
Staphylococcus chromogenes		14.10	0.18	MAN 91	1	95	
Staphylococcus piscifermentans		1.91	0.14		1	1	
607770	**Acceptable identification**						**607770**
					NOV		
Staphylococcus piscifermentans		93.99	0.44	XYL 1	1		
Staphylococcus gallinarum		6.01	0.04		99		
607771	**Low discrimination**						**607771**
					OXI		
Staphylococcus gallinarum		84.95	0.45	ARG 1	1		
Staphylococcus equorum		13.36	0.31		1		
Staphylococcus sciuri		1.69	0.15		99		
607772	**Low discrimination**						**607772**
					NOV		
Staphylococcus piscifermentans		89.16	0.48	XYL 1	1		
Staphylococcus gallinarum		9.31	0.12		99		
Staphylococcus equorum		1.54	-.02		99		
607773	**Low discrimination**						**607773**
					COA	NOV	
Staphylococcus gallinarum		84.47	0.53	ARG 1	1	99	
Staphylococcus equorum		13.95	0.39		1	99	
Staphylococcus aureus subsp. aureus		0.86	0.01		99	4	
610640	**Good identification**						**610640**
Staphylococcus warneri		100.0	0.45	ORN 1			
610642	**Good identification**						**610642**
Staphylococcus warneri		100.0	0.35	ORN 1			
610650	**Good identification**						**610650**
					NOV		
Staphylococcus warneri		96.69	0.42	ORN 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		3.31	-.09		99		
610652	**Low discrimination**						**610652**
					NOV		
Staphylococcus warneri		63.20	0.32	ORN 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		36.81	0.07		99		
610740	**Good identification**						**610740**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		100.0	0.35	ORN 1		
610750	**Good identification**					**610750**
Staphylococcus warneri		100.0	0.32	ORN 1		
612240	**Low discrimination**					**612240**
Staphylococcus hominis subsp. hominis		81.86	0.28	ARG 12		
Staphylococcus epidermidis		11.46	0.07	ORN 1		
Staphylococcus warneri		6.69	0.11			
612242	**Low discrimination**					**612242**
Staphylococcus hominis subsp. hominis		72.60	0.29	ARG 12		
Staphylococcus epidermidis		27.41	0.17	ORN 1		
612253	**Good identification**					**612253**
					COA	
Staphylococcus capitis subsp. ureolyticus		95.44	0.44	ORN 1	1	
Staphylococcus delphini		4.36	0.18		99	
Staphylococcus simulans		0.21	0.02		1	
612340	**Low discrimination**					**612340**
					NOV	
Staphylococcus hominis subsp. hominis		54.09	0.29	ARG 12	1	
Staphylococcus epidermidis		39.59	0.22	ORN 1	1	
Staphylococcus hominis subsp. novobiosepticus		6.32	-.02		99	
612341	**Good identification**					**612341**
Staphylococcus epidermidis		100.0	0.28	VPT 96		
612342	**Low discrimination**					**612342**
					NOV	
Staphylococcus epidermidis		65.21	0.33	VPT 96	1	
Staphylococcus hominis subsp. hominis		33.02	0.30	ORN 12	1	
Staphylococcus hominis subsp. novobiosepticus		1.77	-.07		99	
612343	**Good identification**					**612343**
					YEP	
Staphylococcus epidermidis		98.67	0.38	VPT 96	1	
Staphylococcus chromogenes		1.33	-.04	ORN 12	95	
612402	**Good identification**					**612402**
Staphylococcus piscifermentans		100.0	0.32	ORN 1		
612440	**Low discrimination**					**612440**
Staphylococcus piscifermentans		48.24	0.28	ORN 1		
Staphylococcus auricularis		43.53	0.18	ESL 90		
Staphylococcus warneri		8.23	0.11			
612442	**Good identification**					**612442**
Staphylococcus piscifermentans		100.0	0.32	ORN 1		
612500	**Good identification**					**612500**
Staphylococcus piscifermentans		100.0	0.32	ORN 1		
612502	**Good identification**					**612502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.36	ORN 1		
612512	**Good identification**					**612512**
Staphylococcus piscifermentans		100.0	0.28	ORN 1		
612540	**Good identification**					**612540**
Staphylococcus piscifermentans		100.0	0.32	ORN 1		
612542	**Good identification**					**612542**
Staphylococcus piscifermentans		100.0	0.36	ORN 1		
612552	**Good identification**					**612552**
Staphylococcus piscifermentans		100.0	0.28	ORN 1		
612600	**Low discrimination**					**612600**
Staphylococcus piscifermentans		73.98	0.32	ORN 1		
Staphylococcus auricularis		17.61	0.10	ESL 90		
Staphylococcus warneri		8.41	0.11			
612602	**Low discrimination**					**612602**
Staphylococcus piscifermentans		79.59	0.36	ORN 1		
Staphylococcus simulans		20.41	0.13	ESL 90		
612613	**Good identification**					**612613**
Staphylococcus simulans		98.10	0.36	ORN 1		
Staphylococcus felis		1.90	-.09	BGA 90		
612640	**Low discrimination**					**612640**
Staphylococcus hominis subsp. hominis		53.35	0.38	ARG 12		
Staphylococcus warneri		26.15	0.38	ORN 1		
Staphylococcus piscifermentans		12.11	0.32			
612642	**Low discrimination**					**612642**
Staphylococcus hominis subsp. hominis		68.37	0.40	ARG 12		
Staphylococcus piscifermentans		19.83	0.36	ORN 1		
Staphylococcus warneri		9.52	0.28			
612643	**Low discrimination**					**612643**
Staphylococcus lugdunensis		79.61	0.33	VPT 99		
Staphylococcus petrasii subsp. petrasii		9.31	0.12	ARG 5		
Staphylococcus simulans		8.06	0.16			
612650	**Low discrimination**					**612650**
Staphylococcus warneri		52.66	0.35	ORN 1		
Staphylococcus hominis subsp. hominis		22.17	0.21			
Staphylococcus piscifermentans		14.43	0.25			
612653	**Low discrimination**					**612653**
Staphylococcus simulans		80.13	0.29	ORN 1		
Staphylococcus capitis subsp. ureolyticus		19.87	0.03	BGA 90		
612700	**Good identification**					**612700**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.36	ORN 1		
612702	**Good identification**					**612702**
Staphylococcus piscifermentans		100.0	0.39	ORN 1		
612710	**Good identification**					**612710**
Staphylococcus piscifermentans		100.0	0.28	ORN 1		
612712	**Low discrimination**					**612712**
Staphylococcus piscifermentans		87.81	0.32	ORN 1		
Staphylococcus simulans		12.19	0.03	ESL 90		
612740	**Low discrimination**					**612740**
Staphylococcus hominis subsp. hominis		66.90	0.39	ARG 12		
Staphylococcus piscifermentans		21.03	0.36	ORN 1		
Staphylococcus warneri		9.56	0.27			
612742	**Low discrimination**					**612742**
Staphylococcus hominis subsp. hominis		69.34	0.40	ARG 12		
Staphylococcus piscifermentans		27.85	0.39	ORN 1		
Staphylococcus warneri		2.81	0.17			
612743	**Low discrimination**					**612743**
					YEP	
Staphylococcus lugdunensis		78.47	0.33	VPT 99	50	
Staphylococcus chromogenes		13.85	0.18	ARG 5	95	
Staphylococcus petrasii subsp. petrasii		6.11	0.09		1	
612752	**Low discrimination**					**612752**
Staphylococcus piscifermentans		46.30	0.32	ORN 1		
Staphylococcus hominis subsp. hominis		40.20	0.23	ESL 90		
Staphylococcus warneri		7.90	0.15			
613400	**Good identification**					**613400**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
613402	**Good identification**					**613402**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
613410	**Good identification**					**613410**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		
613412	**Good identification**					**613412**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
613440	**Good identification**					**613440**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
613442	**Good identification**					**613442**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
613450	**Good identification**					**613450**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
613452	**Good identification**					**613452**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
613500	**Good identification**					**613500**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
613502	**Very good identification**					**613502**
Staphylococcus piscifermentans		100.0	0.55	ORN 1		
613510	**Good identification**					**613510**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
613512	**Good identification**					**613512**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
613540	**Good identification**					**613540**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
613542	**Very good identification**					**613542**
Staphylococcus piscifermentans		100.0	0.55	ORN 1		
613550	**Good identification**					**613550**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
613552	**Good identification**					**613552**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
613600	**Good identification**					**613600**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
613602	**Very good identification**					**613602**
Staphylococcus piscifermentans		100.0	0.55	ORN 1		
613610	**Good identification**					**613610**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
613612	**Good identification**					**613612**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
613640	**Good identification**					**613640**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
613642	**Very good identification**					**613642**
Staphylococcus piscifermentans		100.0	0.55	ORN 1		
613650	**Good identification**					**613650**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
613652	**Good identification**					**613652**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
613700	**Very good identification**						**613700**
Staphylococcus piscifermentans		100.0	0.55	ORN 1			
613702	**Very good identification**						**613702**
Staphylococcus piscifermentans		100.0	0.59	ORN 1			
613710	**Good identification**						**613710**
Staphylococcus piscifermentans		100.0	0.48	ORN 1			
613712	**Good identification**						**613712**
Staphylococcus piscifermentans		100.0	0.51	ORN 1			
613740	**Very good identification**						**613740**
Staphylococcus piscifermentans		100.0	0.55	ORN 1			
613742	**Very good identification**						**613742**
Staphylococcus piscifermentans		100.0	0.59	ORN 1			
613750	**Good identification**						**613750**
Staphylococcus piscifermentans		100.0	0.48	ORN 1			
613752	**Good identification**						**613752**
Staphylococcus piscifermentans		100.0	0.51	ORN 1			
616241	**Good identification**						**616241**
					YEP		
Staphylococcus epidermidis		95.03	0.28	VPT 96	1		
Staphylococcus chromogenes		4.97	-.02	ORN 12	95		
616242	**Good identification**						**616242**
Staphylococcus epidermidis		100.0	0.33	VPT 96			
616243	**Low discrimination**						**616243**
					COA	YEP	
Staphylococcus epidermidis		64.57	0.38	VPT 96	1	1	
Staphylococcus delphini		25.82	0.18	ORN 12	99	1	
Staphylococcus chromogenes		9.61	0.17		1	95	
616253	**Very good identification**						**616253**
					COA	YEP	
Staphylococcus delphini		99.67	0.59	ORN 1	99	1	
Staphylococcus capitis subsp. ureolyticus		0.22	0.03		1	77	
Staphylococcus chromogenes		0.11	0.06		1	95	
616303	**Low discrimination**						**616303**
					YEP		
Staphylococcus chromogenes		71.80	0.37	ORN 1	95		
Staphylococcus hyicus		24.27	0.24	TRE 92	1		
Staphylococcus epidermidis		3.07	0.13		1		
616340	**Good identification**						**616340**
Staphylococcus epidermidis		100.0	0.38	VPT 96			
616341	**Acceptable identification**						**616341**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		92.33	0.44	VPT 96	1		
Staphylococcus chromogenes		7.68	0.17	ORN 12	95		
616342	**Good identification**						**616342**
					YEP		
Staphylococcus epidermidis		99.56	0.48	VPT 96	1		
Staphylococcus chromogenes		0.44	-.04	ORN 12	95		
616343	**Low discrimination**						**616343**
					COA	YEP	
Staphylococcus epidermidis		76.52	0.54	VPT 96	1	1	
Staphylococcus chromogenes		18.08	0.37	ORN 12	1	95	
Staphylococcus delphini		5.40	0.18		99	1	
616353	**Very good identification**						**616353**
					COA	YEP	
Staphylococcus delphini		98.65	0.59	ORN 1	99	1	
Staphylococcus chromogenes		0.94	0.26		1	95	
Staphylococcus aureus subsp. aureus		0.23	0.01		99	84	
616402	**Good identification**						**616402**
Staphylococcus piscifermentans		100.0	0.28	ORN 1			
616442	**Good identification**						**616442**
Staphylococcus piscifermentans		100.0	0.28	ORN 1			
616500	**Good identification**						**616500**
Staphylococcus piscifermentans		100.0	0.28	ORN 1			
616502	**Good identification**						**616502**
Staphylococcus piscifermentans		100.0	0.32	ORN 1			
616513	**Low discrimination**						**616513**
					YEP		
Staphylococcus felis		87.36	0.37	ORN 1	1		
Staphylococcus condimenti		11.11	0.14		1		
Staphylococcus chromogenes		1.52	0.07		95		
616540	**Good identification**						**616540**
Staphylococcus piscifermentans		100.0	0.28	ORN 1			
616542	**Good identification**						**616542**
Staphylococcus piscifermentans		100.0	0.32	ORN 1			
616600	**Good identification**						**616600**
Staphylococcus piscifermentans		100.0	0.28	ORN 1			
616602	**Low discrimination**						**616602**
					YEP		
Staphylococcus piscifermentans		80.62	0.32	ORN 1	1		
Staphylococcus simulans		10.34	0.03	ESL 90	1		
Staphylococcus chromogenes		9.04	-.02		95		
616603	**Low discrimination**						**616603**
					YEP		
Staphylococcus chromogenes		69.62	0.39	ORN 1	95		
Staphylococcus hyicus		18.42	0.24	GAL 90	1		
Staphylococcus felis		9.27	0.16		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
616613	**Low discrimination**						**616613**
					YEP		
Staphylococcus felis		67.11	0.33	ORN 1	1		
Staphylococcus chromogenes		21.24	0.28		95		
Staphylococcus simulans		11.65	0.26		1		
616640	**Low discrimination**						**616640**
Staphylococcus piscifermentans		83.03	0.28	ORN 1			
Staphylococcus hominis subsp. hominis		16.97	0.07	ESL 90			
616642	**Low discrimination**						**616642**
					YEP		
Staphylococcus piscifermentans		78.61	0.32	ORN 1	1		
Staphylococcus hominis subsp. hominis		12.58	0.09	ESL 90	49		
Staphylococcus chromogenes		8.81	-.02		95		
616643	**Good identification**						**616643**
					YEP		
Staphylococcus chromogenes		98.29	0.39	ORN 1	95		
Staphylococcus simulans		1.71	0.06	GAL 90	1		
616653	**Low discrimination**						**616653**
					COA	YEP	
Staphylococcus chromogenes		40.02	0.28	ORN 1	1	95	
Staphylococcus delphini		33.16	0.18	GAL 90	99	1	
Staphylococcus aureus subsp. aureus		15.68	0.07		99	84	
616700	**Low discrimination**						**616700**
					YEP		
Staphylococcus piscifermentans		89.92	0.32	ORN 1	1		
Staphylococcus chromogenes		10.08	-.02	ESL 90	95		
616701	**Low discrimination**						**616701**
					YEP		
Staphylococcus chromogenes		79.08	0.39	ORN 1	95		
Staphylococcus hyicus		20.92	0.24	LAC 90	1		
616702	**Low discrimination**						**616702**
					YEP		
Staphylococcus piscifermentans		52.63	0.36	ORN 1	1		
Staphylococcus chromogenes		35.41	0.18	ESL 90	95		
Staphylococcus hyicus		11.97	0.04		1		
616703	**Low discrimination**						**616703**
					YEP		
Staphylococcus chromogenes		71.98	0.59	ORN 1	95		
Staphylococcus hyicus		24.33	0.45		1		
Staphylococcus felis		3.57	0.26		1		
616711	**Acceptable identification**						**616711**
					YEP		
Staphylococcus chromogenes		90.35	0.28	ORN 1	95		
Staphylococcus felis		9.66	0.02	LAC 90	1		
616712	**Low discrimination**						**616712**
					YEP		
Staphylococcus piscifermentans		50.94	0.28	ORN 1	1		
Staphylococcus felis		26.51	0.02	ESL 90	1		
Staphylococcus chromogenes		22.55	0.07		95		
616713	**Low discrimination**						**616713**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus felis		50.20	0.43	ORN 1	1			
Staphylococcus chromogenes		42.71	0.48		95			
Staphylococcus condimenti		4.72	0.18		1			
616740	**Low discrimination**							**616740**
					YEP			
Staphylococcus piscifermentans		79.38	0.32	ORN 1	1			
Staphylococcus hominis subsp. hominis		11.72	0.08	ESL 90	49			
Staphylococcus chromogenes		8.90	-.02		95			
616741	**Good identification**							**616741**
					YEP			
Staphylococcus chromogenes		98.95	0.39	ORN 1	95			
Staphylococcus epidermidis		1.05	0.02	LAC 90	1			
616742	**Low discrimination**							**616742**
					YEP			
Staphylococcus piscifermentans		52.04	0.36	ORN 1	1			
Staphylococcus chromogenes		35.01	0.18	ESL 90	95			
Staphylococcus epidermidis		6.94	0.07		1			
616743	**Very good identification**							**616743**
					COA	YEP		
Staphylococcus chromogenes		98.45	0.59	ORN 1	1	95		
Staphylococcus aureus subsp. aureus		0.60	0.01		99	84		
Staphylococcus epidermidis		0.37	0.13		1	1		
616751	**Low discrimination**							**616751**
					COA	PYR	YEP	
Staphylococcus chromogenes		81.29	0.28	ORN 1	1	45	95	
Staphylococcus aureus subsp. aureus		15.76	0.01	LAC 90	99	10	84	
Staphylococcus intermedius		2.95	0.00		99	90	1	
616752	**Low discrimination**							**616752**
					COA	YEP		
Staphylococcus piscifermentans		38.91	0.28	ORN 1	1	1		
Staphylococcus aureus subsp. aureus		36.75	0.01	ESL 90	99	84		
Staphylococcus chromogenes		17.23	0.07		1	95		
616753	**Low discrimination**							**616753**
					COA	YEP		
Staphylococcus aureus subsp. aureus		64.79	0.42	VPT 87	99	84		
Staphylococcus chromogenes		30.38	0.48	ORN 1	1	95		
Staphylococcus delphini		2.80	0.18		99	1		
617400	**Good identification**							**617400**
Staphylococcus piscifermentans		100.0	0.44	ORN 1				
617402	**Good identification**							**617402**
Staphylococcus piscifermentans		100.0	0.48	ORN 1				
617410	**Good identification**							**617410**
Staphylococcus piscifermentans		100.0	0.37	ORN 1				
617412	**Good identification**							**617412**
Staphylococcus piscifermentans		100.0	0.41	ORN 1				
617440	**Good identification**							**617440**
Staphylococcus piscifermentans		100.0	0.44	ORN 1				

Code		ID	T	tests	Differentiation tests	Code
Taxon		score	index	against		
617442	**Good identification**					**617442**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
617450	**Good identification**					**617450**
Staphylococcus piscifermentans		100.0	0.37	ORN 1		
617452	**Good identification**					**617452**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		
617500	**Good identification**					**617500**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
617502	**Good identification**					**617502**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
617510	**Good identification**					**617510**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		
617512	**Good identification**					**617512**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
617540	**Good identification**					**617540**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
617542	**Good identification**					**617542**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
617550	**Good identification**					**617550**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		
617552	**Good identification**					**617552**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
617600	**Good identification**					**617600**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
617602	**Good identification**					**617602**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
617610	**Good identification**					**617610**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		
617612	**Good identification**					**617612**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
617640	**Good identification**					**617640**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
617642	**Good identification**					**617642**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
617650	**Good identification**					**617650**
Staphylococcus piscifermentans		100.0	0.41	ORN 1		
617652	**Good identification**					**617652**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
617700	**Good identification**					**617700**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
617702	**Very good identification**					**617702**
Staphylococcus piscifermentans		100.0	0.55	ORN 1		
617710	**Good identification**					**617710**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
617712	**Good identification**					**617712**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
617740	**Good identification**					**617740**
Staphylococcus piscifermentans		100.0	0.52	ORN 1		
617742	**Very good identification**					**617742**
Staphylococcus piscifermentans		100.0	0.55	ORN 1		
617750	**Good identification**					**617750**
Staphylococcus piscifermentans		100.0	0.44	ORN 1		
617752	**Good identification**					**617752**
Staphylococcus piscifermentans		100.0	0.48	ORN 1		
620000	**Low discrimination**					**620000**
					OXI	
Kocuria varians		53.65	0.31	ARG 1	13	
Dermacoccus nishinomiyaensis		28.19	0.18		93	
Micrococcus sp.		15.12	0.03		89	
620100	**Acceptable identification**					**620100**
					OXI	
Kocuria varians		93.74	0.35	ARG 1	13	
Dermacoccus nishinomiyaensis		6.26	0.03		93	
620203	**Good identification**					**620203**
Staphylococcus simulans		100.0	0.29	NIT 96		
620212	**Good identification**					**620212**
Staphylococcus simulans		97.30	0.31	NIT 96		
Staphylococcus warneri		2.70	0.08	TRE 95		
620213	**Low discrimination**					**620213**
Staphylococcus simulans		85.68	0.42	NIT 96		
Staphylococcus capitis subsp. ureolyticus		14.32	0.13	TRE 95		
620240	**Good identification**					**620240**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		98.70	0.48	TRE 95			
Staphylococcus auricularis		1.30	0.01				
620242	**Low discrimination**						**620242**
					NOV		
Staphylococcus warneri		81.25	0.38	TRE 95	1		
Staphylococcus simulans		12.54	0.11		1		
Staphylococcus saprophyticus subsp. saprophyticus		6.21	-.06		99		
620250	**Low discrimination**						**620250**
					NOV		
Staphylococcus warneri		87.03	0.45	TRE 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		12.97	0.07		99		
620253	**Low discrimination**						**620253**
Staphylococcus capitis subsp. ureolyticus		89.93	0.41	BGA 12			
Staphylococcus simulans		10.07	0.35	NIT 95			
620340	**Good identification**						**620340**
Staphylococcus warneri		95.12	0.37	TRE 95			
Kocuria varians		4.88	0.04				
620342	**Low discrimination**						**620342**
Staphylococcus warneri		77.72	0.27	TRE 95			
Staphylococcus epidermidis		22.28	0.07				
620350	**Good identification**						**620350**
Staphylococcus warneri		100.0	0.34	TRE 95			
620403	**Good identification**						**620403**
Staphylococcus simulans		100.0	0.27	NIT 96			
620412	**Low discrimination**						**620412**
					NOV		
Staphylococcus simulans		83.21	0.29	NIT 96	1		
Staphylococcus saprophyticus subsp. saprophyticus		13.88	0.02	SUC 96	99		
Staphylococcus warneri		2.92	0.08		1		
620413	**Good identification**						**620413**
Staphylococcus simulans		96.62	0.40	NIT 96			
Staphylococcus felis		3.38	0.01	SUC 96			
620440	**Low discrimination**						**620440**
					NOV		
Staphylococcus warneri		82.77	0.48	SUC 95	1		
Staphylococcus auricularis		13.58	0.23		1		
Staphylococcus saprophyticus subsp. saprophyticus		3.64	-.01		99		
620442	**Low discrimination**						**620442**
					NOV		
Staphylococcus warneri		51.94	0.38	SUC 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		38.88	0.14		99		
Staphylococcus simulans		6.35	0.09		1		
620450	**Low discrimination**						**620450**
					OXI	NOV	
Staphylococcus saprophyticus subsp. saprophyticus		56.69	0.27	ARG 8	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		38.85	0.45	SUC 91	1	1	
Macrococcus equipersicus		2.74	0.11		99	99	
620452	**Acceptable identification**						**620452**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		93.68	0.43	ARG 8	99		
Staphylococcus warneri		3.78	0.35	SUC 91	1		
Staphylococcus simulans		2.55	0.22		1		
620453	**Low discrimination**						**620453**
					NOV		
Staphylococcus simulans		84.67	0.32	NIT 96	1		
Staphylococcus saprophyticus subsp. saprophyticus		9.39	0.02	SUC 96	99		
Staphylococcus cohnii subsp. urealyticum		5.94	0.14		99		
620540	**Acceptable identification**						**620540**
Staphylococcus warneri		94.54	0.37	SUC 95			
Staphylococcus auricularis		5.46	0.03				
620542	**Low discrimination**						**620542**
					NOV		
Staphylococcus warneri		82.91	0.27	SUC 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		17.09	-.08		99		
620550	**Low discrimination**						**620550**
					NOV	PYR	
Staphylococcus warneri		59.73	0.34	SUC 95	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		24.00	0.05		99	7	
Staphylococcus saprophyticus subsp. bovis		6.97	0.05		99	99	
620600	**Acceptable identification**						**620600**
Staphylococcus warneri		92.01	0.48	MLT 95			
Staphylococcus auricularis		5.97	0.15				
Staphylococcus simulans		2.02	0.04				
620602	**Low discrimination**						**620602**
					NOV		
Staphylococcus simulans		85.84	0.45	NIT 96	1		
Staphylococcus warneri		13.15	0.38		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.01	-.06		99		
620603	**Very good identification**						**620603**
Staphylococcus simulans		99.82	0.56	NIT 96			
Staphylococcus felis		0.18	-.10				
620610	**Low discrimination**						**620610**
					NOV		
Staphylococcus warneri		78.71	0.45	MLT 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		11.73	0.07		99		
Staphylococcus simulans		9.56	0.16		1		
620611	**Good identification**						**620611**
Staphylococcus simulans		95.22	0.27	NIT 96			
Staphylococcus warneri		4.78	0.10	LAC 99			
620612	**Good identification**						**620612**
					NOV		
Staphylococcus simulans		91.07	0.57	NIT 96	1		
Staphylococcus saprophyticus subsp. saprophyticus		6.40	0.22		99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		2.53	0.35		1		
620613	**Very good identification**						**620613**
					COA		
Staphylococcus simulans		99.65	0.68	NIT 96	1		
Staphylococcus felis		0.30	0.07		1		
Staphylococcus intermedius		0.06	-.01		99		
620633	**Good identification**						**620633**
Staphylococcus simulans		100.0	0.27	NIT 96			
620640	**Very good identification**						**620640**
					NOV		
Staphylococcus warneri		96.67	0.74		1		
Staphylococcus saprophyticus subsp. saprophyticus		2.27	0.19		99		
Staphylococcus auricularis		0.77	0.22		1		
620641	**Low discrimination**						**620641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		50.18	0.27	VPT 99	1	99	
Staphylococcus warneri		37.43	0.39	NIT 90	1	8	
Kocuria kristinae		7.57	0.10		99	93	
620642	**Low discrimination**						**620642**
					NOV		
Staphylococcus warneri		64.09	0.64		1		
Staphylococcus saprophyticus subsp. saprophyticus		25.53	0.35		99		
Staphylococcus simulans		9.89	0.38		1		
620643	**Low discrimination**						**620643**
					PYR		
Staphylococcus simulans		87.81	0.48	NIT 96	70		
Staphylococcus petrasii subsp. petrasii		7.51	0.21		99		
Staphylococcus warneri		3.47	0.29		8		
620650	**Low discrimination**						**620650**
					NOV		
Staphylococcus warneri		56.14	0.71		1		
Staphylococcus saprophyticus subsp. saprophyticus		43.59	0.48		99		
Staphylococcus simulans		0.16	0.09		1		
620651	**Low discrimination**						**620651**
					NOV		
Staphylococcus warneri		48.52	0.36	MNS 2	1		
Staphylococcus simulans		22.85	0.20		1		
Staphylococcus saprophyticus subsp. saprophyticus		18.65	0.07		99		
620652	**Low discrimination**						**620652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		87.69	0.63	ARG 8	99		
Staphylococcus warneri		6.64	0.61		1		
Staphylococcus simulans		5.66	0.50		1		
620653	**Good identification**						**620653**
					NOV		
Staphylococcus simulans		94.09	0.61	NIT 96	1		
Staphylococcus saprophyticus subsp. saprophyticus		4.40	0.22		99		
Staphylococcus warneri		0.67	0.26		1		
620660	**Good identification**						**620660**
Staphylococcus warneri		100.0	0.33	XYL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
620670	**Low discrimination**						**620670**
					NOV		
Staphylococcus warneri		56.29	0.30	XYL 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		43.71	0.07		99		
620700	**Good identification**						**620700**
Staphylococcus warneri		100.0	0.37	MLT 95			
620703	**Low discrimination**						**620703**
					COA	YEP	
Staphylococcus simulans		45.55	0.34	NIT 96	1	1	
Staphylococcus intermedius		43.03	0.32	GAL 8	99	1	
Staphylococcus chromogenes		4.74	0.10		1	95	
620710	**Good identification**						**620710**
Staphylococcus warneri		100.0	0.34	MLT 95			
620712	**Low discrimination**						**620712**
					NOV		
Staphylococcus simulans		83.85	0.35	NIT 96	1		
Staphylococcus warneri		8.45	0.24	GAL 8	1		
Staphylococcus saprophyticus subsp. saprophyticus		5.89	0.00		99		
620713	**Low discrimination**						**620713**
					COA		
Staphylococcus simulans		51.36	0.46	NIT 96	1		
Staphylococcus intermedius		32.79	0.40	GAL 8	99		
Staphylococcus condimenti		9.61	0.18		1		
620740	**Very good identification**						**620740**
					NOV	PYR	
Staphylococcus warneri		98.46	0.64		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		0.64	-.02		99	7	
Staphylococcus petrasii subsp. petrasii		0.63	0.04		1	99	
620742	**Low discrimination**						**620742**
					NOV		
Staphylococcus warneri		86.02	0.54		1		
Staphylococcus saprophyticus subsp. saprophyticus		9.44	0.13		99		
Staphylococcus simulans		3.66	0.16		1		
620743	**Low discrimination**						**620743**
					COA		
Staphylococcus intermedius		60.83	0.35	NIT 93	99		
Staphylococcus simulans		19.29	0.27	PHS 97	1		
Staphylococcus petrasii subsp. petrasii		12.65	0.18		1		
620750	**Low discrimination**						**620750**
					NOV	PYR	
Staphylococcus warneri		80.75	0.61		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		17.27	0.26		99	7	
Staphylococcus saprophyticus subsp. bovis		1.98	0.18		99	99	
620751	**Low discrimination**						**620751**
					COA	PYR	
Staphylococcus intermedius		71.97	0.28	NIT 93	99	90	
Staphylococcus warneri		16.16	0.26	PHS 97	1	8	
Staphylococcus petrasii subsp. petrasii		7.02	0.04	LAC 86	1	99	
620752	**Low discrimination**						**620752**
					NOV		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus saprophyticus subsp. saprophyticus		74.11	0.42	ARG 8	99		
Staphylococcus warneri		20.39	0.51	GAL 8	1		
Staphylococcus simulans		4.79	0.28		1		
620753	**Low discrimination**						**620753**
					COA	NOV	
Staphylococcus intermedius		65.81	0.44	NIT 93	99	1	
Staphylococcus simulans		30.88	0.39	PHS 97	1	1	
Staphylococcus saprophyticus subsp. saprophyticus		1.44	0.00		1	99	
621613	**Good identification**						**621613**
Staphylococcus simulans		100.0	0.27	ESL 1			
621640	**Acceptable identification**						**621640**
Staphylococcus warneri		93.27	0.33	ESL 1			
Staphylococcus piscifermentans		6.73	0.11				
621650	**Low discrimination**						**621650**
					NOV		
Staphylococcus warneri		56.29	0.30	ESL 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		43.71	0.07		99		
622000	**Low discrimination**						**622000**
					OXI		
Kocuria varians		86.75	0.46	ARG 1	13		
Staphylococcus carnosus subsp. utilis		5.29	0.03		1		
Dermacoccus nishinomiyaensis		5.25	0.13		93		
622002	**Low discrimination**						**622002**
Kocuria varians		81.26	0.34	ARG 1			
Staphylococcus simulans		18.74	0.18				
622003	**Low discrimination**						**622003**
Staphylococcus simulans		85.02	0.29	SUC 96			
Kocuria varians		9.58	0.12	TRE 95			
Staphylococcus carnosus subsp. carnosus		5.41	0.02				
622012	**Good identification**						**622012**
Staphylococcus simulans		100.0	0.31	SUC 96			
622013	**Low discrimination**						**622013**
Staphylococcus simulans		81.73	0.42	SUC 96			
Staphylococcus capitis subsp. ureolyticus		10.81	0.11	TRE 95			
Staphylococcus carnosus subsp. carnosus		5.20	0.14				
622020	**Good identification**						**622020**
Kocuria varians		100.0	0.36	ARG 1			
622040	**Low discrimination**						**622040**
Kocuria varians		71.91	0.32	ARG 1			
Staphylococcus auricularis		23.89	0.21				
Staphylococcus warneri		4.20	0.14				
622053	**Low discrimination**						**622053**
Staphylococcus capitis subsp. ureolyticus		87.60	0.39	BGA 12			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		12.40	0.35	SUC 96		
622100	**Good identification**					**622100**
					OXI	
Kocuria varians		99.24	0.49	ARG 1	13	
Dermacoccus nishinomiyaensis		0.76	-.02		93	
622101	**Acceptable identification**					**622101**
					COA	
Kocuria varians		93.83	0.27	ARG 1	1	
Staphylococcus schleiferi subsp. coagulans		6.17	0.02	MNS 8	99	
622102	**Good identification**					**622102**
Kocuria varians		100.0	0.38	ARG 1		
622120	**Good identification**					**622120**
Kocuria varians		100.0	0.40	ARG 1		
622122	**Good identification**					**622122**
Kocuria varians		100.0	0.28	ARG 1		
622140	**Good identification**					**622140**
Kocuria varians		97.60	0.36	ARG 1		
Staphylococcus auricularis		2.40	0.01			
622200	**Low discrimination**					**622200**
Kocuria varians		70.37	0.28	ARG 1		
Staphylococcus auricularis		14.89	0.13	SUC 12		
Staphylococcus simulans		8.14	0.06			
622202	**Good identification**					**622202**
Staphylococcus simulans		97.60	0.47	TRE 95		
Kocuria varians		2.41	0.16			
622203	**Very good identification**					**622203**
					YEP	
Staphylococcus simulans		96.68	0.58	TRE 95	1	
Staphylococcus capitis subsp. ureolyticus		2.13	0.11		77	
Staphylococcus chromogenes		0.66	0.10		95	
622212	**Very good identification**					**622212**
Staphylococcus simulans		99.56	0.59	TRE 95		
Staphylococcus capitis subsp. ureolyticus		0.45	-.02			
622213	**Low discrimination**					**622213**
Staphylococcus simulans		88.06	0.70	TRE 95		
Staphylococcus capitis subsp. ureolyticus		11.65	0.39			
Staphylococcus felis		0.21	0.07			
622233	**Low discrimination**					**622233**
Staphylococcus simulans		88.32	0.29	TRE 95		
Staphylococcus capitis subsp. ureolyticus		11.68	-.02	XYL 1		
622240	**Low discrimination**					**622240**
Staphylococcus warneri		48.26	0.40	TRE 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		23.87	0.28				
Staphylococcus auricularis		13.35	0.20				
622242	**Low discrimination**						**622242**
Staphylococcus simulans		65.83	0.40	TRE 95			
Staphylococcus hominis subsp. hominis		13.26	0.29				
Staphylococcus epidermidis		9.18	0.23				
622243	**Low discrimination**						**622243**
Staphylococcus capitis subsp. ureolyticus		51.74	0.39	BGA 12			
Staphylococcus simulans		43.93	0.51	MAN 96			
Staphylococcus epidermidis		3.40	0.28				
622250	**Low discrimination**						**622250**
Staphylococcus warneri		52.39	0.37	TRE 95			
Staphylococcus capitis subsp. ureolyticus		27.17	0.09				
Staphylococcus simulans		8.42	0.11				
622251	**Good identification**						**622251**
Staphylococcus capitis subsp. ureolyticus		98.96	0.50	BGA 12			
Staphylococcus simulans		1.04	0.22	LAC 88			
622252	**Low discrimination**						**622252**
Staphylococcus simulans		78.90	0.52	TRE 95			
Staphylococcus capitis subsp. ureolyticus		18.86	0.26				
Staphylococcus warneri		1.65	0.27				
622253	**Low discrimination**						**622253**
					COA		
Staphylococcus capitis subsp. ureolyticus		87.32	0.67	BGA 12	1		
Staphylococcus simulans		12.36	0.63		1		
Staphylococcus delphini		0.30	0.18		99		
622300	**Good identification**						**622300**
Kocuria varians		100.0	0.31	ARG 1			
622303	**Low discrimination**						**622303**
					COA	YEP	
Staphylococcus simulans		32.21	0.36	GAL 8	1	1	
Staphylococcus intermedius		24.08	0.32	TRE 95	99	1	
Staphylococcus chromogenes		22.83	0.30		1	95	
622312	**Good identification**						**622312**
					COA		
Staphylococcus simulans		98.32	0.37	GAL 8	1		
Staphylococcus intermedius		1.68	-.01	TRE 95	99		
622313	**Low discrimination**						**622313**
					COA		
Staphylococcus simulans		55.02	0.48	GAL 8	1		
Staphylococcus intermedius		27.81	0.40	TRE 95	99		
Staphylococcus condimenti		8.15	0.18		1		
622340	**Low discrimination**						**622340**
Staphylococcus epidermidis		37.72	0.28	VPT 96			
Staphylococcus hominis subsp. hominis		28.11	0.29				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		16.57	0.30					
622341	**Low discrimination**							**622341**
					COA	YEP	PYR	
Staphylococcus epidermidis		70.40	0.33	VPT 96	1	1	1	
Staphylococcus intermedius		17.97	0.19		99	1	90	
Staphylococcus chromogenes		7.75	0.10		1	95	45	
622342	**Low discrimination**							**622342**
Staphylococcus epidermidis		67.37	0.38	VPT 96				
Staphylococcus hominis subsp. hominis		18.61	0.30					
Staphylococcus simulans		7.41	0.18					
622343	**Low discrimination**							**622343**
					COA	YEP	PYR	
Staphylococcus epidermidis		48.95	0.44	VPT 96	1	1	1	
Staphylococcus intermedius		24.24	0.35		99	1	90	
Staphylococcus chromogenes		15.32	0.30		1	95	45	
622351	**Low discrimination**							**622351**
					COA	NOV	PYR	
Staphylococcus intermedius		64.23	0.28	PHS 97	99	1	90	
Staphylococcus capitis subsp. ureolyticus		26.18	0.09	TRE 93	1	1	1	
Staphylococcus saprophyticus subsp. bovis		4.34	0.05	LAC 86	1	99	99	
622352	**Low discrimination**							**622352**
Staphylococcus simulans		77.58	0.30	GAL 8				
Staphylococcus hominis subsp. hominis		7.28	0.13	TRE 95				
Staphylococcus warneri		5.90	0.17					
622353	**Low discrimination**							**622353**
					COA	PYR		
Staphylococcus intermedius		43.19	0.44	PHS 97	99	90		
Staphylococcus simulans		25.59	0.41	TRE 93	1	70		
Staphylococcus capitis subsp. ureolyticus		21.01	0.26		1	1		
622400	**Low discrimination**							**622400**
Staphylococcus auricularis		42.91	0.35	URE 1				
Staphylococcus carnosus subsp. utilis		38.20	0.18	BGA 14				
Kocuria varians		8.99	0.22					
622402	**Low discrimination**							**622402**
Staphylococcus simulans		89.33	0.45	SUC 96				
Staphylococcus piscifermentans		6.25	0.32					
Staphylococcus auricularis		2.89	0.16					
622403	**Good identification**							**622403**
Staphylococcus simulans		92.06	0.56	SUC 96				
Staphylococcus felis		7.94	0.25					
622411	**Low discrimination**							**622411**
Staphylococcus simulans		87.38	0.27	SUC 96				
Staphylococcus felis		12.62	0.01	LAC 99				
622412	**Very good identification**							**622412**
Staphylococcus simulans		98.78	0.57	SUC 96				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		0.74	0.25				
Staphylococcus felis		0.48	0.01				
622413	**Low discrimination**						**622413**
Staphylococcus simulans		86.46	0.68	SUC 96			
Staphylococcus felis		12.48	0.42				
Staphylococcus condimenti		0.94	0.14				
622433	**Low discrimination**						**622433**
Staphylococcus simulans		87.38	0.27	SUC 96			
Staphylococcus felis		12.62	0.01	XYL 1			
622440	**Low discrimination**						**622440**
Staphylococcus auricularis		72.53	0.43	URE 1			
Staphylococcus warneri		21.05	0.40	BGA 14			
Staphylococcus piscifermentans		4.99	0.28				
622442	**Low discrimination**						**622442**
Staphylococcus simulans		68.04	0.38	SUC 96			
Staphylococcus auricularis		11.42	0.23				
Staphylococcus piscifermentans		10.60	0.32				
622443	**Good identification**						**622443**
Staphylococcus simulans		98.68	0.48	SUC 96			
Staphylococcus petrasii subsp. petrasii		0.77	0.00				
Staphylococcus auricularis		0.55	0.03				
622450	**Low discrimination**						**622450**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		61.79	0.34	ARG 1	99	99	
Staphylococcus warneri		29.04	0.37		1	8	
Staphylococcus piscifermentans		4.07	0.21		1	15	
622452	**Acceptable identification**						**622452**
					NOV	PYR	
Staphylococcus simulans		90.84	0.50	SUC 96	1	70	
Staphylococcus saprophyticus subsp. bovis		3.84	0.22		99	99	
Staphylococcus warneri		2.40	0.27		1	8	
622453	**Very good identification**						**622453**
Staphylococcus simulans		99.00	0.61	SUC 96			
Staphylococcus capitis subsp. ureolyticus		0.37	-.02				
Staphylococcus felis		0.32	0.01				
622502	**Low discrimination**						**622502**
Staphylococcus piscifermentans		46.16	0.36	BGA 1			
Staphylococcus simulans		38.22	0.23	ESL 90			
Kocuria varians		11.33	0.15				
622503	**Low discrimination**						**622503**
Staphylococcus felis		64.75	0.36	PHS 99			
Staphylococcus simulans		19.49	0.34	MAN 87			
Staphylococcus condimenti		9.73	0.14				
622512	**Low discrimination**						**622512**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
	Staphylococcus simulans	57.63	0.35	GAL 8			
	Staphylococcus condimenti	24.08	0.14	SUC 96			
	Staphylococcus felis	10.83	0.12				
622513	**Low discrimination**						**622513**
	Staphylococcus condimenti	65.00	0.55	PHS 99			
	Staphylococcus felis	29.25	0.53				
	Staphylococcus simulans	5.26	0.46				
622542	**Low discrimination**						**622542**
	Staphylococcus piscifermentans	60.63	0.36	BGA 1			
	Staphylococcus simulans	22.56	0.16	ESL 90			
	Staphylococcus warneri	11.97	0.20				
622543	**Low discrimination**						**622543**
					COA	YEP	
	Staphylococcus simulans	55.90	0.27	GAL 8	1	1	
	Staphylococcus intermedius	23.66	0.17	SUC 96	99	1	
	Staphylococcus chromogenes	12.94	0.10		1	95	
622550	**Acceptable identification**						**622550**
					NOV	PYR	
	Staphylococcus saprophyticus subsp. bovis	94.18	0.47	ARG 1	99	99	
	Staphylococcus warneri	3.50	0.27		1	8	
	Staphylococcus piscifermentans	2.33	0.25		1	15	
622551	**Acceptable identification**						**622551**
					COA	NOV	
	Staphylococcus saprophyticus subsp. bovis	92.57	0.34	ARG 1	1	99	
	Staphylococcus intermedius	7.43	0.10		99	1	
622552	**Low discrimination**						**622552**
					NOV		
	Staphylococcus saprophyticus subsp. bovis	58.39	0.34	ARG 1	99		
	Staphylococcus simulans	30.07	0.28		1		
	Staphylococcus piscifermentans	8.66	0.28		1		
622553	**Low discrimination**						**622553**
					COA		
	Staphylococcus simulans	54.05	0.39	GAL 8	1		
	Staphylococcus intermedius	15.46	0.26	SUC 96	99		
	Staphylococcus condimenti	15.01	0.14		1		
622600	**Low discrimination**						**622600**
	Staphylococcus auricularis	34.15	0.34	URE 1			
	Staphylococcus simulans	30.83	0.32	BGA 14			
	Staphylococcus warneri	25.05	0.40				
622601	**Low discrimination**						**622601**
					YEP		
	Staphylococcus simulans	87.34	0.43	LAC 99	1		
	Staphylococcus chromogenes	3.98	0.12		95		
	Staphylococcus auricularis	3.21	0.15		1		
622602	**Very good identification**						**622602**
	Staphylococcus simulans	99.15	0.73				
	Staphylococcus piscifermentans	0.43	0.36				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		0.27	0.30			
622603	**Very good identification**					**622603**
					YEP	
Staphylococcus simulans		98.57	0.84		1	
Staphylococcus felis		0.72	0.31		1	
Staphylococcus chromogenes		0.41	0.32		95	
622610	**Low discrimination**					**622610**
Staphylococcus simulans		83.99	0.45	LAC 99		
Staphylococcus warneri		12.35	0.37			
Staphylococcus piscifermentans		2.60	0.25			
622611	**Very good identification**					**622611**
					COA	
Staphylococcus simulans		98.00	0.55	LAC 99	1	
Staphylococcus felis		1.20	0.07		1	
Staphylococcus intermedius		0.49	0.06		99	
622612	**Very good identification**					**622612**
Staphylococcus simulans		99.86	0.86			
Staphylococcus warneri		0.05	0.27			
Staphylococcus piscifermentans		0.05	0.28			
622613	**Very good identification**					**622613**
Staphylococcus simulans		98.66	0.96			
Staphylococcus felis		1.21	0.48			
Staphylococcus condimenti		0.07	0.18			
622622	**Good identification**					**622622**
Staphylococcus simulans		100.0	0.32	XYL 1		
622623	**Good identification**					**622623**
Staphylococcus simulans		99.28	0.43	XYL 1		
Staphylococcus felis		0.73	-.10			
622632	**Good identification**					**622632**
Staphylococcus simulans		100.0	0.45	XYL 1		
622633	**Very good identification**					**622633**
Staphylococcus simulans		98.79	0.55	XYL 1		
Staphylococcus felis		1.21	0.07			
622640	**Low discrimination**					**622640**
Staphylococcus warneri		71.40	0.66			
Staphylococcus auricularis		11.95	0.42			
Staphylococcus hominis subsp. hominis		5.89	0.38			
622641	**Low discrimination**					**622641**
					PYR	
Staphylococcus petrasii subsp. petrasii		79.25	0.47	VPT 99	99	
Staphylococcus simulans		13.44	0.36		70	
Staphylococcus warneri		2.82	0.31		8	
622642	**Low discrimination**					**622642**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus simulans		84.35	0.66					
Staphylococcus warneri		9.76	0.56					
Staphylococcus hominis subsp. hominis		2.83	0.40					
622643	**Very good identification**							**622643**
					YEP			
Staphylococcus simulans		95.72	0.77		1			
Staphylococcus petrasii subsp. petrasii		3.07	0.41		1			
Staphylococcus chromogenes		0.88	0.32		95			
622650	**Low discrimination**							**622650**
					NOV	PYR		
Staphylococcus warneri		58.78	0.63		1	8		
Staphylococcus saprophyticus subsp. bovis		26.33	0.47		99	99		
Staphylococcus simulans		9.45	0.37		1	70		
622651	**Low discrimination**							**622651**
					NOV			
Staphylococcus simulans		67.37	0.48	LAC 99	1			
Staphylococcus saprophyticus subsp. bovis		14.02	0.34		99			
Staphylococcus petrasii subsp. petrasii		11.04	0.27		1			
622652	**Very good identification**							**622652**
					NOV	PYR		
Staphylococcus simulans		96.33	0.78		1	70		
Staphylococcus warneri		2.02	0.54		1	8		
Staphylococcus saprophyticus subsp. bovis		0.68	0.34		99	99		
622653	**Very good identification**							**622653**
					COA	PYR		
Staphylococcus simulans		99.27	0.89		1	70		
Staphylococcus capitis subsp. ureolyticus		0.37	0.26		1	1		
Staphylococcus intermedius		0.10	0.26		99	90		
622663	**Good identification**							**622663**
Staphylococcus simulans		96.89	0.36	XYL 1				
Staphylococcus petrasii subsp. petrasii		3.11	0.00					
622672	**Good identification**							**622672**
Staphylococcus simulans		97.95	0.37	XYL 1				
Staphylococcus warneri		2.05	0.12					
622673	**Good identification**							**622673**
					NOV			
Staphylococcus simulans		97.47	0.48	XYL 1	1			
Staphylococcus xylosus		2.53	0.12		99			
622700	**Low discrimination**							**622700**
Staphylococcus piscifermentans		44.39	0.36	BGA 1				
Staphylococcus warneri		26.29	0.30	ESL 90				
Staphylococcus auricularis		12.61	0.15					
622701	**Low discrimination**							**622701**
					YEP	PYR	COA	
Staphylococcus intermedius		48.38	0.38	PHS 97	1	90	99	
Staphylococcus chromogenes		27.09	0.32	LAC 86	95	45	1	
Staphylococcus hyicus		16.00	0.23		1	1	24	
622702	**Low discrimination**							**622702**
					COA			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		87.87	0.51	GAL 8	1		
Staphylococcus piscifermentans		6.63	0.39		1		
Staphylococcus intermedius		1.55	0.14		99		
622703	**Low discrimination**						**622703**
					COA	YEP	
Staphylococcus simulans		38.63	0.62	GAL 8	1	1	
Staphylococcus intermedius		20.20	0.55		99	1	
Staphylococcus chromogenes		16.57	0.51		1	95	
622711	**Low discrimination**						**622711**
					COA		
Staphylococcus intermedius		68.51	0.47	PHS 97	99		
Staphylococcus simulans		12.03	0.34	LAC 86	1		
Staphylococcus condimenti		9.29	0.18		1		
622712	**Good identification**						**622712**
Staphylococcus simulans		93.78	0.64	GAL 8			
Staphylococcus condimenti		2.45	0.18				
Staphylococcus felis		1.49	0.18				
622713	**Low discrimination**						**622713**
Staphylococcus simulans		38.07	0.75	GAL 8			
Staphylococcus condimenti		29.39	0.59				
Staphylococcus felis		17.90	0.59				
622733	**Low discrimination**						**622733**
Staphylococcus simulans		36.66	0.34	GAL 8			
Staphylococcus condimenti		28.30	0.18	XYL 1			
Staphylococcus felis		17.24	0.18				
622740	**Low discrimination**						**622740**
					PYR		
Staphylococcus warneri		57.14	0.56		8		
Staphylococcus hominis subsp. hominis		16.16	0.39		1		
Staphylococcus petrasii subsp. petrasii		7.70	0.23		99		
622741	**Low discrimination**						**622741**
					COA	YEP	
Staphylococcus petrasii subsp. petrasii		52.30	0.43	VPT 99	1	1	
Staphylococcus intermedius		32.54	0.42		99	1	
Staphylococcus chromogenes		12.15	0.32		1	95	
622742	**Low discrimination**						**622742**
Staphylococcus simulans		43.46	0.44	GAL 8			
Staphylococcus warneri		18.26	0.46				
Staphylococcus hominis subsp. hominis		18.18	0.40				
622743	**Low discrimination**						**622743**
					COA	YEP	
Staphylococcus intermedius		44.00	0.58	PHS 97	99	1	
Staphylococcus simulans		25.20	0.55		1	1	
Staphylococcus chromogenes		24.07	0.51		1	95	
622750	**Low discrimination**						**622750**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		81.27	0.59	ARG 1	99	99	
Staphylococcus warneri		14.32	0.53		1	8	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus haemolyticus		1.48	0.26		2	98	
622751	**Low discrimination**						**622751**
					COA	NOV	
Staphylococcus intermedius		59.78	0.51	PHS 97	99	1	
Staphylococcus saprophyticus subsp. bovis		30.08	0.47	LAC 86	1	99	
Staphylococcus petrasii subsp. petrasii		3.95	0.23		1	1	
622752	**Low discrimination**						**622752**
					NOV	PYR	
Staphylococcus simulans		66.76	0.57	GAL 8	1	70	
Staphylococcus saprophyticus subsp. bovis		21.61	0.47		99	99	
Staphylococcus warneri		5.08	0.43		1	8	
622753	**Low discrimination**						**622753**
					COA	YEP	
Staphylococcus intermedius		50.29	0.67	PHS 97	99	1	
Staphylococcus simulans		42.62	0.67		1	1	
Staphylococcus chromogenes		2.87	0.40		1	95	
622773	**Low discrimination**						**622773**
					COA	NOV	
Staphylococcus xylosus		48.53	0.33	ARG 1	1	99	
Staphylococcus intermedius		18.97	0.26	GLR 95	99	1	
Staphylococcus simulans		16.07	0.26		1	1	
623213	**Low discrimination**						**623213**
Staphylococcus simulans		88.32	0.29	ESL 1			
Staphylococcus capitis subsp. ureolyticus		11.68	-.02	TRE 95			
623400	**Good identification**						**623400**
Staphylococcus piscifermentans		100.0	0.48	BGA 1			
623402	**Good identification**						**623402**
Staphylococcus piscifermentans		98.42	0.52	BGA 1			
Staphylococcus simulans		1.58	0.04				
623410	**Good identification**						**623410**
Staphylococcus piscifermentans		100.0	0.41	BGA 1			
623412	**Low discrimination**						**623412**
Staphylococcus piscifermentans		86.99	0.44	BGA 1			
Staphylococcus simulans		13.02	0.16				
623413	**Low discrimination**						**623413**
Staphylococcus simulans		87.38	0.27	ESL 1			
Staphylococcus felis		12.62	0.01	SUC 96			
623440	**Good identification**						**623440**
Staphylococcus piscifermentans		98.39	0.48	BGA 1			
Staphylococcus auricularis		1.61	0.02				
623442	**Good identification**						**623442**
Staphylococcus piscifermentans		100.0	0.52	BGA 1			
623450	**Good identification**						**623450**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.41	BGA 1		
623452	**Acceptable identification**					**623452**
Staphylococcus piscifermentans		93.70	0.44	BGA 1		
Staphylococcus simulans		6.30	0.09			
623500	**Good identification**					**623500**
Staphylococcus piscifermentans		100.0	0.52	BGA 1		
623502	**Very good identification**					**623502**
Staphylococcus piscifermentans		100.0	0.55	BGA 1		
623510	**Good identification**					**623510**
Staphylococcus piscifermentans		100.0	0.44	BGA 1		
623512	**Good identification**					**623512**
Staphylococcus piscifermentans		100.0	0.48	BGA 1		
623540	**Good identification**					**623540**
Staphylococcus piscifermentans		100.0	0.52	BGA 1		
623542	**Very good identification**					**623542**
Staphylococcus piscifermentans		100.0	0.55	BGA 1		
623550	**Good identification**					**623550**
					NOV	
Staphylococcus piscifermentans		95.66	0.44	BGA 1	1	
Staphylococcus saprophyticus subsp. bovis		4.34	0.05		99	
623552	**Good identification**					**623552**
Staphylococcus piscifermentans		100.0	0.48	BGA 1		
623600	**Good identification**					**623600**
Staphylococcus piscifermentans		100.0	0.52	BGA 1		
623602	**Low discrimination**					**623602**
Staphylococcus piscifermentans		79.59	0.55	BGA 1		
Staphylococcus simulans		20.41	0.32			
623603	**Good identification**					**623603**
Staphylococcus simulans		98.13	0.43	ESL 1		
Staphylococcus piscifermentans		1.15	0.14			
Staphylococcus felis		0.72	-.10			
623610	**Good identification**					**623610**
Staphylococcus piscifermentans		96.50	0.44	BGA 1		
Staphylococcus simulans		3.50	0.03			
623612	**Low discrimination**					**623612**
Staphylococcus simulans		70.54	0.45	ESL 1		
Staphylococcus piscifermentans		29.47	0.48			
623613	**Very good identification**					**623613**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		98.79	0.55	ESL 1			
Staphylococcus felis		1.21	0.07				
623640	**Acceptable identification**						**623640**
Staphylococcus piscifermentans		93.46	0.52	BGA 1			
Staphylococcus warneri		5.61	0.25				
Staphylococcus auricularis		0.94	0.01				
623642	**Low discrimination**						**623642**
Staphylococcus piscifermentans		88.61	0.55	BGA 1			
Staphylococcus simulans		10.21	0.25				
Staphylococcus warneri		1.18	0.15				
623643	**Acceptable identification**						**623643**
Staphylococcus simulans		94.50	0.36	ESL 1			
Staphylococcus petrasii subsp. petrasii		3.03	0.00				
Staphylococcus piscifermentans		2.47	0.14				
623650	**Low discrimination**						**623650**
					NOV	PYR	
Staphylococcus piscifermentans		87.20	0.44	BGA 1	1	15	
Staphylococcus warneri		8.84	0.22		1	8	
Staphylococcus saprophyticus subsp. bovis		3.96	0.05		99	99	
623652	**Low discrimination**						**623652**
Staphylococcus simulans		51.26	0.37	ESL 1			
Staphylococcus piscifermentans		47.66	0.48				
Staphylococcus warneri		1.07	0.12				
623653	**Good identification**						**623653**
Staphylococcus simulans		100.0	0.48	ESL 1			
623700	**Very good identification**						**623700**
Staphylococcus piscifermentans		100.0	0.55	BGA 1			
623702	**Very good identification**						**623702**
Staphylococcus piscifermentans		98.54	0.59	BGA 1			
Staphylococcus simulans		1.47	0.10				
623710	**Good identification**						**623710**
Staphylococcus piscifermentans		100.0	0.48	BGA 1			
623712	**Low discrimination**						**623712**
Staphylococcus piscifermentans		87.81	0.51	BGA 1			
Staphylococcus simulans		12.19	0.23				
623713	**Low discrimination**						**623713**
Staphylococcus simulans		36.87	0.34	ESL 1			
Staphylococcus condimenti		28.47	0.18	GAL 8			
Staphylococcus felis		17.34	0.18				
623740	**Very good identification**						**623740**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus piscifermentans		98.75	0.55	BGA 1				
Staphylococcus warneri		1.25	0.15					
623742	**Very good identification**							**623742**
Staphylococcus piscifermentans		99.34	0.59	BGA 1				
Staphylococcus simulans		0.66	0.03					
623750	**Low discrimination**							**623750**
					NOV	PYR		
Staphylococcus piscifermentans		87.53	0.48	BGA 1	1	15		
Staphylococcus saprophyticus subsp. bovis		10.60	0.18		99	99		
Staphylococcus warneri		1.87	0.12		1	8		
623752	**Acceptable identification**							**623752**
					NOV			
Staphylococcus piscifermentans		92.38	0.51	BGA 1	1			
Staphylococcus simulans		5.76	0.16		1			
Staphylococcus saprophyticus subsp. bovis		1.86	0.05		99			
623773	**Low discrimination**							**623773**
					OXI			
Staphylococcus equorum		69.60	0.34	ARG 1	1			
Staphylococcus xylosus		17.66	0.20	GLR 90	1			
Staphylococcus lentus		6.85	0.18		99			
624213	**Low discrimination**							**624213**
					COA			
Staphylococcus simulans		78.04	0.32	NIT 96	1			
Staphylococcus felis		13.17	0.07	TRE 95	1			
Staphylococcus intermedius		6.07	0.07		99			
624303	**Low discrimination**							**624303**
					YEP	PYR	COA	
Staphylococcus intermedius		67.87	0.39	NIT 93	1	90	99	
Staphylococcus chromogenes		26.44	0.30	TRE 93	95	45	1	
Staphylococcus hyicus		2.32	0.04		1	1	24	
624311	**Good identification**							**624311**
					COA	YEP		
Staphylococcus intermedius		97.30	0.32	NIT 93	99	1		
Staphylococcus chromogenes		2.70	-.01	TRE 93	1	95		
624313	**Low discrimination**							**624313**
					COA			
Staphylococcus intermedius		79.83	0.48	NIT 93	99			
Staphylococcus condimenti		9.62	0.18	TRE 93	1			
Staphylococcus felis		5.86	0.18		1			
624341	**Low discrimination**							**624341**
					COA	YEP	PYR	
Staphylococcus intermedius		68.53	0.27	NIT 93	99	1	90	
Staphylococcus epidermidis		19.33	0.18	TRE 93	1	1	1	
Staphylococcus chromogenes		12.15	0.10	LAC 86	1	95	45	
624343	**Low discrimination**							**624343**
					COA	YEP	PYR	
Staphylococcus intermedius		71.17	0.43	NIT 93	99	1	90	
Staphylococcus chromogenes		18.48	0.30	TRE 93	1	95	45	
Staphylococcus epidermidis		10.35	0.28		1	1	1	
624351	**Good identification**							**624351**
					COA	YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus intermedius		98.19	0.36	NIT 93	99	1	
Staphylococcus chromogenes		1.82	-.01	TRE 93	1	95	
624353	**Acceptable identification**						**624353**
					COA	YEP	
Staphylococcus intermedius		91.12	0.52	NIT 93	99	1	
Staphylococcus delphini		6.10	0.18	TRE 93	99	1	
Staphylococcus chromogenes		2.47	0.18		1	95	
624413	**Low discrimination**						**624413**
Staphylococcus felis		84.65	0.42	NIT 99			
Staphylococcus simulans		8.14	0.30				
Staphylococcus condimenti		6.36	0.14				
624450	**Good identification**						**624450**
Staphylococcus kloosii		100.0	0.46	ARG 1			
624452	**Low discrimination**						**624452**
					NOV	YEP	
Staphylococcus kloosii		77.88	0.47	ARG 1	99	99	
Staphylococcus saprophyticus subsp. saprophyticus		11.66	0.02		99	16	
Staphylococcus simulans		10.46	0.12		1	1	
624470	**Good identification**						**624470**
Staphylococcus kloosii		100.0	0.34	ARG 1			
624472	**Good identification**						**624472**
Staphylococcus kloosii		100.0	0.35	ARG 1			
624503	**Low discrimination**						**624503**
					COA		
Staphylococcus felis		78.47	0.36	NIT 99	1		
Staphylococcus condimenti		11.79	0.14	MAN 87	1		
Staphylococcus intermedius		7.29	0.22		99		
624513	**Low discrimination**						**624513**
					COA		
Staphylococcus condimenti		68.10	0.55	NIT 99	1		
Staphylococcus felis		30.64	0.53		1		
Staphylococcus intermedius		1.15	0.30		99		
624550	**Good identification**						**624550**
Staphylococcus kloosii		100.0	0.52	ARG 1			
624552	**Very good identification**						**624552**
Staphylococcus kloosii		100.0	0.53	ARG 1			
624553	**Low discrimination**						**624553**
					COA		
Staphylococcus intermedius		58.73	0.34	NIT 93	99		
Staphylococcus condimenti		23.43	0.14	SUC 99	1		
Staphylococcus felis		10.54	0.12		1		
624570	**Good identification**						**624570**
Staphylococcus kloosii		100.0	0.40	ARG 1			
624572	**Good identification**						**624572**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus kloosii		100.0	0.41	ARG 1				
624602	**Good identification**							**624602**
Staphylococcus simulans		98.25	0.35	NIT 96				
Staphylococcus felis		1.75	-.10					
624603	**Low discrimination**							**624603**
					YEP			
Staphylococcus simulans		51.95	0.46	NIT 96	1			
Staphylococcus felis		27.30	0.31		1			
Staphylococcus chromogenes		15.51	0.32		95			
624612	**Good identification**							**624612**
Staphylococcus simulans		97.11	0.47	NIT 96				
Staphylococcus felis		2.89	0.07					
624613	**Low discrimination**							**624613**
					COA			
Staphylococcus simulans		49.91	0.58	NIT 96	1			
Staphylococcus felis		43.89	0.48		1			
Staphylococcus intermedius		2.72	0.30		99			
624640	**Good identification**							**624640**
Staphylococcus warneri		100.0	0.33	PHS 1				
624642	**Low discrimination**							**624642**
					NOV			
Staphylococcus simulans		78.46	0.28	NIT 96	1			
Staphylococcus warneri		15.40	0.23		1			
Staphylococcus saprophyticus subsp. saprophyticus		6.14	-.06		99			
624643	**Low discrimination**							**624643**
					COA	YEP		
Staphylococcus simulans		51.42	0.39	NIT 96	1	1		
Staphylococcus chromogenes		34.16	0.32		1	95		
Staphylococcus intermedius		13.82	0.25		99	1		
624650	**Low discrimination**							**624650**
					NOV			
Staphylococcus warneri		46.80	0.30	PHS 1	1			
Staphylococcus saprophyticus subsp. saprophyticus		36.34	0.07		99			
Staphylococcus kloosii		12.43	0.26		99			
624652	**Low discrimination**							**624652**
					NOV			
Staphylococcus simulans		65.13	0.40	NIT 96	1			
Staphylococcus saprophyticus subsp. saprophyticus		30.56	0.22		99			
Staphylococcus warneri		2.32	0.20		1			
624653	**Low discrimination**							**624653**
					COA	YEP		
Staphylococcus simulans		79.81	0.51	NIT 96	1	1		
Staphylococcus intermedius		14.49	0.34		99	1		
Staphylococcus chromogenes		3.74	0.20		1	95		
624701	**Low discrimination**							**624701**
					YEP	PYR	COA	
Staphylococcus intermedius		79.39	0.46	NIT 93	1	90	99	
Staphylococcus chromogenes		18.27	0.32	LAC 86	95	45	1	
Staphylococcus hyicus		1.25	0.04		1	1	24	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
624703	**Low discrimination**							**624703**
					COA	YEP		
Staphylococcus intermedius		61.61	0.63	NIT 93	99	1		
Staphylococcus chromogenes		20.77	0.51		1	95		
Staphylococcus felis		13.61	0.42		1	1		
624711	**Low discrimination**							**624711**
					COA			
Staphylococcus intermedius		89.80	0.55	NIT 93	99			
Staphylococcus condimenti		5.00	0.18	LAC 86	1			
Staphylococcus felis		3.05	0.18		1			
624712	**Low discrimination**							**624712**
					COA			
Staphylococcus intermedius		33.82	0.30	NIT 93	99			
Staphylococcus condimenti		30.36	0.18	MNS 99	1			
Staphylococcus felis		18.49	0.18		1			
624713	**Low discrimination**							**624713**
					COA			
Staphylococcus intermedius		40.07	0.72	NIT 93	99			
Staphylococcus condimenti		35.97	0.59		1			
Staphylococcus felis		21.91	0.59		1			
624733	**Low discrimination**							**624733**
					COA			
Staphylococcus intermedius		40.33	0.30	NIT 93	99			
Staphylococcus condimenti		36.20	0.18	XYL 1	1			
Staphylococcus felis		22.05	0.18		1			
624741	**Low discrimination**							**624741**
					COA	YEP		
Staphylococcus intermedius		86.70	0.50	NIT 93	99	1		
Staphylococcus chromogenes		13.30	0.32	LAC 86	1	95		
624743	**Low discrimination**							**624743**
					COA	YEP		
Staphylococcus intermedius		81.33	0.66	NIT 93	99	1		
Staphylococcus chromogenes		18.28	0.51		1	95		
Staphylococcus simulans		0.27	0.17		1	1		
624751	**Very good identification**							**624751**
					COA	YEP		
Staphylococcus intermedius		98.43	0.59	NIT 93	99	1		
Staphylococcus chromogenes		1.58	0.20	LAC 86	1	95		
624752	**Low discrimination**							**624752**
					COA	NOV	YEP	
Staphylococcus intermedius		74.08	0.34	NIT 93	99	1	1	
Staphylococcus simulans		10.59	0.18	MNS 99	1	1	1	
Staphylococcus kloosii		7.25	0.32		1	99	99	
624753	**Very good identification**							**624753**
					COA	PYR	YEP	
Staphylococcus intermedius		95.13	0.75	NIT 93	99	90	1	
Staphylococcus chromogenes		2.23	0.40		1	45	95	
Staphylococcus aureus subsp. aureus		1.23	0.22		99	10	84	
624773	**Low discrimination**							**624773**
					COA	PYR	NOV	
Staphylococcus intermedius		71.81	0.34	NIT 93	99	90	1	
Staphylococcus lutrae		23.88	0.18	XYL 1	99	10	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus xylosus		2.63	0.03		1	65	99	
625450	**Good identification**							**625450**
Staphylococcus kloosii		100.0	0.46	ARG 1				
625452	**Good identification**							**625452**
Staphylococcus kloosii		100.0	0.47	ARG 1				
625470	**Good identification**							**625470**
Staphylococcus kloosii		100.0	0.34	ARG 1				
625472	**Good identification**							**625472**
Staphylococcus kloosii		100.0	0.35	ARG 1				
625512	**Good identification**							**625512**
Staphylococcus kloosii		100.0	0.27	ARG 1				
625550	**Good identification**							**625550**
Staphylococcus kloosii		100.0	0.52	ARG 1				
625552	**Very good identification**							**625552**
Staphylococcus kloosii		100.0	0.53	ARG 1				
625570	**Good identification**							**625570**
Staphylococcus kloosii		100.0	0.40	ARG 1				
625572	**Good identification**							**625572**
Staphylococcus kloosii		100.0	0.41	ARG 1				
625713	**Low discrimination**							**625713**
					COA			
Staphylococcus intermedius		40.33	0.30	ESL 1	99			
Staphylococcus condimenti		36.20	0.18	NIT 93	1			
Staphylococcus felis		22.05	0.18		1			
625750	**Good identification**							**625750**
Staphylococcus kloosii		100.0	0.32	ARG 1				
625752	**Good identification**							**625752**
Staphylococcus kloosii		100.0	0.32	ARG 1				
625753	**Good identification**							**625753**
					COA	YEP		
Staphylococcus intermedius		97.71	0.34	ESL 1	99	1		
Staphylococcus chromogenes		2.29	-.01	NIT 93	1	95		
626001	**Acceptable identification**							**626001**
					COA			
Staphylococcus carnosus subsp. carnosus		90.56	0.31	URE 1	1			
Staphylococcus schleiferi subsp. schleiferi		7.17	0.08		1			
Staphylococcus schleiferi subsp. coagulans		2.27	0.02		99			
626003	**Low discrimination**							**626003**
Staphylococcus carnosus subsp. carnosus		72.57	0.43	URE 1				

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus felis		18.90	0.25					
Staphylococcus simulans		3.84	0.19					
626011	**Good identification**							**626011**
					COA			
Staphylococcus carnosus subsp. carnosus		97.72	0.43	URE 1	1			
Staphylococcus felis		1.72	0.01		1			
Staphylococcus schleiferi subsp. coagulans		0.56	0.01		99			
626013	**Low discrimination**							**626013**
Staphylococcus carnosus subsp. carnosus		65.26	0.55	URE 1				
Staphylococcus felis		28.44	0.42					
Staphylococcus simulans		3.46	0.32					
626101	**Low discrimination**							**626101**
					COA	YEP		
Staphylococcus schleiferi subsp. coagulans		82.60	0.43	VPT 99	99	1		
Staphylococcus carnosus subsp. carnosus		8.34	0.18	LAC 86	1	1		
Staphylococcus chromogenes		3.48	0.10		1	95		
626103	**Low discrimination**							**626103**
					COA			
Staphylococcus schleiferi subsp. coagulans		70.53	0.59	VPT 99	99			
Staphylococcus felis		16.17	0.36		1			
Staphylococcus carnosus subsp. carnosus		4.64	0.31		1			
626111	**Low discrimination**							**626111**
					COA			
Staphylococcus schleiferi subsp. coagulans		54.20	0.42	VPT 99	99			
Staphylococcus carnosus subsp. carnosus		23.71	0.31	LAC 86	1			
Staphylococcus condimenti		12.43	0.14		1			
626113	**Low discrimination**							**626113**
					COA			
Staphylococcus condimenti		54.91	0.55	TRE 99	1			
Staphylococcus felis		24.71	0.53		1			
Staphylococcus schleiferi subsp. coagulans		14.86	0.58		99			
626143	**Low discrimination**							**626143**
					COA	YEP	PYR	
Staphylococcus chromogenes		52.61	0.30	SUC 99	1	95	45	
Staphylococcus intermedius		27.19	0.25	TRE 92	99	1	90	
Staphylococcus epidermidis		9.62	0.18		1	1	1	
626153	**Low discrimination**							**626153**
					COA			
Staphylococcus intermedius		40.12	0.34	SUC 99	99			
Staphylococcus delphini		20.01	0.18	TRE 93	99			
Staphylococcus condimenti		16.01	0.14		1			
626201	**Low discrimination**							**626201**
					YEP	PYR	COA	
Staphylococcus chromogenes		86.00	0.31	GAL 90	95	45	1	
Staphylococcus hyicus		4.62	0.01	TRE 92	1	1	24	
Staphylococcus intermedius		3.95	0.05	LAC 90	1	90	99	
626202	**Acceptable identification**							**626202**
					YEP			
Staphylococcus simulans		92.35	0.37	TRE 95	1			
Staphylococcus chromogenes		6.35	0.10		95			
Staphylococcus felis		1.30	-.10		1			

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
626203	**Low discrimination**							**626203**
					YEP			
Staphylococcus chromogenes		55.32	0.51	GAL 90	95			
Staphylococcus simulans		27.20	0.48	TRE 92	1			
Staphylococcus felis		11.32	0.31		1			
626212	**Good identification**							**626212**
					YEP			
Staphylococcus simulans		97.24	0.49	TRE 95	1			
Staphylococcus felis		2.29	0.07		1			
Staphylococcus chromogenes		0.47	-.01		95			
626213	**Low discrimination**							**626213**
					YEP			
Staphylococcus simulans		50.82	0.60	TRE 95	1			
Staphylococcus felis		35.38	0.48		1			
Staphylococcus chromogenes		7.29	0.40		95			
626240	**Good identification**							**626240**
Staphylococcus epidermidis		100.0	0.28	VPT 96				
626241	**Low discrimination**							**626241**
					COA	YEP	PYR	
Staphylococcus chromogenes		52.23	0.31	GAL 90	1	95	45	
Staphylococcus epidermidis		43.12	0.33	TRE 92	1	1	1	
Staphylococcus intermedius		3.60	0.09	LAC 90	99	1	90	
626242	**Low discrimination**							**626242**
					YEP			
Staphylococcus epidermidis		67.28	0.38	VPT 96	1			
Staphylococcus simulans		28.38	0.30		1			
Staphylococcus chromogenes		4.35	0.10		95			
626243	**Low discrimination**							**626243**
					YEP			
Staphylococcus chromogenes		61.29	0.51	GAL 90	95			
Staphylococcus epidermidis		17.80	0.44	TRE 92	1			
Staphylococcus simulans		13.54	0.41		1			
626252	**Low discrimination**							**626252**
					COA	YEP		
Staphylococcus simulans		79.95	0.42	TRE 95	1	1		
Staphylococcus delphini		19.19	0.18		99	1		
Staphylococcus chromogenes		0.86	-.01		1	95		
626253	**Low discrimination**							**626253**
					COA	YEP		
Staphylococcus delphini		80.68	0.59	BGA 1	99	1		
Staphylococcus simulans		11.37	0.53		1	1		
Staphylococcus chromogenes		3.63	0.40		1	95		
626301	**Low discrimination**							**626301**
					COA	YEP		
Staphylococcus chromogenes		60.20	0.51	TRE 92	1	95		
Staphylococcus intermedius		30.39	0.46	LAC 90	99	1		
Staphylococcus schleiferi subsp. coagulans		4.55	0.32		99	1		
626302	**Low discrimination**							**626302**
					COA	YEP		
Staphylococcus chromogenes		55.59	0.30	TRE 92	1	95		
Staphylococcus intermedius		19.15	0.22	MNS 99	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus simulans		7.81	0.15		1	1		
626303	**Low discrimination**							**626303**
					YEP	PYR	COA	
Staphylococcus chromogenes		63.50	0.71	TRE 92	95	45	1	
Staphylococcus intermedius		21.88	0.63		1	90	99	
Staphylococcus hyicus		5.57	0.45		1	1	24	
626311	**Low discrimination**							**626311**
					COA	YEP		
Staphylococcus intermedius		74.22	0.55	TRE 93	99	1		
Staphylococcus chromogenes		15.34	0.40	LAC 86	1	95		
Staphylococcus condimenti		4.13	0.18		1	1		
626312	**Low discrimination**							**626312**
					COA			
Staphylococcus intermedius		29.40	0.30	TRE 93	99			
Staphylococcus condimenti		26.39	0.18	MNS 99	1			
Staphylococcus simulans		17.73	0.28		1			
626313	**Low discrimination**							**626313**
					COA			
Staphylococcus intermedius		35.36	0.72	TRE 93	99			
Staphylococcus condimenti		31.75	0.59		1			
Staphylococcus felis		19.33	0.59		1			
626323	**Low discrimination**							**626323**
					YEP	PYR	COA	
Staphylococcus chromogenes		64.11	0.30	TRE 92	95	45	1	
Staphylococcus intermedius		22.09	0.22	XYL 1	1	90	99	
Staphylococcus hyicus		5.62	0.04		1	1	24	
626333	**Low discrimination**							**626333**
					COA			
Staphylococcus intermedius		35.74	0.30	TRE 93	99			
Staphylococcus condimenti		32.08	0.18	XYL 1	1			
Staphylococcus felis		19.54	0.18		1			
626340	**Acceptable identification**							**626340**
					COA	YEP	PYR	
Staphylococcus epidermidis		94.04	0.43	VPT 96	1	1	1	
Staphylococcus chromogenes		3.39	0.10		1	95	45	
Staphylococcus intermedius		2.57	0.09		99	1	90	
626341	**Low discrimination**							**626341**
					COA	YEP	PYR	
Staphylococcus chromogenes		43.92	0.51	TRE 92	1	95	45	
Staphylococcus intermedius		33.25	0.50	LAC 90	99	1	90	
Staphylococcus epidermidis		22.83	0.49		1	1	1	
626342	**Low discrimination**							**626342**
					COA	YEP	PYR	
Staphylococcus epidermidis		86.06	0.54	VPT 96	1	1	1	
Staphylococcus chromogenes		8.83	0.30		1	95	45	
Staphylococcus intermedius		4.56	0.25		99	1	90	
626343	**Low discrimination**							**626343**
					COA	YEP	PYR	
Staphylococcus chromogenes		58.44	0.71	TRE 92	1	95	45	
Staphylococcus intermedius		30.20	0.66		99	1	90	
Staphylococcus epidermidis		10.69	0.59		1	1	1	
626351	**Low discrimination**							**626351**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	YEP		
Staphylococcus intermedius		85.38	0.59	TRE 93	99	1		
Staphylococcus chromogenes		11.76	0.40	LAC 86	1	95		
Staphylococcus delphini		2.64	0.18		99	1		
626352	**Low discrimination**							**626352**
					COA	YEP		
Staphylococcus intermedius		51.24	0.34	TRE 93	99	1		
Staphylococcus delphini		25.55	0.18	MNS 99	99	1		
Staphylococcus chromogenes		10.34	0.18		1	95		
626353	**Low discrimination**							**626353**
					COA	YEP		
Staphylococcus intermedius		57.76	0.75	TRE 93	99	1		
Staphylococcus delphini		28.81	0.59		99	1		
Staphylococcus chromogenes		11.66	0.59		1	95		
626363	**Low discrimination**							**626363**
					COA	PYR	YEP	
Staphylococcus chromogenes		51.57	0.30	TRE 92	1	45	95	
Staphylococcus intermedius		26.65	0.25	XYL 1	99	90	1	
Staphylococcus lutrae		12.34	0.12		99	10	1	
626373	**Low discrimination**							**626373**
					PYR			
Staphylococcus intermedius		49.19	0.34	TRE 93	90			
Staphylococcus delphini		24.53	0.18	XYL 1	15			
Staphylococcus lutrae		16.35	0.18		10			
626402	**Low discrimination**							**626402**
Staphylococcus simulans		49.83	0.35	SUC 96				
Staphylococcus felis		43.19	0.25					
Staphylococcus piscifermentans		6.98	0.28					
626403	**Good identification**							**626403**
Staphylococcus felis		93.15	0.66	MAN 87				
Staphylococcus simulans		3.63	0.46					
Staphylococcus carnosus subsp. carnosus		1.55	0.35					
626411	**Low discrimination**							**626411**
Staphylococcus felis		74.09	0.42	LAC 99				
Staphylococcus carnosus subsp. carnosus		18.21	0.35					
Staphylococcus condimenti		5.57	0.14					
626412	**Low discrimination**							**626412**
Staphylococcus felis		56.02	0.42	MNS 99				
Staphylococcus simulans		38.63	0.47					
Staphylococcus condimenti		4.21	0.14					
626413	**Good identification**							**626413**
Staphylococcus felis		90.17	0.83					
Staphylococcus condimenti		6.78	0.55					
Staphylococcus simulans		2.10	0.58					
626433	**Acceptable identification**							**626433**
Staphylococcus felis		90.22	0.42	XYL 1				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus condimenti		6.78	0.14				
Staphylococcus simulans		2.10	0.17				
626442	**Low discrimination**						**626442**
Staphylococcus simulans		76.24	0.28	SUC 96			
Staphylococcus piscifermentans		23.76	0.28				
626443	**Low discrimination**						**626443**
					YEP		
Staphylococcus simulans		43.93	0.39	SUC 96	1		
Staphylococcus chromogenes		29.18	0.32		95		
Staphylococcus felis		25.31	0.25		1		
626452	**Acceptable identification**						**626452**
Staphylococcus simulans		91.10	0.40	SUC 96			
Staphylococcus piscifermentans		3.04	0.21				
Staphylococcus felis		2.97	0.01				
626453	**Low discrimination**						**626453**
Staphylococcus simulans		47.12	0.51	SUC 96			
Staphylococcus felis		45.43	0.42				
Staphylococcus condimenti		3.42	0.14				
626500	**Low discrimination**						**626500**
Staphylococcus piscifermentans		82.69	0.28	BGA 1			
Staphylococcus felis		17.31	-.06	ESL 90			
626501	**Low discrimination**						**626501**
					COA	YEP	
Staphylococcus felis		54.79	0.36	MAN 87	1	1	
Staphylococcus chromogenes		18.87	0.32	LAC 99	1	95	
Staphylococcus intermedius		11.00	0.28		99	1	
626502	**Low discrimination**						**626502**
Staphylococcus felis		77.14	0.36	MAN 87			
Staphylococcus condimenti		11.59	0.14	MNS 99			
Staphylococcus piscifermentans		5.58	0.32				
626503	**Low discrimination**						**626503**
					YEP		
Staphylococcus felis		83.37	0.77	MAN 87	1		
Staphylococcus condimenti		12.53	0.55		1		
Staphylococcus chromogenes		2.61	0.51		95		
626511	**Low discrimination**						**626511**
					COA		
Staphylococcus condimenti		66.59	0.55	LAC 99	1		
Staphylococcus felis		29.96	0.53		1		
Staphylococcus intermedius		2.43	0.37		99		
626512	**Low discrimination**						**626512**
Staphylococcus condimenti		68.36	0.55	MNS 99			
Staphylococcus felis		30.76	0.53				
Staphylococcus simulans		0.55	0.26				
626513	**Low discrimination**						**626513**
					COA		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus condimenti		68.80	0.96		1			
Staphylococcus felis		30.96	0.94		1			
Staphylococcus intermedius		0.16	0.54		99			
626523	**Low discrimination**							**626523**
					COA			
Staphylococcus felis		79.46	0.36	MAN 87	1			
Staphylococcus condimenti		11.94	0.14	XYL 1	1			
Staphylococcus lutrae		5.13	0.12		99			
626533	**Low discrimination**							**626533**
					COA			
Staphylococcus condimenti		68.44	0.55	XYL 1	1			
Staphylococcus felis		30.79	0.53		1			
Staphylococcus lutrae		0.58	0.18		99			
626540	**Good identification**							**626540**
Staphylococcus piscifermentans		100.0	0.28	BGA 1				
626541	**Low discrimination**							**626541**
					COA	YEP		
Staphylococcus chromogenes		52.52	0.32	SUC 99	1	95		
Staphylococcus intermedius		45.94	0.32	LAC 90	99	1		
Staphylococcus felis		1.54	-.06		1	1		
626542	**Low discrimination**							**626542**
					COA	YEP		
Staphylococcus piscifermentans		49.60	0.32	BGA 1	1	1		
Staphylococcus chromogenes		21.45	0.10	ESL 90	1	95		
Staphylococcus intermedius		12.81	0.07		99	1		
626543	**Low discrimination**							**626543**
					COA	YEP		
Staphylococcus chromogenes		49.84	0.51	SUC 99	1	95		
Staphylococcus intermedius		29.76	0.48		99	1		
Staphylococcus felis		16.08	0.36		1	1		
626550	**Acceptable identification**							**626550**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		91.56	0.34	ARG 1	1	99		
Staphylococcus piscifermentans		6.04	0.21		1	1		
Staphylococcus intermedius		2.40	0.00		99	1		
626551	**Low discrimination**							**626551**
					COA	YEP		
Staphylococcus intermedius		66.94	0.41	SUC 99	99	1		
Staphylococcus condimenti		12.35	0.14	LAC 86	1	1		
Staphylococcus chromogenes		7.98	0.20		1	95		
626553	**Low discrimination**							**626553**
					COA			
Staphylococcus condimenti		52.34	0.55	MLT 1	1			
Staphylococcus felis		23.55	0.53		1			
Staphylococcus intermedius		17.60	0.57		99			
626563	**Very good identification**							**626563**
					COA	PYR	YEP	
Staphylococcus lutrae		99.07	0.53	ARG 1	99	10	1	
Staphylococcus chromogenes		0.49	0.10		1	45	95	
Staphylococcus intermedius		0.29	0.07		99	90	1	
626573	**Very good identification**							**626573**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COA	
Staphylococcus lutrae		97.81	0.59	ARG 1	99	
Staphylococcus condimenti		1.19	0.14		1	
Staphylococcus felis		0.53	0.12		1	
626601	**Low discrimination**					**626601**
					YEP	
Staphylococcus chromogenes		74.59	0.53	GAL 90	95	
Staphylococcus felis		11.94	0.31	LAC 90	1	
Staphylococcus simulans		5.51	0.33		1	
626602	**Low discrimination**					**626602**
					YEP	
Staphylococcus simulans		88.65	0.63		1	
Staphylococcus felis		6.50	0.31		1	
Staphylococcus chromogenes		3.69	0.32		95	
626603	**Low discrimination**					**626603**
					YEP	
Staphylococcus felis		47.79	0.72	MAN 87	1	
Staphylococcus chromogenes		27.14	0.73		95	
Staphylococcus simulans		22.04	0.74		1	
626610	**Low discrimination**					**626610**
Staphylococcus simulans		82.18	0.35	LAC 99		
Staphylococcus felis		10.08	0.07			
Staphylococcus piscifermentans		5.09	0.21			
626611	**Low discrimination**					**626611**
					YEP	
Staphylococcus felis		57.86	0.48	LAC 99	1	
Staphylococcus simulans		15.95	0.46		1	
Staphylococcus chromogenes		15.23	0.42		95	
626612	**Low discrimination**					**626612**
Staphylococcus simulans		88.16	0.76			
Staphylococcus felis		10.81	0.48			
Staphylococcus condimenti		0.60	0.18			
626613	**Low discrimination**					**626613**
Staphylococcus felis		73.34	0.89			
Staphylococcus simulans		20.22	0.87			
Staphylococcus condimenti		4.07	0.59			
626623	**Low discrimination**					**626623**
					YEP	
Staphylococcus felis		47.88	0.31	MAN 87	1	
Staphylococcus chromogenes		27.19	0.32	XYL 1	95	
Staphylococcus simulans		22.08	0.33		1	
626632	**Low discrimination**					**626632**
Staphylococcus simulans		89.07	0.35	XYL 1		
Staphylococcus felis		10.93	0.07			
626633	**Low discrimination**					**626633**
Staphylococcus felis		73.34	0.48	XYL 1		
Staphylococcus simulans		20.22	0.46			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus condimenti		4.07	0.18				
626640	**Low discrimination**						**626640**
					YEP		
Staphylococcus piscifermentans		26.88	0.28	BGA 1	1		
Staphylococcus warneri		21.77	0.25	ESL 90	43		
Staphylococcus chromogenes		21.31	0.12		95		
626641	**Low discrimination**						**626641**
					COA	YEP	
Staphylococcus chromogenes		89.27	0.53	GAL 90	1	95	
Staphylococcus intermedius		7.10	0.32	LAC 90	99	1	
Staphylococcus simulans		2.96	0.26		1	1	
626642	**Low discrimination**						**626642**
					YEP		
Staphylococcus simulans		88.85	0.56		1		
Staphylococcus chromogenes		8.23	0.32		95		
Staphylococcus piscifermentans		1.73	0.32		1		
626643	**Low discrimination**						**626643**
					COA	YEP	
Staphylococcus chromogenes		69.51	0.73	GAL 90	1	95	
Staphylococcus simulans		25.37	0.67		1	1	
Staphylococcus intermedius		3.77	0.48		99	1	
626650	**Low discrimination**						**626650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		58.99	0.34	ARG 1	99	99	
Staphylococcus simulans		28.23	0.28		1	70	
Staphylococcus warneri		5.32	0.22		1	8	
626651	**Low discrimination**						**626651**
					COA	YEP	
Staphylococcus chromogenes		42.49	0.42	GAL 90	1	95	
Staphylococcus intermedius		32.39	0.41	LAC 90	99	1	
Staphylococcus simulans		19.99	0.38		1	1	
626652	**Very good identification**						**626652**
					NOV	YEP	
Staphylococcus simulans		97.91	0.69		1	1	
Staphylococcus chromogenes		0.64	0.20		1	95	
Staphylococcus saprophyticus subsp. bovis		0.52	0.22		99	80	
626653	**Low discrimination**						**626653**
					COA	YEP	
Staphylococcus simulans		71.24	0.79		1	1	
Staphylococcus chromogenes		13.77	0.62		1	95	
Staphylococcus intermedius		7.16	0.57		99	1	
626663	**Low discrimination**						**626663**
					COA	YEP	
Staphylococcus chromogenes		61.57	0.32	GAL 90	1	95	
Staphylococcus simulans		22.47	0.26	XYL 1	1	1	
Staphylococcus lutrae		11.53	0.12		99	1	
626672	**Good identification**						**626672**
Staphylococcus simulans		100.0	0.28	XYL 1			
626673	**Low discrimination**						**626673**
					COA	YEP	
Staphylococcus simulans		60.02	0.38	XYL 1	1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus lutrae		14.95	0.18		99	1		
Staphylococcus chromogenes		11.60	0.20		1	95		
626700	**Low discrimination**							**626700**
					COA	YEP		
Staphylococcus chromogenes		51.48	0.32	MNS 99	1	95		
Staphylococcus intermedius		30.02	0.28	LAC 90	99	1		
Staphylococcus piscifermentans		10.82	0.32		1	1		
626701	**Low discrimination**							**626701**
					YEP	PYR	COA	
Staphylococcus chromogenes		58.19	0.73	LAC 90	95	45	1	
Staphylococcus intermedius		33.93	0.70		1	90	99	
Staphylococcus hyicus		3.99	0.45		1	1	24	
626702	**Low discrimination**							**626702**
					COA	YEP		
Staphylococcus chromogenes		40.31	0.51	MNS 99	1	95		
Staphylococcus felis		26.41	0.42		1	1		
Staphylococcus intermedius		16.05	0.45		99	1		
626703	**Low discrimination**							**626703**
					COA	YEP		
Staphylococcus chromogenes		45.02	0.92		1	95		
Staphylococcus felis		29.49	0.83		1	1		
Staphylococcus intermedius		17.92	0.86		99	1		
626710	**Low discrimination**							**626710**
					COA			
Staphylococcus intermedius		50.49	0.37	MNS 99	99			
Staphylococcus condimenti		20.96	0.18	LAC 86	1			
Staphylococcus felis		12.76	0.18		1			
626711	**Low discrimination**							**626711**
					COA			
Staphylococcus intermedius		54.08	0.78	LAC 86	99			
Staphylococcus condimenti		22.45	0.59		1			
Staphylococcus felis		13.67	0.59		1			
626712	**Low discrimination**							**626712**
					COA			
Staphylococcus condimenti		51.85	0.59	MNS 99	1			
Staphylococcus felis		31.58	0.59		1			
Staphylococcus intermedius		7.75	0.54		99			
626713	**Low discrimination**							**626713**
					COA			
Staphylococcus condimenti		55.48	1.00		1			
Staphylococcus felis		33.79	1.00		1			
Staphylococcus intermedius		8.29	0.95		99			
626721	**Low discrimination**							**626721**
					YEP	PYR	COA	
Staphylococcus chromogenes		58.44	0.32	XYL 1	95	45	1	
Staphylococcus intermedius		34.08	0.28	LAC 90	1	90	99	
Staphylococcus hyicus		4.01	0.04		1	1	24	
626723	**Low discrimination**							**626723**
					COA	YEP		
Staphylococcus chromogenes		44.60	0.51	XYL 1	1	95		
Staphylococcus felis		29.22	0.42		1	1		
Staphylococcus intermedius		17.75	0.45		99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
626731	**Low discrimination**							**626731**
					COA			
Staphylococcus intermedius		54.15	0.37	XYL 1	99			
Staphylococcus condimenti		22.48	0.18	LAC 86	1			
Staphylococcus felis		13.69	0.18		1			
626733	**Low discrimination**							**626733**
					COA			
Staphylococcus condimenti		55.31	0.59	XYL 1	1			
Staphylococcus felis		33.68	0.59		1			
Staphylococcus intermedius		8.27	0.54		99			
626740	**Low discrimination**							**626740**
					COA	YEP		
Staphylococcus chromogenes		45.19	0.32	MNS 99	1	95		
Staphylococcus intermedius		39.53	0.32	LAC 90	99	1		
Staphylococcus piscifermentans		9.50	0.32		1	1		
626741	**Low discrimination**							**626741**
					YEP	PYR	COA	
Staphylococcus chromogenes		53.27	0.73	LAC 90	95	45	1	
Staphylococcus intermedius		46.60	0.73		1	90	99	
Staphylococcus hyicus		0.04	0.04		1	1	24	
626742	**Low discrimination**							**626742**
					COA	YEP		
Staphylococcus chromogenes		56.96	0.51	MNS 99	1	95		
Staphylococcus intermedius		34.01	0.48		99	1		
Staphylococcus simulans		5.94	0.34		1	1		
626743	**Low discrimination**							**626743**
					COA	YEP		
Staphylococcus chromogenes		62.08	0.92		1	95		
Staphylococcus intermedius		37.07	0.89		99	1		
Staphylococcus felis		0.41	0.42		1	1		
626750	**Low discrimination**							**626750**
					COA	NOV	YEP	
Staphylococcus saprophyticus subsp. bovis		56.54	0.47	ARG 1	1	99	80	
Staphylococcus intermedius		36.69	0.41		99	1	1	
Staphylococcus chromogenes		4.38	0.20		1	1	95	
626751	**Low discrimination**							**626751**
					COA	PYR	YEP	
Staphylococcus intermedius		88.17	0.82	LAC 86	99	90	1	
Staphylococcus chromogenes		10.52	0.62		1	45	95	
Staphylococcus aureus subsp. aureus		0.53	0.22		99	10	84	
626752	**Low discrimination**							**626752**
					COA	YEP		
Staphylococcus intermedius		59.67	0.57	MNS 99	99	1		
Staphylococcus simulans		15.42	0.47		1	1		
Staphylococcus chromogenes		10.43	0.40		1	95		
626753	**Low discrimination**							**626753**
					COA	PYR	YEP	
Staphylococcus intermedius		73.88	0.98		99	90	1	
Staphylococcus chromogenes		12.91	0.81		1	45	95	
Staphylococcus aureus subsp. aureus		7.14	0.63		99	10	84	
626761	**Low discrimination**							**626761**
					COA	PYR	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus chromogenes		48.50	0.32	XYL 1	1	45	95	
Staphylococcus intermedius		42.42	0.32	LAC 90	99	90	1	
Staphylococcus lutrae		9.08	0.12		99	10	1	
626763	**Low discrimination**							**626763**
					COA	PYR	YEP	
Staphylococcus lutrae		56.14	0.53	ARG 1	99	10	1	
Staphylococcus chromogenes		27.25	0.51		1	45	95	
Staphylococcus intermedius		16.27	0.48		99	90	1	
626771	**Low discrimination**							**626771**
					COA	PYR	YEP	
Staphylococcus intermedius		74.76	0.41	XYL 1	99	90	1	
Staphylococcus lutrae		11.49	0.18	LAC 86	99	10	1	
Staphylococcus chromogenes		8.92	0.20		1	45	95	
626773	**Low discrimination**							**626773**
					COA	PYR	YEP	
Staphylococcus lutrae		64.20	0.59	ARG 1	99	10	1	
Staphylococcus intermedius		25.91	0.57		99	90	1	
Staphylococcus chromogenes		4.53	0.40		1	45	95	
627013	**Low discrimination**							**627013**
					COA			
Staphylococcus carnosus subsp. carnosus		89.58	0.36	URE 1	1			
Staphylococcus schleiferi subsp. coagulans		6.87	0.17	ESL 10	99			
Staphylococcus felis		3.55	0.01		1			
627101	**Good identification**							**627101**
					COA			
Staphylococcus schleiferi subsp. coagulans		98.85	0.42	VPT 99	99			
Staphylococcus carnosus subsp. carnosus		1.15	-.02	LAC 86	1			
627103	**Very good identification**							**627103**
					COA			
Staphylococcus schleiferi subsp. coagulans		99.01	0.59	VPT 99	99			
Staphylococcus carnosus subsp. carnosus		0.75	0.11		1			
Staphylococcus felis		0.24	-.06		1			
627111	**Good identification**							**627111**
					COA			
Staphylococcus schleiferi subsp. coagulans		95.19	0.42	VPT 99	99			
Staphylococcus carnosus subsp. carnosus		4.82	0.11	LAC 86	1			
627113	**Good identification**							**627113**
					COA			
Staphylococcus schleiferi subsp. coagulans		91.81	0.58	VPT 99	99			
Staphylococcus condimenti		3.57	0.14		1			
Staphylococcus carnosus subsp. carnosus		3.02	0.23		1			
627301	**Low discrimination**							**627301**
					COA	YEP		
Staphylococcus schleiferi subsp. coagulans		82.70	0.32	VPT 99	99	1		
Staphylococcus chromogenes		11.50	0.10	LAC 86	1	95		
Staphylococcus intermedius		5.80	0.05		99	1		
627303	**Low discrimination**							**627303**
					COA	YEP		
Staphylococcus schleiferi subsp. coagulans		76.50	0.48	VPT 99	99	1		
Staphylococcus chromogenes		15.58	0.30		1	95		
Staphylococcus intermedius		5.37	0.22		99	1		
627311	**Low discrimination**							**627311**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	YEP		
Staphylococcus schleiferi subsp. coagulans		80.13	0.31	VPT 99	99	1		
Staphylococcus intermedius		16.47	0.14	LAC 86	99	1		
Staphylococcus chromogenes		3.40	-.01		1	95		
627313	**Low discrimination**							**627313**
					COA			
Staphylococcus schleiferi subsp. coagulans		63.91	0.48	VPT 99	99			
Staphylococcus intermedius		13.14	0.30		99			
Staphylococcus condimenti		11.79	0.18		1			
627343	**Low discrimination**							**627343**
					COA	YEP	PYR	
Staphylococcus chromogenes		57.17	0.30	ESL 1	1	95	45	
Staphylococcus intermedius		29.54	0.25	TRE 92	99	1	90	
Staphylococcus epidermidis		10.46	0.18		1	1	1	
627353	**Low discrimination**							**627353**
					COA	YEP		
Staphylococcus intermedius		57.69	0.34	ESL 1	99	1		
Staphylococcus delphini		28.77	0.18	TRE 93	99	1		
Staphylococcus chromogenes		11.65	0.18		1	95		
627400	**Good identification**							**627400**
Staphylococcus piscifermentans		100.0	0.44	BGA 1				
627402	**Good identification**							**627402**
Staphylococcus piscifermentans		100.0	0.48	BGA 1				
627410	**Good identification**							**627410**
Staphylococcus piscifermentans		100.0	0.37	BGA 1				
627412	**Low discrimination**							**627412**
Staphylococcus piscifermentans		84.51	0.41	BGA 1				
Staphylococcus felis		9.17	0.01					
Staphylococcus simulans		6.32	0.06					
627413	**Low discrimination**							**627413**
Staphylococcus felis		82.80	0.42	ESL 1				
Staphylococcus carnosus subsp. carnosus		9.05	0.28					
Staphylococcus condimenti		6.22	0.14					
627440	**Good identification**							**627440**
Staphylococcus piscifermentans		100.0	0.44	BGA 1				
627442	**Good identification**							**627442**
Staphylococcus piscifermentans		100.0	0.48	BGA 1				
627450	**Good identification**							**627450**
Staphylococcus piscifermentans		100.0	0.37	BGA 1				
627452	**Good identification**							**627452**
Staphylococcus piscifermentans		96.75	0.41	BGA 1				
Staphylococcus simulans		3.25	-.01					
627500	**Good identification**							**627500**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
	Staphylococcus piscifermentans	100.0	0.48	BGA 1		
627502	**Good identification**					**627502**
	Staphylococcus piscifermentans	98.47	0.52	BGA 1		
	Staphylococcus felis	1.53	-.06			
627503	**Low discrimination**					**627503**
					COA	
	Staphylococcus felis	79.58	0.36	ESL 1	1	
	Staphylococcus condimenti	11.96	0.14	MAN 87	1	
	Staphylococcus schleiferi subsp. coagulans	3.37	0.18		99	
627510	**Good identification**					**627510**
	Staphylococcus piscifermentans	100.0	0.41	BGA 1		
627512	**Low discrimination**					**627512**
	Staphylococcus piscifermentans	56.17	0.44	BGA 1		
	Staphylococcus condimenti	30.23	0.14			
	Staphylococcus felis	13.60	0.12			
627513	**Low discrimination**					**627513**
	Staphylococcus condimenti	68.54	0.55	ESL 1		
	Staphylococcus felis	30.84	0.53			
	Staphylococcus carnosus subsp. carnosus	0.25	0.16			
627540	**Good identification**					**627540**
	Staphylococcus piscifermentans	100.0	0.48	BGA 1		
627542	**Good identification**					**627542**
	Staphylococcus piscifermentans	100.0	0.52	BGA 1		
627550	**Good identification**					**627550**
	Staphylococcus piscifermentans	100.0	0.41	BGA 1		
627552	**Good identification**					**627552**
	Staphylococcus piscifermentans	100.0	0.44	BGA 1		
627600	**Good identification**					**627600**
	Staphylococcus piscifermentans	100.0	0.48	BGA 1		
627602	**Low discrimination**					**627602**
	Staphylococcus piscifermentans	87.90	0.52	BGA 1		
	Staphylococcus simulans	11.27	0.22			
	Staphylococcus felis	0.83	-.10			
627603	**Low discrimination**					**627603**
					YEP	
	Staphylococcus felis	47.63	0.31	ESL 1	1	
	Staphylococcus chromogenes	27.05	0.32	MAN 87	95	
	Staphylococcus simulans	21.97	0.33		1	
627610	**Good identification**					**627610**
	Staphylococcus piscifermentans	100.0	0.41	BGA 1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
627612	**Low discrimination**							**627612**
Staphylococcus simulans		51.07	0.35	ESL 1				
Staphylococcus piscifermentans		42.67	0.44					
Staphylococcus felis		6.26	0.07					
627613	**Low discrimination**							**627613**
Staphylococcus felis		73.34	0.48	ESL 1				
Staphylococcus simulans		20.22	0.46					
Staphylococcus condimenti		4.07	0.18					
627640	**Good identification**							**627640**
Staphylococcus piscifermentans		100.0	0.48	BGA 1				
627642	**Acceptable identification**							**627642**
Staphylococcus piscifermentans		94.55	0.52	BGA 1				
Staphylococcus simulans		5.45	0.15					
627643	**Low discrimination**							**627643**
					COA	YEP		
Staphylococcus chromogenes		68.68	0.32	ESL 1	1	95		
Staphylococcus simulans		25.06	0.26	GAL 90	1	1		
Staphylococcus intermedius		3.73	0.07		99	1		
627650	**Good identification**							**627650**
Staphylococcus piscifermentans		100.0	0.41	BGA 1				
627652	**Low discrimination**							**627652**
Staphylococcus piscifermentans		65.03	0.44	BGA 1				
Staphylococcus simulans		34.97	0.28					
627653	**Low discrimination**							**627653**
					COA	YEP		
Staphylococcus simulans		71.69	0.38	ESL 1	1	1		
Staphylococcus chromogenes		13.85	0.20		1	95		
Staphylococcus intermedius		7.21	0.16		99	1		
627700	**Good identification**							**627700**
Staphylococcus piscifermentans		100.0	0.52	BGA 1				
627701	**Low discrimination**							**627701**
					YEP	PYR	COA	
Staphylococcus chromogenes		57.79	0.32	ESL 1	95	45	1	
Staphylococcus intermedius		33.70	0.28	LAC 90	1	90	99	
Staphylococcus hyicus		3.96	0.04		1	1	24	
627702	**Good identification**							**627702**
					YEP			
Staphylococcus piscifermentans		93.18	0.55	BGA 1	1			
Staphylococcus chromogenes		2.99	0.10		95			
Staphylococcus felis		1.96	0.01		1			
627703	**Low discrimination**							**627703**
					COA	YEP		
Staphylococcus chromogenes		44.87	0.51	ESL 1	1	95		
Staphylococcus felis		29.39	0.42		1	1		
Staphylococcus intermedius		17.86	0.45		99	1		

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
627710	**Good identification**							**627710**
	Staphylococcus piscifermentans	100.0	0.44	BGA 1				
627711	**Low discrimination**							**627711**
					COA			
	Staphylococcus intermedius	54.15	0.37	ESL 1	99			
	Staphylococcus condimenti	22.48	0.18	LAC 86	1			
	Staphylococcus felis	13.69	0.18		1			
627712	**Low discrimination**							**627712**
	Staphylococcus piscifermentans	49.10	0.48	BGA 1				
	Staphylococcus condimenti	26.42	0.18					
	Staphylococcus felis	16.09	0.18					
627713	**Low discrimination**							**627713**
					COA			
	Staphylococcus condimenti	55.48	0.59	ESL 1	1			
	Staphylococcus felis	33.78	0.59		1			
	Staphylococcus intermedius	8.29	0.54		99			
627740	**Good identification**							**627740**
	Staphylococcus piscifermentans	100.0	0.52	BGA 1				
627741	**Low discrimination**							**627741**
					COA	YEP		
	Staphylococcus chromogenes	52.80	0.32	ESL 1	1	95		
	Staphylococcus intermedius	46.19	0.32	LAC 90	99	1		
	Staphylococcus piscifermentans	1.01	0.11		1	1		
627742	**Very good identification**							**627742**
					COA	YEP		
	Staphylococcus piscifermentans	95.13	0.55	BGA 1	1	1		
	Staphylococcus chromogenes	3.05	0.10		1	95		
	Staphylococcus intermedius	1.82	0.07		99	1		
627743	**Low discrimination**							**627743**
					COA	YEP		
	Staphylococcus chromogenes	62.10	0.51	ESL 1	1	95		
	Staphylococcus intermedius	37.08	0.48		99	1		
	Staphylococcus felis	0.41	0.01		1	1		
627750	**Acceptable identification**							**627750**
					COA	NOV		
	Staphylococcus piscifermentans	93.04	0.44	BGA 1	1	1		
	Staphylococcus saprophyticus subsp. bovis	4.22	0.05		1	99		
	Staphylococcus intermedius	2.74	0.00		99	1		
627751	**Low discrimination**							**627751**
					COA	NOV	YEP	
	Staphylococcus intermedius	88.04	0.41	ESL 1	99	1	1	
	Staphylococcus chromogenes	10.50	0.20	LAC 86	1	1	95	
	Staphylococcus gallinarum	1.46	0.01		1	99	78	
627752	**Low discrimination**							**627752**
					COA			
	Staphylococcus piscifermentans	85.26	0.48	BGA 1	1			
	Staphylococcus intermedius	10.29	0.16		99			
	Staphylococcus simulans	2.66	0.06		1			
627753	**Low discrimination**							**627753**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					COA	PYR	YEP	
Staphylococcus intermedius		73.47	0.57	ESL 1	99	90	1	
Staphylococcus chromogenes		12.84	0.40		1	45	95	
Staphylococcus aureus subsp. aureus		7.10	0.22		99	10	84	
627763	**Low discrimination**							**627763**
					COA	NOV	YEP	
Staphylococcus gallinarum		73.83	0.29	ARG 1	1	99	78	
Staphylococcus lutrae		14.75	0.12	MAN 91	99	1	1	
Staphylococcus chromogenes		7.16	0.10		1	1	95	
627771	**Low discrimination**							**627771**
Staphylococcus gallinarum		84.45	0.42	ARG 1				
Staphylococcus equorum		14.41	0.29					
Staphylococcus xylosus		0.62	-.01					
627773	**Low discrimination**							**627773**
					COA	NOV		
Staphylococcus gallinarum		80.18	0.50	ARG 1	1	99		
Staphylococcus equorum		14.37	0.38		1	99		
Staphylococcus lutrae		3.08	0.18		99	1		
630613	**Good identification**							**630613**
Staphylococcus simulans		100.0	0.27	ORN 1				
630640	**Good identification**							**630640**
Staphylococcus warneri		100.0	0.33	ORN 1				
630650	**Low discrimination**							**630650**
					NOV			
Staphylococcus warneri		56.29	0.30	ORN 1	1			
Staphylococcus saprophyticus subsp. saprophyticus		43.71	0.07		99			
632213	**Low discrimination**							**632213**
Staphylococcus simulans		88.32	0.29	ORN 1				
Staphylococcus capitis subsp. ureolyticus		11.68	-.02	TRE 95				
632413	**Low discrimination**							**632413**
Staphylococcus simulans		87.38	0.27	ORN 1				
Staphylococcus felis		12.62	0.01	SUC 96				
632602	**Good identification**							**632602**
Staphylococcus simulans		100.0	0.32	ORN 1				
632603	**Good identification**							**632603**
Staphylococcus simulans		99.28	0.43	ORN 1				
Staphylococcus felis		0.73	-.10					
632612	**Good identification**							**632612**
Staphylococcus simulans		100.0	0.45	ORN 1				
632613	**Very good identification**							**632613**
Staphylococcus simulans		98.79	0.55	ORN 1				
Staphylococcus felis		1.21	0.07					
632643	**Low discrimination**							**632643**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus simulans		76.54	0.36	ORN 1				
Staphylococcus lugdunensis		21.01	0.20					
Staphylococcus petrasii subsp. petrasii		2.46	0.00					
632652	**Good identification**							**632652**
Staphylococcus simulans		97.95	0.37	ORN 1				
Staphylococcus warneri		2.05	0.12					
632653	**Good identification**							**632653**
Staphylococcus simulans		100.0	0.48	ORN 1				
632713	**Low discrimination**							**632713**
Staphylococcus simulans		38.08	0.34	ORN 1				
Staphylococcus condimenti		29.40	0.18	GAL 8				
Staphylococcus felis		17.91	0.18					
634713	**Low discrimination**							**634713**
					COA			
Staphylococcus intermedius		40.33	0.30	ORN 1	99			
Staphylococcus condimenti		36.20	0.18	NIT 93	1			
Staphylococcus felis		22.05	0.18		1			
634753	**Good identification**							**634753**
					COA	YEP		
Staphylococcus intermedius		97.71	0.34	ORN 1	99	1		
Staphylococcus chromogenes		2.29	-.01	NIT 93	1	95		
636303	**Low discrimination**							**636303**
					YEP	PYR	COA	
Staphylococcus chromogenes		63.10	0.30	ORN 1	95	45	1	
Staphylococcus intermedius		21.74	0.22	TRE 92	1	90	99	
Staphylococcus hyicus		5.53	0.04		1	1	24	
636313	**Low discrimination**							**636313**
					COA			
Staphylococcus intermedius		35.74	0.30	ORN 1	99			
Staphylococcus condimenti		32.08	0.18	TRE 93	1			
Staphylococcus felis		19.54	0.18		1			
636341	**Low discrimination**							**636341**
					COA	YEP	PYR	
Staphylococcus epidermidis		79.97	0.31	VPT 96	1	1	1	
Staphylococcus chromogenes		11.40	0.10	ORN 12	1	95	45	
Staphylococcus intermedius		8.63	0.09		99	1	90	
636342	**Good identification**							**636342**
Staphylococcus epidermidis		100.0	0.36	VPT 96				
636343	**Low discrimination**							**636343**
					COA	YEP	PYR	
Staphylococcus epidermidis		61.95	0.41	VPT 96	1	1	1	
Staphylococcus chromogenes		25.09	0.30	ORN 12	1	95	45	
Staphylococcus intermedius		12.97	0.25		99	1	90	
636353	**Low discrimination**							**636353**
					COA	YEP		
Staphylococcus intermedius		58.19	0.34	ORN 1	99	1		
Staphylococcus delphini		29.02	0.18	TRE 93	99	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus chromogenes		11.75	0.18		1	95		
636413	**Acceptable identification**							**636413**
Staphylococcus felis		90.22	0.42	ORN 1				
Staphylococcus condimenti		6.78	0.14					
Staphylococcus simulans		2.10	0.17					
636503	**Low discrimination**							**636503**
					YEP			
Staphylococcus felis		83.75	0.36	ORN 1	1			
Staphylococcus condimenti		12.58	0.14	MAN 87	1			
Staphylococcus chromogenes		2.62	0.10		95			
636513	**Low discrimination**							**636513**
					COA			
Staphylococcus condimenti		68.83	0.55	ORN 1	1			
Staphylococcus felis		30.97	0.53		1			
Staphylococcus intermedius		0.16	0.12		99			
636603	**Low discrimination**							**636603**
					YEP			
Staphylococcus felis		47.88	0.31	ORN 1	1			
Staphylococcus chromogenes		27.19	0.32	MAN 87	95			
Staphylococcus simulans		22.08	0.33		1			
636612	**Low discrimination**							**636612**
Staphylococcus simulans		89.07	0.35	ORN 1				
Staphylococcus felis		10.93	0.07					
636613	**Low discrimination**							**636613**
Staphylococcus felis		73.34	0.48	ORN 1				
Staphylococcus simulans		20.22	0.46					
Staphylococcus condimenti		4.07	0.18					
636643	**Low discrimination**							**636643**
					COA	YEP		
Staphylococcus chromogenes		69.59	0.32	ORN 1	1	95		
Staphylococcus simulans		25.40	0.26	GAL 90	1	1		
Staphylococcus intermedius		3.78	0.07		99	1		
636652	**Good identification**							**636652**
Staphylococcus simulans		100.0	0.28	ORN 1				
636653	**Low discrimination**							**636653**
					COA	YEP		
Staphylococcus simulans		71.69	0.38	ORN 1	1	1		
Staphylococcus chromogenes		13.85	0.20		1	95		
Staphylococcus intermedius		7.21	0.16		99	1		
636701	**Low discrimination**							**636701**
					YEP	PYR	COA	
Staphylococcus chromogenes		58.44	0.32	ORN 1	95	45	1	
Staphylococcus intermedius		34.08	0.28	LAC 90	1	90	99	
Staphylococcus hyicus		4.01	0.04		1	1	24	
636703	**Low discrimination**							**636703**
					COA	YEP		
Staphylococcus chromogenes		45.02	0.51	ORN 1	1	95		
Staphylococcus felis		29.49	0.42		1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		17.92	0.45		99	1		
636711	**Low discrimination**							**636711**
					COA			
Staphylococcus intermedius		54.15	0.37	ORN 1	99			
Staphylococcus condimenti		22.48	0.18	LAC 86	1			
Staphylococcus felis		13.69	0.18		1			
636713	**Low discrimination**							**636713**
					COA			
Staphylococcus condimenti		55.49	0.59	ORN 1	1			
Staphylococcus felis		33.79	0.59		1			
Staphylococcus intermedius		8.30	0.54		99			
636741	**Low discrimination**							**636741**
					COA	YEP		
Staphylococcus chromogenes		53.34	0.32	ORN 1	1	95		
Staphylococcus intermedius		46.66	0.32	LAC 90	99	1		
636743	**Low discrimination**							**636743**
					COA	YEP		
Staphylococcus chromogenes		62.14	0.51	ORN 1	1	95		
Staphylococcus intermedius		37.10	0.48		99	1		
Staphylococcus felis		0.41	0.01		1	1		
636751	**Low discrimination**							**636751**
					COA	YEP		
Staphylococcus intermedius		89.35	0.41	ORN 1	99	1		
Staphylococcus chromogenes		10.66	0.20	LAC 86	1	95		
636753	**Low discrimination**							**636753**
					COA	PYR	YEP	
Staphylococcus intermedius		73.93	0.57	ORN 1	99	90	1	
Staphylococcus chromogenes		12.92	0.40		1	45	95	
Staphylococcus aureus subsp. aureus		7.14	0.22		99	10	84	
640040	**Good identification**							**640040**
Staphylococcus warneri		100.0	0.46	SUC 95				
640042	**Good identification**							**640042**
Staphylococcus warneri		100.0	0.36	SUC 95				
640050	**Low discrimination**							**640050**
					OXI	NOV		
Staphylococcus warneri		89.39	0.43	SUC 95	1	1		
Micrococcus equipersicus		7.48	0.11	TRE 95	99	99		
Micrococcus bovicus		3.13	0.02		99	99		
640052	**Good identification**							**640052**
Staphylococcus warneri		100.0	0.33	SUC 95				
640140	**Good identification**							**640140**
Staphylococcus warneri		100.0	0.36	SUC 95				
640150	**Low discrimination**							**640150**
					OXI	NOV		
Staphylococcus warneri		89.80	0.33	SUC 95	1	1		
Micrococcus equipersicus		10.20	0.04	TRE 95	99	99		
640200	**Good identification**							**640200**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		96.23	0.46	TRE 95		
Staphylococcus pasteuri		3.77	0.02	MLT 95		
640202	**Low discrimination**					**640202**
					YEP	
Staphylococcus warneri		89.60	0.36	TRE 95	43	
Staphylococcus simulans		6.09	0.02	MLT 95	1	
Staphylococcus pasteuri		4.30	-.06		86	
640210	**Low discrimination**					**640210**
Staphylococcus warneri		75.05	0.43	TRE 95		
Staphylococcus pasteuri		24.95	0.19	MLT 95		
640212	**Low discrimination**					**640212**
					YEP	
Staphylococcus warneri		56.08	0.33	TRE 95	43	
Staphylococcus pasteuri		22.85	0.11	MLT 95	86	
Staphylococcus simulans		21.06	0.15		1	
640240	**Very good identification**					**640240**
					PYR	
Staphylococcus warneri		97.79	0.72	TRE 95	8	
Staphylococcus pasteuri		1.24	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		0.97	0.05		90	
640241	**Good identification**					**640241**
Staphylococcus warneri		100.0	0.38	TRE 95		
640242	**Very good identification**					**640242**
Staphylococcus warneri		98.47	0.62	TRE 95		
Staphylococcus pasteuri		1.53	0.11			
640243	**Low discrimination**					**640243**
Staphylococcus warneri		69.53	0.28	TRE 95		
Staphylococcus simulans		18.34	0.06	MNS 2		
Staphylococcus capitis subsp. ureolyticus		12.13	-.11			
640250	**Good identification**					**640250**
Staphylococcus warneri		90.29	0.69	TRE 95		
Staphylococcus pasteuri		9.71	0.35			
640251	**Low discrimination**					**640251**
Staphylococcus warneri		79.19	0.35	TRE 95		
Staphylococcus capitis subsp. ureolyticus		20.81	0.00	MNS 2		
640252	**Low discrimination**					**640252**
					YEP	
Staphylococcus warneri		87.67	0.60	TRE 95	43	
Staphylococcus pasteuri		11.55	0.27		86	
Staphylococcus simulans		0.78	0.08		1	
640260	**Good identification**					**640260**
Staphylococcus warneri		100.0	0.31	TRE 95		
640270	**Good identification**					**640270**
Staphylococcus warneri		100.0	0.28	TRE 95		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
640300	**Good identification**						**640300**
Staphylococcus warneri		100.0	0.36	TRE 95			
640310	**Good identification**						**640310**
Staphylococcus warneri		100.0	0.33	TRE 95			
640340	**Very good identification**						**640340**
Staphylococcus warneri		100.0	0.62	TRE 95			
640341	**Good identification**						**640341**
Staphylococcus warneri		100.0	0.27	TRE 95			
640342	**Good identification**						**640342**
Staphylococcus warneri		100.0	0.52	TRE 95			
640350	**Very good identification**						**640350**
Staphylococcus warneri		100.0	0.59	TRE 95			
640352	**Good identification**						**640352**
Staphylococcus warneri		100.0	0.49	TRE 95			
640400	**Low discrimination**						**640400**
					YEP		
Staphylococcus warneri		84.57	0.46	SUC 95	43		
Staphylococcus carnosus subsp. utilis		10.78	0.03	MLT 95	10		
Staphylococcus pasteuri		3.32	0.02		86		
640402	**Acceptable identification**						**640402**
					YEP		
Staphylococcus warneri		90.76	0.36	SUC 95	43		
Staphylococcus simulans		4.89	0.00	MLT 95	1		
Staphylococcus pasteuri		4.36	-.06		86		
640410	**Low discrimination**						**640410**
					OXI	NOV	
Staphylococcus warneri		72.15	0.43	SUC 95	1	1	
Staphylococcus pasteuri		23.99	0.19	MLT 95	1	1	
Micrococcus equipersicus		2.23	0.02		99	99	
640412	**Low discrimination**						**640412**
					YEP		
Staphylococcus warneri		58.66	0.33	SUC 95	43		
Staphylococcus pasteuri		23.90	0.11	MLT 95	86		
Staphylococcus simulans		17.44	0.13		1		
640413	**Low discrimination**						**640413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		68.78	0.36	ARG 1	99		
Staphylococcus simulans		31.22	0.24		1		
640440	**Very good identification**						**640440**
					PYR		
Staphylococcus warneri		97.61	0.72	SUC 95	8		
Staphylococcus pasteuri		1.24	0.19		1		
Staphylococcus petrasii subsp. croceilyticus		0.96	0.05		90		
640441	**Acceptable identification**						**640441**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		92.97	0.38	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		7.03	0.10	MNS 2	99		
640442	**Very good identification**						**640442**
Staphylococcus warneri		98.47	0.62	SUC 95			
Staphylococcus pasteuri		1.53	0.11				
640443	**Low discrimination**						**640443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		65.50	0.31	ARG 1	99		
Staphylococcus warneri		28.54	0.28	MAN 91	1		
Staphylococcus simulans		5.96	0.04		1		
640450	**Low discrimination**						**640450**
					OXI	NOV	
Staphylococcus warneri		89.71	0.69	SUC 95	1	1	
Staphylococcus pasteuri		9.65	0.35		1	1	
Macrococcus equipersicus		0.40	0.11		99	99	
640451	**Low discrimination**						**640451**
					NOV		
Staphylococcus cohnii subsp. urealyticum		51.37	0.31	ARG 1	99		
Staphylococcus warneri		48.63	0.35	LAC 91	1		
640452	**Low discrimination**						**640452**
					YEP		
Staphylococcus warneri		86.89	0.60	SUC 95	43		
Staphylococcus pasteuri		11.45	0.27		86		
Staphylococcus simulans		0.61	0.06		1		
640453	**Acceptable identification**						**640453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.70	0.52	ARG 1	99		
Staphylococcus simulans		3.37	0.16		1		
Staphylococcus warneri		2.92	0.25		1		
640460	**Good identification**						**640460**
Staphylococcus warneri		100.0	0.31	SUC 95			
640470	**Good identification**						**640470**
Staphylococcus warneri		100.0	0.28	SUC 95			
640500	**Good identification**						**640500**
Staphylococcus warneri		100.0	0.36	SUC 95			
640510	**Good identification**						**640510**
Staphylococcus warneri		100.0	0.33	SUC 95			
640513	**Acceptable identification**						**640513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		91.57	0.29	ARG 1	99		
Staphylococcus simulans		8.43	0.02		1		
640540	**Very good identification**						**640540**
Staphylococcus warneri		100.0	0.62	SUC 95			
640541	**Good identification**						**640541**
Staphylococcus warneri		100.0	0.27	SUC 95			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
640542	**Good identification**						**640542**
Staphylococcus warneri		100.0	0.52	SUC 95			
640550	**Very good identification**						**640550**
					OXI	NOV	
Staphylococcus warneri		99.04	0.59	SUC 95	1	1	
Macrococcus equipersicus		0.59	0.04		99	99	
Macrococcus bovicus		0.37	-.02		99	99	
640552	**Good identification**						**640552**
Staphylococcus warneri		100.0	0.49	SUC 95			
640553	**Good identification**						**640553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		97.75	0.44	ARG 1	99		
Staphylococcus warneri		2.25	0.14		1		
640600	**Low discrimination**						**640600**
					PYR		
Staphylococcus warneri		82.36	0.72	MLT 95	8		
Staphylococcus pasteuri		16.82	0.43		1		
Staphylococcus petrasii subsp. croceilyticus		0.81	0.05		90		
640601	**Low discrimination**						**640601**
					YEP		
Staphylococcus warneri		87.09	0.38	MLT 95	43		
Staphylococcus pasteuri		8.80	0.02	MNS 2	86		
Staphylococcus hyicus		4.11	-.07		1		
640602	**Low discrimination**						**640602**
					YEP		
Staphylococcus warneri		75.86	0.62	MLT 95	43		
Staphylococcus pasteuri		18.99	0.35		86		
Staphylococcus simulans		5.16	0.29		1		
640603	**Low discrimination**						**640603**
Staphylococcus simulans		80.21	0.40	BGA 90			
Staphylococcus hyicus		11.71	0.15	NIT 96			
Staphylococcus warneri		7.19	0.28				
640610	**Low discrimination**						**640610**
					YEP		
Staphylococcus pasteuri		63.38	0.60	VPT 86	86		
Staphylococcus warneri		36.58	0.69	MLT 86	43		
Staphylococcus simulans		0.05	0.00		1		
640611	**Low discrimination**						**640611**
					YEP		
Staphylococcus warneri		48.43	0.35	MLT 95	43		
Staphylococcus pasteuri		41.53	0.19	MNS 2	86		
Staphylococcus simulans		10.05	0.11		1		
640612	**Low discrimination**						**640612**
					YEP		
Staphylococcus pasteuri		60.68	0.52	VPT 86	86		
Staphylococcus warneri		28.58	0.60	MLT 86	43		
Staphylococcus simulans		10.74	0.41		1		
640613	**Good identification**						**640613**
					YEP		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		96.51	0.52	BGA 90	1	
Staphylococcus pasteuri		1.65	0.11	NIT 96	86	
Staphylococcus warneri		1.57	0.25		43	
640620	**Low discrimination**					**640620**
Staphylococcus warneri		83.04	0.31	XYL 1		
Staphylococcus pasteuri		16.96	0.02	MLT 95		
640640	**Low discrimination**					**640640**
					PYR	
Staphylococcus warneri		89.48	0.99		8	
Staphylococcus pasteuri		5.91	0.60		1	
Staphylococcus petrasii subsp. croceilyticus		4.61	0.47		90	
640641	**Good identification**					**640641**
					PYR	
Staphylococcus warneri		94.20	0.64	MNS 2	8	
Staphylococcus pasteuri		3.08	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		2.40	0.05		90	
640642	**Good identification**					**640642**
					YEP	
Staphylococcus warneri		92.24	0.89		43	
Staphylococcus pasteuri		7.46	0.52		86	
Staphylococcus simulans		0.15	0.22		1	
640643	**Low discrimination**					**640643**
					YEP	
Staphylococcus warneri		76.70	0.54	MNS 2	43	
Staphylococcus simulans		20.23	0.32		1	
Staphylococcus pasteuri		3.07	0.11		86	
640650	**Low discrimination**					**640650**
					PYR	
Staphylococcus warneri		64.06	0.96		8	
Staphylococcus pasteuri		35.88	0.76		1	
Staphylococcus petrasii subsp. croceilyticus		0.05	0.05		90	
640651	**Low discrimination**					**640651**
					YEP	
Staphylococcus warneri		77.99	0.61	MNS 2	43	
Staphylococcus pasteuri		21.62	0.35		86	
Staphylococcus simulans		0.38	0.04		1	
640652	**Low discrimination**					**640652**
					YEP	
Staphylococcus warneri		58.88	0.86		43	
Staphylococcus pasteuri		40.42	0.68		86	
Staphylococcus simulans		0.52	0.34		1	
640653	**Low discrimination**					**640653**
					YEP	
Staphylococcus simulans		51.14	0.45	BGA 90	1	
Staphylococcus warneri		35.10	0.51	NIT 96	43	
Staphylococcus pasteuri		11.93	0.27		86	
640660	**Low discrimination**					**640660**
					PYR	
Staphylococcus warneri		89.48	0.58	XYL 1	8	
Staphylococcus pasteuri		5.91	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		4.61	0.05		90	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
640662	**Acceptable identification**					**640662**
Staphylococcus warneri		92.51	0.48	XYL 1		
Staphylococcus pasteuri		7.49	0.11			
640670	**Low discrimination**					**640670**
Staphylococcus warneri		64.10	0.55	XYL 1		
Staphylococcus pasteuri		35.90	0.35			
640672	**Low discrimination**					**640672**
Staphylococcus warneri		59.30	0.45	XYL 1		
Staphylococcus pasteuri		40.70	0.27			
640700	**Very good identification**					**640700**
Staphylococcus warneri		99.35	0.62	MLT 95		
Staphylococcus pasteuri		0.65	0.02			
640702	**Good identification**					**640702**
Staphylococcus warneri		96.25	0.52	MLT 95		
Staphylococcus simulans		1.80	0.07			
Staphylococcus hyicus		1.18	-.04			
640703	**Acceptable identification**					**640703**
Staphylococcus hyicus		93.58	0.37	NIT 99		
Staphylococcus simulans		4.85	0.18	PHS 92		
Staphylococcus warneri		1.58	0.17			
640710	**Good identification**					**640710**
Staphylococcus warneri		94.75	0.59	MLT 95		
Staphylococcus pasteuri		5.25	0.19			
640712	**Low discrimination**					**640712**
					YEP	
Staphylococcus warneri		85.37	0.49	MLT 95	43	
Staphylococcus simulans		8.83	0.19		1	
Staphylococcus pasteuri		5.80	0.11		86	
640713	**Acceptable identification**					**640713**
Staphylococcus simulans		90.28	0.30	BGA 90		
Staphylococcus warneri		5.32	0.14	NIT 96		
Staphylococcus hyicus		4.40	-.04	GAL 8		
640740	**Very good identification**					**640740**
					PYR	
Staphylococcus warneri		99.63	0.88		8	
Staphylococcus pasteuri		0.21	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		0.16	0.05		90	
640741	**Very good identification**					**640741**
					PYR	
Staphylococcus warneri		99.29	0.54	MNS 2	8	
Staphylococcus petrasii subsp. petrasii		0.71	-.05		99	
640742	**Very good identification**					**640742**
					YEP	
Staphylococcus warneri		99.70	0.78		43	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		0.26	0.11		86		
Staphylococcus simulans		0.04	0.00		1		
640743	**Acceptable identification**						**640743**
Staphylococcus warneri		90.57	0.44	MNS 2			
Staphylococcus simulans		6.58	0.11				
Staphylococcus hyicus		2.86	-.04				
640750	**Very good identification**						**640750**
					NOV	PYR	
Staphylococcus warneri		98.22	0.85		1	8	
Staphylococcus pasteuri		1.76	0.35		1	1	
Staphylococcus saprophyticus subsp. bovis		0.03	0.02		99	99	
640751	**Good identification**						**640751**
					NOV		
Staphylococcus warneri		98.96	0.51	MNS 2	1		
Staphylococcus xylosus		1.04	-.03		99		
640752	**Very good identification**						**640752**
					YEP		
Staphylococcus warneri		97.54	0.76		43		
Staphylococcus pasteuri		2.14	0.27		86		
Staphylococcus simulans		0.24	0.12		1		
640753	**Low discrimination**						**640753**
					NOV		
Staphylococcus warneri		64.03	0.41	MNS 2	1		
Staphylococcus simulans		25.69	0.23		1		
Staphylococcus xylosus		5.76	0.06		99		
640760	**Good identification**						**640760**
Staphylococcus warneri		100.0	0.47	XYL 1			
640762	**Good identification**						**640762**
Staphylococcus warneri		100.0	0.37	XYL 1			
640770	**Good identification**						**640770**
Staphylococcus warneri		100.0	0.44	XYL 1			
640772	**Low discrimination**						**640772**
					NOV		
Staphylococcus warneri		89.33	0.35	XYL 1	1		
Staphylococcus arlettae		10.67	0.12		99		
640773	**Acceptable identification**						**640773**
Staphylococcus xylosus		92.96	0.28	ARG 1			
Staphylococcus arlettae		7.04	0.14	NIT 89			
641240	**Good identification**						**641240**
Staphylococcus warneri		100.0	0.31	ESL 1			
641250	**Good identification**						**641250**
Staphylococcus warneri		100.0	0.28	ESL 1			
641413	**Good identification**						**641413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ARG 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
641440	**Good identification**						**641440**
Staphylococcus warneri		100.0	0.31	ESL 1			
641450	**Low discrimination**						**641450**
					OXI	NOV	
Staphylococcus warneri		51.90	0.28	ESL 1	1	1	
Macrococcus equipersicus		48.10	0.18	SUC 95	99	99	
641453	**Good identification**						**641453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ARG 1			
641553	**Good identification**						**641553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.36	ARG 1			
641600	**Low discrimination**						**641600**
					YEP		
Staphylococcus warneri		77.52	0.31	ESL 1	43		
Staphylococcus pasteuri		15.83	0.02	MLT 95	86		
Staphylococcus piscifermentans		6.64	0.11		1		
641640	**Low discrimination**						**641640**
					PYR		
Staphylococcus warneri		89.13	0.58	ESL 1	8		
Staphylococcus pasteuri		5.89	0.19		1		
Staphylococcus petrasii subsp. croceilyticus		4.59	0.05		90		
641642	**Acceptable identification**						**641642**
					YEP		
Staphylococcus warneri		90.81	0.48	ESL 1	43		
Staphylococcus pasteuri		7.35	0.11		86		
Staphylococcus piscifermentans		1.84	0.14		1		
641650	**Low discrimination**						**641650**
Staphylococcus warneri		64.10	0.55	ESL 1			
Staphylococcus pasteuri		35.90	0.35				
641652	**Low discrimination**						**641652**
Staphylococcus warneri		59.30	0.45	ESL 1			
Staphylococcus pasteuri		40.70	0.27				
641740	**Good identification**						**641740**
Staphylococcus warneri		97.90	0.47	ESL 1			
Staphylococcus piscifermentans		2.10	0.14				
641742	**Acceptable identification**						**641742**
Staphylococcus warneri		91.21	0.37	ESL 1			
Staphylococcus piscifermentans		8.79	0.18				
641750	**Good identification**						**641750**
Staphylococcus warneri		100.0	0.44	ESL 1			
641752	**Acceptable identification**						**641752**
Staphylococcus warneri		94.60	0.35	ESL 1			
Staphylococcus piscifermentans		5.40	0.10				
641773	**Low discrimination**						**641773**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					YEP	
Staphylococcus equorum		83.13	0.34	ARG 1	1	
Staphylococcus xylosus		12.95	0.15	NIT 92	40	
Staphylococcus arlettae		3.92	0.14		99	
642040	**Acceptable identification**					**642040**
Staphylococcus warneri		93.67	0.38	SUC 95		
Staphylococcus auricularis		6.33	0.06	TRE 95		
642042	**Good identification**					**642042**
Staphylococcus warneri		100.0	0.29	SUC 95		
642050	**Low discrimination**					**642050**
Staphylococcus warneri		89.46	0.36	SUC 95		
Staphylococcus pasteuri		10.54	0.02	TRE 95		
642140	**Low discrimination**					**642140**
Staphylococcus warneri		86.64	0.28	SUC 95		
Kocuria varians		13.36	0.04	TRE 95		
642200	**Low discrimination**					**642200**
Staphylococcus warneri		81.70	0.38	TRE 95		
Staphylococcus pasteuri		18.30	0.10	MLT 95		
642202	**Low discrimination**					**642202**
					YEP	
Staphylococcus simulans		73.08	0.31	BGA 90	1	
Staphylococcus warneri		19.19	0.29	TRE 95	43	
Staphylococcus pasteuri		5.27	0.02		86	
642203	**Low discrimination**					**642203**
Staphylococcus simulans		49.88	0.42	BGA 90		
Staphylococcus hyicus		49.63	0.35	TRE 95		
Staphylococcus capitis subsp. ureolyticus		0.49	-.13			
642210	**Low discrimination**					**642210**
					YEP	
Staphylococcus pasteuri		63.95	0.27	VPT 86	86	
Staphylococcus warneri		33.67	0.36	TRE 99	43	
Staphylococcus simulans		2.38	0.02	MLT 86	1	
642212	**Low discrimination**					**642212**
					YEP	
Staphylococcus simulans		86.34	0.43	BGA 90	1	
Staphylococcus pasteuri		9.56	0.19	TRE 95	86	
Staphylococcus warneri		4.11	0.26		43	
642213	**Good identification**					**642213**
Staphylococcus simulans		94.22	0.54	BGA 90		
Staphylococcus capitis subsp. ureolyticus		5.54	0.16	TRE 95		
Staphylococcus hyicus		0.24	-.07			
642240	**Low discrimination**					**642240**
					PYR	
Staphylococcus warneri		83.17	0.65	TRE 95	8	
Staphylococcus petrasii subsp. croceilyticus		7.67	0.18		90	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus pasteuri		6.02	0.27		1	
642241	**Low discrimination**					**642241**
Staphylococcus warneri		74.00	0.30	TRE 95		
Staphylococcus capitis subsp. ureolyticus		26.01	-.02	MNS 2		
642242	**Low discrimination**					**642242**
Staphylococcus warneri		77.30	0.55	TRE 95		
Staphylococcus hominis subsp. hominis		8.41	0.29			
Staphylococcus simulans		6.96	0.24			
642243	**Low discrimination**					**642243**
Staphylococcus simulans		61.26	0.35	BGA 90		
Staphylococcus capitis subsp. ureolyticus		32.07	0.16	TRE 95		
Staphylococcus warneri		4.15	0.20			
642250	**Low discrimination**					**642250**
Staphylococcus warneri		61.60	0.62	TRE 95		
Staphylococcus pasteuri		37.83	0.43			
Staphylococcus hominis subsp. hominis		0.39	0.11			
642252	**Low discrimination**					**642252**
					YEP	
Staphylococcus warneri		43.01	0.52	TRE 95	43	
Staphylococcus pasteuri		32.37	0.35		86	
Staphylococcus simulans		21.39	0.36		1	
642253	**Low discrimination**					**642253**
					COA	
Staphylococcus capitis subsp. ureolyticus		73.10	0.44	GLR 1	1	
Staphylococcus simulans		23.27	0.47		1	
Staphylococcus delphini		3.34	0.18		99	
642300	**Low discrimination**					**642300**
Staphylococcus warneri		88.09	0.28	TRE 95		
Staphylococcus hyicus		11.91	-.07	MLT 95		
642301	**Good identification**					**642301**
Staphylococcus hyicus		100.0	0.35	PHS 92		
642303	**Very good identification**					**642303**
					YEP	
Staphylococcus hyicus		99.17	0.56	PHS 92	1	
Staphylococcus simulans		0.75	0.20	TRE 92	1	
Staphylococcus chromogenes		0.08	-.04		95	
642313	**Low discrimination**					**642313**
Staphylococcus simulans		75.06	0.32	BGA 90		
Staphylococcus hyicus		24.94	0.15	GAL 8		
642340	**Low discrimination**					**642340**
					PYR	
Staphylococcus warneri		88.39	0.54	TRE 95	8	
Staphylococcus hominis subsp. hominis		9.37	0.29		1	
Staphylococcus haemolyticus		1.14	0.08		98	
642342	**Low discrimination**					**642342**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		66.91	0.45	TRE 95		
Staphylococcus hominis subsp. hominis		24.98	0.30			
Staphylococcus epidermidis		3.65	0.10			
642350	**Acceptable identification**					**642350**
Staphylococcus warneri		94.05	0.52	TRE 95		
Staphylococcus hominis subsp. hominis		2.06	0.12			
Staphylococcus pasteuri		1.85	0.02			
642352	**Low discrimination**					**642352**
Staphylococcus warneri		82.37	0.42	TRE 95		
Staphylococcus simulans		11.28	0.14			
Staphylococcus hominis subsp. hominis		6.35	0.13			
642402	**Low discrimination**					**642402**
Staphylococcus simulans		53.20	0.29	BGA 90		
Staphylococcus piscifermentans		22.35	0.32	SUC 96		
Staphylococcus warneri		17.65	0.29			
642403	**Low discrimination**					**642403**
Staphylococcus simulans		87.26	0.40	BGA 90		
Staphylococcus hyicus		12.74	0.15	SUC 96		
642410	**Low discrimination**					**642410**
					YEP	
Staphylococcus pasteuri		60.84	0.27	VPT 86	86	
Staphylococcus warneri		32.03	0.36	SUC 99	43	
Staphylococcus piscifermentans		5.33	0.21	MLT 86	1	
642412	**Low discrimination**					**642412**
					YEP	
Staphylococcus simulans		80.33	0.41	BGA 90	1	
Staphylococcus pasteuri		11.23	0.19	SUC 96	86	
Staphylococcus warneri		4.82	0.26		43	
642413	**Good identification**					**642413**
					NOV	
Staphylococcus simulans		98.41	0.52	BGA 90	1	
Staphylococcus cohnii subsp. urealyticum		1.59	0.21	SUC 96	99	
642440	**Low discrimination**					**642440**
					PYR	
Staphylococcus warneri		81.94	0.65	SUC 95	8	
Staphylococcus petrasii subsp. croceilyticus		7.56	0.18		90	
Staphylococcus pasteuri		5.93	0.27		1	
642441	**Low discrimination**					**642441**
Staphylococcus warneri		81.78	0.30	SUC 95		
Staphylococcus auricularis		18.22	0.08	MNS 2		
642442	**Low discrimination**					**642442**
					YEP	
Staphylococcus warneri		80.63	0.55	SUC 95	43	
Staphylococcus pasteuri		7.15	0.19		86	
Staphylococcus simulans		5.75	0.22		1	
642443	**Low discrimination**					**642443**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus simulans		85.74	0.32	BGA 90	1		
Staphylococcus warneri		7.33	0.20	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		6.93	0.16		99		
642450	**Low discrimination**						**642450**
					NOV	PYR	
Staphylococcus warneri		60.79	0.62	SUC 95	1	8	
Staphylococcus pasteuri		37.33	0.43		1	1	
Staphylococcus saprophyticus subsp. bovis		1.35	0.18		99	99	
642451	**Low discrimination**						**642451**
					NOV		
Staphylococcus warneri		44.89	0.27	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		19.52	0.16	MNS 2	99		
Staphylococcus pasteuri		13.64	0.02		1		
642452	**Low discrimination**						**642452**
					YEP		
Staphylococcus warneri		45.41	0.52	SUC 95	43		
Staphylococcus pasteuri		34.17	0.35		86		
Staphylococcus simulans		17.88	0.34		1		
642453	**Low discrimination**						**642453**
					NOV		
Staphylococcus simulans		81.98	0.45	BGA 90	1		
Staphylococcus cohnii subsp. urealyticum		16.75	0.36	SUC 96	99		
Staphylococcus warneri		1.27	0.17		1		
642500	**Low discrimination**						**642500**
Staphylococcus piscifermentans		54.47	0.32	GLR 1			
Staphylococcus warneri		40.75	0.28	ESL 90			
Staphylococcus auricularis		4.78	0.01				
642502	**Low discrimination**						**642502**
Staphylococcus piscifermentans		71.72	0.36	GLR 1			
Staphylococcus warneri		11.92	0.18	ESL 90			
Staphylococcus simulans		9.90	0.07				
642503	**Good identification**						**642503**
Staphylococcus hyicus		95.08	0.37	PHS 92			
Staphylococcus simulans		4.93	0.18	SUC 99			
642513	**Low discrimination**						**642513**
					NOV		
Staphylococcus simulans		84.17	0.30	BGA 90	1		
Staphylococcus cohnii subsp. urealyticum		6.72	0.13	GAL 8	99		
Staphylococcus felis		5.00	-.04	SUC 96	1		
642540	**Good identification**						**642540**
Staphylococcus warneri		92.19	0.54	SUC 95			
Staphylococcus piscifermentans		6.49	0.32				
Staphylococcus auricularis		1.33	0.08				
642542	**Low discrimination**						**642542**
Staphylococcus warneri		74.84	0.45	SUC 95			
Staphylococcus piscifermentans		23.69	0.36				

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		1.47	0.00				
642550	**Low discrimination**						**642550**
					NOV	PYR	
Staphylococcus warneri		74.52	0.52	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		20.92	0.31		99	99	
Staphylococcus piscifermentans		3.10	0.25		1	15	
642552	**Low discrimination**						**642552**
					NOV	PYR	
Staphylococcus warneri		66.39	0.42	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		13.98	0.18		99	99	
Staphylococcus piscifermentans		12.44	0.28		1	15	
642553	**Low discrimination**						**642553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		44.80	0.29	ARG 1	99		
Staphylococcus simulans		44.50	0.23		1		
Staphylococcus saprophyticus subsp. bovis		6.45	0.06		99		
642600	**Low discrimination**						**642600**
					PYR		
Staphylococcus pasteuri		50.46	0.51	VPT 86	1		
Staphylococcus warneri		43.24	0.65	MAN 86	8		
Staphylococcus petrasii subsp. croceilyticus		3.99	0.18	MLT 86	90		
642601	**Low discrimination**						**642601**
Staphylococcus hyicus		74.76	0.35	PHS 92			
Staphylococcus simulans		14.42	0.27	GAL 92			
Staphylococcus warneri		6.86	0.30	LAC 92			
642602	**Low discrimination**						**642602**
					YEP		
Staphylococcus simulans		59.36	0.57	BGA 90	1		
Staphylococcus pasteuri		22.29	0.43		86		
Staphylococcus warneri		15.59	0.55		43		
642603	**Low discrimination**						**642603**
Staphylococcus simulans		62.29	0.68	BGA 90			
Staphylococcus hyicus		37.51	0.56				
Staphylococcus warneri		0.10	0.20				
642610	**Good identification**						**642610**
					YEP		
Staphylococcus pasteuri		90.13	0.68	VPT 86	86		
Staphylococcus warneri		9.11	0.62	MLT 86	43		
Staphylococcus simulans		0.65	0.29		1		
642611	**Low discrimination**						**642611**
					YEP		
Staphylococcus simulans		65.75	0.39	BGA 90	1		
Staphylococcus pasteuri		27.73	0.27	LAC 99	86		
Staphylococcus warneri		5.66	0.27		43		
642612	**Low discrimination**						**642612**
					YEP		
Staphylococcus simulans		61.46	0.70	BGA 90	1		
Staphylococcus pasteuri		35.44	0.60		86		
Staphylococcus warneri		2.92	0.52		43		
642613	**Very good identification**						**642613**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					YEP	
Staphylococcus simulans		99.63	0.80	BGA 90	1	
Staphylococcus pasteuri		0.17	0.19		86	
Staphylococcus hyicus		0.15	0.15		1	
642623	**Low discrimination**					**642623**
Staphylococcus simulans		62.42	0.27	BGA 90		
Staphylococcus hyicus		37.58	0.15	XYL 1		
642630	**Acceptable identification**					**642630**
Staphylococcus pasteuri		90.82	0.27	VPT 86		
Staphylococcus warneri		9.18	0.21	XYL 1		
642632	**Low discrimination**					**642632**
					YEP	
Staphylococcus simulans		61.57	0.29	BGA 90	1	
Staphylococcus pasteuri		35.50	0.19	XYL 1	86	
Staphylococcus warneri		2.93	0.11		43	
642633	**Good identification**					**642633**
					NOV	
Staphylococcus simulans		97.56	0.39	BGA 90	1	
Staphylococcus xylosus		2.44	0.03	XYL 1	99	
642640	**Low discrimination**					**642640**
					PYR	
Staphylococcus warneri		53.50	0.91		8	
Staphylococcus petrasii subsp. croceilyticus		25.71	0.59		90	
Staphylococcus pasteuri		20.18	0.68		1	
642641	**Low discrimination**					**642641**
					PYR	
Staphylococcus warneri		63.81	0.56	MNS 2	8	
Staphylococcus petrasii subsp. croceilyticus		15.18	0.18		90	
Staphylococcus pasteuri		11.91	0.27		1	
642642	**Low discrimination**					**642642**
					YEP	
Staphylococcus warneri		62.69	0.81		43	
Staphylococcus pasteuri		28.98	0.60		86	
Staphylococcus simulans		5.64	0.50		1	
642643	**Good identification**					**642643**
					YEP	
Staphylococcus simulans		90.33	0.61	BGA 90	1	
Staphylococcus warneri		6.12	0.46		43	
Staphylococcus pasteuri		1.40	0.19		86	
642650	**Low discrimination**					**642650**
					PYR	
Staphylococcus pasteuri		75.89	0.84	VPT 86	1	
Staphylococcus warneri		23.71	0.88		8	
Staphylococcus petrasii subsp. croceilyticus		0.16	0.18		90	
642651	**Low discrimination**					**642651**
					YEP	
Staphylococcus pasteuri		54.12	0.43	VPT 86	86	
Staphylococcus warneri		34.17	0.53	MNS 1	43	
Staphylococcus simulans		9.39	0.32		1	
642652	**Low discrimination**					**642652**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					YEP	
Staphylococcus pasteuri		72.18	0.76	VPT 86	86	
Staphylococcus warneri		18.41	0.78		43	
Staphylococcus simulans		9.16	0.62		1	
642653	**Very good identification**					**642653**
					YEP	
Staphylococcus simulans		96.23	0.73	BGA 90	1	
Staphylococcus pasteuri		2.29	0.35		86	
Staphylococcus warneri		1.18	0.44		43	
642660	**Low discrimination**					**642660**
					PYR	
Staphylococcus warneri		53.83	0.50	XYL 1	8	
Staphylococcus petrasii subsp. croceilyticus		25.87	0.18		90	
Staphylococcus pasteuri		20.31	0.27		1	
642662	**Low discrimination**					**642662**
					YEP	
Staphylococcus warneri		64.42	0.40	XYL 1	43	
Staphylococcus pasteuri		29.78	0.19		86	
Staphylococcus simulans		5.80	0.09		1	
642670	**Low discrimination**					**642670**
Staphylococcus pasteuri		76.19	0.43	VPT 86		
Staphylococcus warneri		23.81	0.47	XYL 1		
642672	**Low discrimination**					**642672**
					YEP	
Staphylococcus pasteuri		72.37	0.35	VPT 86	86	
Staphylococcus warneri		18.46	0.37	XYL 1	43	
Staphylococcus simulans		9.18	0.21		1	
642673	**Low discrimination**					**642673**
					NOV	
Staphylococcus simulans		57.47	0.32	BGA 90	1	
Staphylococcus xylosus		42.53	0.26	XYL 1	99	
642700	**Low discrimination**					**642700**
Staphylococcus warneri		81.65	0.54	MLT 95		
Staphylococcus piscifermentans		8.62	0.36			
Staphylococcus hyicus		6.68	0.15			
642701	**Very good identification**					**642701**
Staphylococcus hyicus		99.51	0.56	PHS 92		
Staphylococcus warneri		0.25	0.20	LAC 92		
Staphylococcus simulans		0.15	0.05			
642702	**Low discrimination**					**642702**
Staphylococcus hyicus		52.36	0.37	PHS 92		
Staphylococcus simulans		19.44	0.35	MNS 99		
Staphylococcus warneri		18.54	0.45			
642703	**Very good identification**					**642703**
					YEP	
Staphylococcus hyicus		98.68	0.78	PHS 92	1	
Staphylococcus simulans		1.24	0.46		1	
Staphylococcus chromogenes		0.08	0.18		95	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
642710	**Low discrimination**						**642710**
					YEP		
Staphylococcus warneri		70.93	0.52	MLT 95	43		
Staphylococcus pasteuri		22.46	0.27		86		
Staphylococcus piscifermentans		4.43	0.28		1		
642712	**Low discrimination**						**642712**
					YEP		
Staphylococcus simulans		77.22	0.48	BGA 90	1		
Staphylococcus warneri		13.33	0.42	GAL 8	43		
Staphylococcus pasteuri		5.17	0.19		86		
642713	**Low discrimination**						**642713**
Staphylococcus simulans		82.44	0.59	BGA 90			
Staphylococcus hyicus		16.58	0.37	GAL 8			
Staphylococcus felis		0.42	0.02				
642723	**Good identification**						**642723**
					NOV		
Staphylococcus hyicus		97.86	0.37	PHS 92	1		
Staphylococcus simulans		1.23	0.05	XYL 1	1		
Staphylococcus xylosus		0.91	-.01		99		
642740	**Good identification**						**642740**
					PYR		
Staphylococcus warneri		94.24	0.81		8		
Staphylococcus hominis subsp. hominis		1.67	0.39		1		
Staphylococcus petrasii subsp. croceilyticus		1.45	0.18		90		
642741	**Low discrimination**						**642741**
					PYR		
Staphylococcus warneri		71.47	0.46	MNS 2	8		
Staphylococcus hyicus		15.09	0.15		1		
Staphylococcus petrasii subsp. petrasii		10.73	0.14		99		
642742	**Low discrimination**						**642742**
Staphylococcus warneri		87.45	0.71				
Staphylococcus hominis subsp. hominis		5.44	0.40				
Staphylococcus piscifermentans		2.19	0.39				
642743	**Low discrimination**						**642743**
Staphylococcus hyicus		54.83	0.37	PHS 92			
Staphylococcus simulans		30.62	0.39	MLT 1			
Staphylococcus warneri		7.53	0.36				
642750	**Low discrimination**						**642750**
					NOV	PYR	
Staphylococcus warneri		85.02	0.78		1	8	
Staphylococcus pasteuri		8.70	0.43		1	1	
Staphylococcus saprophyticus subsp. bovis		5.03	0.43		99	99	
642751	**Low discrimination**						**642751**
					NOV	PYR	
Staphylococcus warneri		48.26	0.43	MNS 2	1	8	
Staphylococcus saprophyticus subsp. bovis		34.94	0.31		99	99	
Staphylococcus xylosus		9.60	0.15		99	65	
642752	**Low discrimination**						**642752**
					YEP		

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		74.26	0.68		43		
Staphylococcus simulans		10.17	0.41		1		
Staphylococcus pasteuri		9.31	0.35		86		
642753	**Low discrimination**						**642753**
					NOV		
Staphylococcus simulans		84.85	0.51	BGA 90	1		
Staphylococcus xylosus		6.41	0.25	GAL 8	99		
Staphylococcus warneri		3.78	0.33		1		
642760	**Good identification**						**642760**
Staphylococcus warneri		100.0	0.40	XYL 1			
642762	**Good identification**						**642762**
Staphylococcus warneri		100.0	0.30	XYL 1			
642770	**Low discrimination**						**642770**
					NOV	PYR	
Staphylococcus warneri		82.77	0.37	XYL 1	1	8	
Staphylococcus pasteuri		8.47	0.02		1	1	
Staphylococcus saprophyticus subsp. bovis		4.89	0.02		99	99	
642771	**Good identification**						**642771**
Staphylococcus xylosus		97.51	0.37	ARG 1			
Staphylococcus equorum		2.49	0.06				
642772	**Low discrimination**						**642772**
					NOV		
Staphylococcus warneri		65.13	0.27	XYL 1	1		
Staphylococcus xylosus		25.95	0.05		99		
Staphylococcus simulans		8.92	-.01		1		
642773	**Good identification**						**642773**
					NOV		
Staphylococcus xylosus		96.65	0.47	ARG 1	99		
Staphylococcus equorum		2.23	0.14		99		
Staphylococcus simulans		1.12	0.10		1		
643400	**Good identification**						**643400**
Staphylococcus piscifermentans		100.0	0.48	GLR 1			
643402	**Good identification**						**643402**
Staphylococcus piscifermentans		100.0	0.52	GLR 1			
643410	**Good identification**						**643410**
Staphylococcus piscifermentans		100.0	0.41	GLR 1			
643412	**Good identification**						**643412**
Staphylococcus piscifermentans		97.57	0.44	GLR 1			
Staphylococcus simulans		2.43	0.00				
643440	**Acceptable identification**						**643440**
Staphylococcus piscifermentans		92.96	0.48	GLR 1			
Staphylococcus warneri		7.04	0.24				
643442	**Good identification**						**643442**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		98.34	0.52	GLR 1		
Staphylococcus warneri		1.66	0.14			
643450	**Low discrimination**					**643450**
					YEP	
Staphylococcus piscifermentans		82.87	0.41	GLR 1	1	
Staphylococcus warneri		10.61	0.21		43	
Staphylococcus pasteuri		6.52	0.02		86	
643452	**Good identification**					**643452**
Staphylococcus piscifermentans		97.23	0.44	GLR 1		
Staphylococcus warneri		2.77	0.11			
643453	**Low discrimination**					**643453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		88.72	0.28	ARG 1	99	
Staphylococcus simulans		11.28	0.04		1	
643500	**Good identification**					**643500**
Staphylococcus piscifermentans		100.0	0.52	GLR 1		
643502	**Very good identification**					**643502**
Staphylococcus piscifermentans		100.0	0.55	GLR 1		
643510	**Good identification**					**643510**
Staphylococcus piscifermentans		100.0	0.44	GLR 1		
643512	**Good identification**					**643512**
Staphylococcus piscifermentans		100.0	0.48	GLR 1		
643540	**Good identification**					**643540**
Staphylococcus piscifermentans		98.43	0.52	GLR 1		
Staphylococcus warneri		1.57	0.13			
643542	**Very good identification**					**643542**
Staphylococcus piscifermentans		100.0	0.55	GLR 1		
643550	**Good identification**					**643550**
Staphylococcus piscifermentans		97.38	0.44	GLR 1		
Staphylococcus warneri		2.63	0.10			
643552	**Good identification**					**643552**
Staphylococcus piscifermentans		100.0	0.48	GLR 1		
643600	**Acceptable identification**					**643600**
					YEP	
Staphylococcus piscifermentans		90.13	0.52	GLR 1	1	
Staphylococcus pasteuri		5.31	0.10		86	
Staphylococcus warneri		4.55	0.24		43	
643602	**Good identification**					**643602**
					YEP	
Staphylococcus piscifermentans		93.46	0.55	GLR 1	1	
Staphylococcus simulans		4.00	0.16		1	
Staphylococcus pasteuri		1.50	0.02		86	
643603	**Low discrimination**					**643603**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		59.78	0.27	BGA 90			
Staphylococcus hyicus		36.00	0.15	ESL 1			
Staphylococcus piscifermentans		4.22	0.14				
643610	**Low discrimination**						**643610**
					YEP		
Staphylococcus piscifermentans		51.81	0.44	GLR 1	1		
Staphylococcus pasteuri		43.77	0.27		86		
Staphylococcus warneri		4.42	0.21		43		
643612	**Low discrimination**						**643612**
					YEP		
Staphylococcus piscifermentans		60.68	0.48	GLR 1	1		
Staphylococcus simulans		24.21	0.29		1		
Staphylococcus pasteuri		13.96	0.19		86		
643613	**Good identification**						**643613**
Staphylococcus simulans		100.0	0.39	BGA 90			
643640	**Low discrimination**						**643640**
					YEP	PYR	
Staphylococcus piscifermentans		35.93	0.52	GLR 1	1	15	
Staphylococcus warneri		34.49	0.50		43	8	
Staphylococcus petrasii subsp. croceilyticus		16.57	0.18		99	90	
643642	**Low discrimination**						**643642**
					YEP		
Staphylococcus piscifermentans		75.13	0.55	GLR 1	1		
Staphylococcus warneri		16.03	0.40		43		
Staphylococcus pasteuri		7.41	0.19		86		
643650	**Low discrimination**						**643650**
					YEP		
Staphylococcus pasteuri		66.44	0.43	VPT 86	86		
Staphylococcus warneri		20.76	0.47	ESL 1	43		
Staphylococcus piscifermentans		12.80	0.44		1		
643652	**Low discrimination**						**643652**
					YEP		
Staphylococcus pasteuri		47.86	0.35	VPT 86	86		
Staphylococcus piscifermentans		33.87	0.48	ESL 1	1		
Staphylococcus warneri		12.21	0.37		43		
643653	**Good identification**						**643653**
Staphylococcus simulans		100.0	0.32	BGA 90			
643673	**Acceptable identification**						**643673**
Staphylococcus equorum		90.12	0.35	ARG 1			
Staphylococcus xylosus		9.88	0.14	GAL 91			
643700	**Very good identification**						**643700**
Staphylococcus piscifermentans		98.95	0.55	GLR 1			
Staphylococcus warneri		1.05	0.13				
643702	**Very good identification**						**643702**
Staphylococcus piscifermentans		99.34	0.59	GLR 1			
Staphylococcus hyicus		0.66	-.04				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
643703	**Good identification**					**643703**
Staphylococcus hyicus		97.29	0.37	ESL 1		
Staphylococcus piscifermentans		1.49	0.18	PHS 92		
Staphylococcus simulans		1.22	0.05			
643710	**Good identification**					**643710**
Staphylococcus piscifermentans		98.24	0.48	GLR 1		
Staphylococcus warneri		1.77	0.10			
643712	**Good identification**					**643712**
Staphylococcus piscifermentans		97.74	0.51	GLR 1		
Staphylococcus simulans		2.26	0.07			
643733	**Acceptable identification**					**643733**
Staphylococcus equorum		92.30	0.35	ARG 1		
Staphylococcus xylosus		7.70	0.11	MLT 91		
643740	**Low discrimination**					**643740**
Staphylococcus piscifermentans		83.19	0.55	GLR 1		
Staphylococcus warneri		16.81	0.40			
643742	**Very good identification**					**643742**
Staphylococcus piscifermentans		95.70	0.59	GLR 1		
Staphylococcus warneri		4.30	0.30			
643750	**Low discrimination**					**643750**
					YEP	
Staphylococcus piscifermentans		71.60	0.48	GLR 1	1	
Staphylococcus warneri		24.45	0.37		43	
Staphylococcus pasteuri		2.50	0.02		86	
643752	**Acceptable identification**					**643752**
Staphylococcus piscifermentans		92.06	0.51	GLR 1		
Staphylococcus warneri		6.99	0.27			
Staphylococcus simulans		0.96	-.01			
643771	**Low discrimination**					**643771**
Staphylococcus equorum		89.23	0.47	ARG 1		
Staphylococcus xylosus		8.83	0.25			
Staphylococcus gallinarum		1.94	0.10			
643773	**Low discrimination**					**643773**
Staphylococcus equorum		88.47	0.55	ARG 1		
Staphylococcus xylosus		9.70	0.34			
Staphylococcus gallinarum		1.83	0.18			
644250	**Good identification**					**644250**
Staphylococcus warneri		100.0	0.28	PHS 1		
644303	**Good identification**					**644303**
					YEP	
Staphylococcus hyicus		99.35	0.37	NIT 99	1	
Staphylococcus chromogenes		0.65	-.04	TRE 92	95	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
644413	**Acceptable identification**					**644413**
					NOV	
Staphylococcus cohnii subsp. urealyticum		90.84	0.40	ARG 1	99	
Staphylococcus simulans		9.16	0.14		1	
644440	**Good identification**					**644440**
Staphylococcus warneri		100.0	0.31	PHS 1		
644443	**Good identification**					**644443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ARG 1		
644450	**Low discrimination**					**644450**
					NOV	
Staphylococcus kloosii		76.84	0.47	ARG 1	99	
Staphylococcus warneri		23.16	0.28		1	
644451	**Good identification**					**644451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ARG 1		
644452	**Low discrimination**					**644452**
					NOV	
Staphylococcus kloosii		84.48	0.48	ARG 1	99	
Staphylococcus warneri		7.83	0.19		1	
Staphylococcus cohnii subsp. urealyticum		7.69	0.14		99	
644453	**Very good identification**					**644453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		99.21	0.55	ARG 1	99	
Staphylococcus simulans		0.79	0.07		1	
644470	**Good identification**					**644470**
Staphylococcus kloosii		100.0	0.34	ARG 1		
644472	**Good identification**					**644472**
Staphylococcus kloosii		100.0	0.35	ARG 1		
644512	**Good identification**					**644512**
Staphylococcus kloosii		100.0	0.27	ARG 1		
644513	**Acceptable identification**					**644513**
					NOV	
Staphylococcus cohnii subsp. urealyticum		91.94	0.32	ARG 1	99	
Staphylococcus felis		8.06	-.04		1	
644543	**Good identification**					**644543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1		
644550	**Good identification**					**644550**
					NOV	
Staphylococcus kloosii		95.33	0.53	ARG 1	99	
Staphylococcus warneri		4.67	0.18		1	
644551	**Good identification**					**644551**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1		
644552	**Very good identification**					**644552**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus kloosii		98.03	0.53	ARG 1		
Staphylococcus cohnii subsp. urealyticum		1.97	0.07			
644553	**Good identification**					**644553**
					NOV	
Staphylococcus cohnii subsp. urealyticum		97.63	0.48	ARG 1	99	
Staphylococcus caprae		2.37	-.02		1	
644570	**Good identification**					**644570**
Staphylococcus kloosii		100.0	0.40	ARG 1		
644572	**Good identification**					**644572**
Staphylococcus kloosii		100.0	0.41	ARG 1		
644600	**Low discrimination**					**644600**
Staphylococcus warneri		83.04	0.31	PHS 1		
Staphylococcus pasteuri		16.96	0.02	MLT 95		
644603	**Low discrimination**					**644603**
					YEP	
Staphylococcus hyicus		82.85	0.37	NIT 99	1	
Staphylococcus simulans		16.45	0.30	GAL 92	1	
Staphylococcus chromogenes		0.69	-.02		95	
644612	**Low discrimination**					**644612**
					YEP	
Staphylococcus simulans		79.87	0.31	BGA 90	1	
Staphylococcus pasteuri		13.68	0.11	NIT 96	86	
Staphylococcus warneri		6.44	0.19		43	
644613	**Good identification**					**644613**
					NOV	
Staphylococcus simulans		96.63	0.42	BGA 90	1	
Staphylococcus cohnii subsp. urealyticum		1.23	0.09	NIT 96	99	
Staphylococcus hyicus		1.23	-.04		1	
644640	**Low discrimination**					**644640**
					PYR	
Staphylococcus warneri		89.48	0.58	PHS 1	8	
Staphylococcus pasteuri		5.91	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		4.61	0.05		90	
644642	**Low discrimination**					**644642**
					YEP	
Staphylococcus warneri		88.19	0.48	PHS 1	43	
Staphylococcus pasteuri		7.14	0.11		86	
Staphylococcus simulans		4.68	0.12		1	
644650	**Low discrimination**					**644650**
					NOV	
Staphylococcus warneri		63.39	0.55	PHS 1	1	
Staphylococcus pasteuri		35.51	0.35		1	
Staphylococcus kloosii		1.10	0.26		99	
644652	**Low discrimination**					**644652**
					YEP	
Staphylococcus warneri		49.13	0.45	PHS 1	43	
Staphylococcus pasteuri		33.72	0.27		86	
Staphylococcus simulans		14.40	0.24		1	
644653	**Low discrimination**					**644653**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus simulans		84.61	0.35	BGA 90	1		
Staphylococcus cohnii subsp. urealyticum		13.63	0.24	NIT 96	99		
Staphylococcus warneri		1.76	0.10		1		
644701	**Good identification**						**644701**
					YEP		
Staphylococcus hyicus		99.17	0.37	NIT 99	1		
Staphylococcus chromogenes		0.83	-.02	LAC 92	95		
644703	**Very good identification**						**644703**
					YEP		
Staphylococcus hyicus		99.20	0.59	NIT 99	1		
Staphylococcus chromogenes		0.65	0.18		95		
Staphylococcus simulans		0.15	0.08		1		
644740	**Good identification**						**644740**
Staphylococcus warneri		100.0	0.47	PHS 1			
644742	**Good identification**						**644742**
Staphylococcus warneri		100.0	0.37	PHS 1			
644750	**Acceptable identification**						**644750**
					NOV		
Staphylococcus warneri		90.40	0.44	PHS 1	1		
Staphylococcus kloosii		9.60	0.32		99		
644752	**Low discrimination**						**644752**
					NOV	YEP	
Staphylococcus warneri		70.14	0.35	PHS 1	1	43	
Staphylococcus kloosii		24.20	0.33		99	99	
Staphylococcus simulans		5.66	0.02		1	1	
645413	**Good identification**						**645413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1			
645450	**Good identification**						**645450**
Staphylococcus kloosii		100.0	0.47	ARG 1			
645452	**Good identification**						**645452**
Staphylococcus kloosii		96.58	0.48	ARG 1			
Staphylococcus cohnii subsp. urealyticum		3.42	0.06				
645453	**Good identification**						**645453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	ARG 1			
645470	**Good identification**						**645470**
Staphylococcus kloosii		100.0	0.34	ARG 1			
645472	**Good identification**						**645472**
Staphylococcus kloosii		100.0	0.35	ARG 1			
645512	**Good identification**						**645512**
Staphylococcus kloosii		100.0	0.27	ARG 1			
645550	**Good identification**						**645550**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.53	ARG 1			
645552	**Very good identification**						**645552**
Staphylococcus kloosii		100.0	0.53	ARG 1			
645553	**Good identification**						**645553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ARG 1			
645570	**Good identification**						**645570**
Staphylococcus kloosii		100.0	0.40	ARG 1			
645572	**Good identification**						**645572**
Staphylococcus kloosii		100.0	0.41	ARG 1			
645652	**Good identification**						**645652**
Staphylococcus kloosii		100.0	0.27	ARG 1			
645750	**Good identification**						**645750**
Staphylococcus kloosii		100.0	0.32	ARG 1			
645752	**Good identification**						**645752**
Staphylococcus kloosii		100.0	0.33	ARG 1			
645771	**Low discrimination**						**645771**
Staphylococcus equorum		85.72	0.29	ARG 1			
Staphylococcus gallinarum		14.28	0.10	NIT 92			
645773	**Low discrimination**						**645773**
					YEP		
Staphylococcus equorum		80.96	0.37	ARG 1	1		
Staphylococcus gallinarum		12.84	0.18	NIT 92	78		
Staphylococcus arlettae		3.82	0.18		99		
646103	**Good identification**						**646103**
					YEP	COA	
Staphylococcus hyicus		96.99	0.37	SUC 99	1	24	
Staphylococcus schleiferi subsp. coagulans		2.38	0.12	TRE 92	1	99	
Staphylococcus chromogenes		0.64	-.04		95	1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
646201	**Good identification**						**646201**
					YEP		
Staphylococcus hyicus		98.94	0.35	GAL 92	1		
Staphylococcus chromogenes		1.06	-.02	TRE 92	95		
646203	**Very good identification**						**646203**
					YEP		
Staphylococcus hyicus		96.38	0.56	GAL 92	1		
Staphylococcus simulans		2.81	0.32	TRE 92	1		
Staphylococcus chromogenes		0.81	0.17		95		
646212	**Good identification**						**646212**
Staphylococcus simulans		100.0	0.33	BGA 90			
646213	**Low discrimination**						**646213**
					COA		
Staphylococcus simulans		76.47	0.44	BGA 90	1		
Staphylococcus delphini		14.78	0.18	TRE 95	99		
Staphylococcus hyicus		6.63	0.15		24		
646253	**Very good identification**						**646253**
					COA		
Staphylococcus delphini		97.39	0.59	GLR 1	99		
Staphylococcus simulans		2.29	0.37		1		
Staphylococcus capitis subsp. ureolyticus		0.22	0.03		1		
646301	**Very good identification**						**646301**
					YEP		
Staphylococcus hyicus		99.17	0.56	TRE 92	1		
Staphylococcus chromogenes		0.83	0.17	LAC 92	95		
646302	**Good identification**						**646302**
					YEP		
Staphylococcus hyicus		98.71	0.37	TRE 92	1		
Staphylococcus chromogenes		0.65	-.04	MNS 99	95		
Staphylococcus simulans		0.64	-.01		1		
646303	**Very good identification**						**646303**
					YEP		
Staphylococcus hyicus		99.32	0.78	TRE 92	1		
Staphylococcus chromogenes		0.65	0.37		95		
Staphylococcus simulans		0.02	0.10		1		
646313	**Low discrimination**						**646313**
					YEP	COA	
Staphylococcus hyicus		66.47	0.37	TRE 92	1	24	
Staphylococcus delphini		12.89	0.18	MAN 1	1	99	
Staphylococcus chromogenes		12.17	0.26		95	1	
646323	**Good identification**						**646323**
					YEP		
Staphylococcus hyicus		99.35	0.37	TRE 92	1		
Staphylococcus chromogenes		0.65	-.04	XYL 1	95		
646343	**Low discrimination**						**646343**
					YEP		
Staphylococcus hyicus		48.62	0.37	TRE 92	1		
Staphylococcus chromogenes		31.57	0.37	MLT 1	95		
Staphylococcus epidermidis		9.90	0.31		1		
646353	**Very good identification**						**646353**
					COA	YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus delphini		98.48	0.59	GLR 1	99	1	
Staphylococcus chromogenes		0.94	0.26		1	95	
Staphylococcus aureus subsp. aureus		0.23	0.01		99	84	
646403	**Low discrimination**						**646403**
Staphylococcus hyicus		79.28	0.37	GAL 92			
Staphylococcus simulans		15.74	0.30	SUC 99			
Staphylococcus felis		4.32	0.09				
646412	**Acceptable identification**						**646412**
Staphylococcus simulans		91.74	0.31	BGA 90			
Staphylococcus piscifermentans		8.26	0.21	SUC 96			
646413	**Low discrimination**						**646413**
					NOV		
Staphylococcus simulans		64.76	0.42	BGA 90	1		
Staphylococcus felis		29.70	0.26	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		4.72	0.24		99		
646442	**Low discrimination**						**646442**
Staphylococcus piscifermentans		56.75	0.28	GLR 1			
Staphylococcus simulans		30.35	0.12	ESL 90			
Staphylococcus warneri		12.90	0.14				
646453	**Low discrimination**						**646453**
					NOV		
Staphylococcus simulans		52.10	0.35	BGA 90	1		
Staphylococcus cohnii subsp. urealyticum		47.90	0.40	SUC 96	99		
646500	**Low discrimination**						**646500**
Staphylococcus piscifermentans		83.15	0.28	GLR 1			
Staphylococcus hyicus		16.85	-.04	ESL 90			
646501	**Good identification**						**646501**
					YEP		
Staphylococcus hyicus		99.17	0.37	SUC 99	1		
Staphylococcus chromogenes		0.83	-.02	LAC 92	95		
646503	**Very good identification**						**646503**
					YEP		
Staphylococcus hyicus		97.67	0.59	SUC 99	1		
Staphylococcus felis		1.55	0.20		1		
Staphylococcus chromogenes		0.64	0.18		95		
646513	**Low discrimination**						**646513**
Staphylococcus felis		75.94	0.37	GLR 1			
Staphylococcus condimenti		9.66	0.14				
Staphylococcus hyicus		7.23	0.18				
646540	**Low discrimination**						**646540**
Staphylococcus piscifermentans		82.28	0.28	GLR 1			
Staphylococcus warneri		17.72	0.13	ESL 90			
646542	**Good identification**						**646542**
Staphylococcus piscifermentans		100.0	0.32	GLR 1			
646601	**Very good identification**						**646601**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					YEP	
Staphylococcus hyicus		98.40	0.56	GAL 92	1	
Staphylococcus chromogenes		1.05	0.20	LAC 92	95	
Staphylococcus simulans		0.55	0.17		1	
646602	**Low discrimination**					**646602**
Staphylococcus simulans		56.17	0.47	BGA 90		
Staphylococcus hyicus		39.46	0.37			
Staphylococcus piscifermentans		2.95	0.32			
646603	**Good identification**					**646603**
					YEP	
Staphylococcus hyicus		94.55	0.78	GAL 92	1	
Staphylococcus simulans		4.55	0.58		1	
Staphylococcus chromogenes		0.79	0.39		95	
646610	**Low discrimination**					**646610**
					YEP	
Staphylococcus pasteuri		70.18	0.27	VPT 86	86	
Staphylococcus simulans		16.57	0.19	PHS 1	1	
Staphylococcus warneri		7.09	0.21	MLT 86	43	
646611	**Low discrimination**					**646611**
					YEP	
Staphylococcus simulans		61.52	0.30	BGA 90	1	
Staphylococcus hyicus		27.78	0.15	LAC 99	1	
Staphylococcus chromogenes		8.31	0.08		95	
646612	**Very good identification**					**646612**
					YEP	
Staphylococcus simulans		97.31	0.60	BGA 90	1	
Staphylococcus pasteuri		1.70	0.19		86	
Staphylococcus piscifermentans		0.55	0.25		1	
646613	**Good identification**					**646613**
Staphylococcus simulans		90.62	0.71	BGA 90		
Staphylococcus hyicus		4.75	0.37			
Staphylococcus felis		3.52	0.33			
646623	**Acceptable identification**					**646623**
					YEP	
Staphylococcus hyicus		94.65	0.37	GAL 92	1	
Staphylococcus simulans		4.56	0.17	XYL 1	1	
Staphylococcus chromogenes		0.79	-.02		95	
646633	**Acceptable identification**					**646633**
Staphylococcus simulans		91.64	0.30	BGA 90		
Staphylococcus hyicus		4.81	-.04	XYL 1		
Staphylococcus felis		3.56	-.09			
646640	**Low discrimination**					**646640**
					PYR	
Staphylococcus warneri		51.00	0.50	PHS 1	8	
Staphylococcus petrasii subsp. croceilyticus		24.51	0.18		90	
Staphylococcus pasteuri		19.24	0.27		1	
646642	**Low discrimination**					**646642**
					YEP	
Staphylococcus simulans		60.36	0.40	BGA 90	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		20.32	0.40		43			
Staphylococcus pasteuri		9.39	0.19		86			
646643	**Low discrimination**							**646643**
					YEP			
Staphylococcus simulans		53.92	0.51	BGA 90	1			
Staphylococcus hyicus		25.18	0.37		1			
Staphylococcus chromogenes		20.90	0.39		95			
646650	**Low discrimination**							**646650**
					NOV	PYR		
Staphylococcus pasteuri		72.46	0.43	VPT 86	1	1		
Staphylococcus warneri		22.64	0.47	PHS 1	1	8		
Staphylococcus saprophyticus subsp. bovis		2.62	0.18		99	99		
646652	**Low discrimination**							**646652**
					YEP			
Staphylococcus simulans		75.89	0.53	BGA 90	1			
Staphylococcus pasteuri		18.13	0.35		86			
Staphylococcus warneri		4.62	0.37		43			
646653	**Good identification**							**646653**
					COA	YEP		
Staphylococcus simulans		93.99	0.63	BGA 90	1	1		
Staphylococcus chromogenes		2.57	0.28		1	95		
Staphylococcus delphini		2.13	0.18		99	1		
646700	**Acceptable identification**							**646700**
Staphylococcus hyicus		91.42	0.37	MNS 99				
Staphylococcus piscifermentans		6.83	0.32	LAC 92				
Staphylococcus warneri		0.98	0.13					
646701	**Very good identification**							**646701**
					YEP	PYR	COA	
Staphylococcus hyicus		99.16	0.78	LAC 92	1	1	24	
Staphylococcus chromogenes		0.83	0.39		95	45	1	
Staphylococcus intermedius		0.01	-.03		1	90	99	
646702	**Very good identification**							**646702**
Staphylococcus hyicus		97.36	0.59	MNS 99				
Staphylococcus simulans		1.05	0.26					
Staphylococcus piscifermentans		0.95	0.36					
646703	**Excellent identification**							**646703**
					YEP			
Staphylococcus hyicus		99.28	1.00		1			
Staphylococcus chromogenes		0.65	0.59		95			
Staphylococcus simulans		0.04	0.36		1			
646711	**Low discrimination**							**646711**
					YEP			
Staphylococcus hyicus		77.33	0.37	MAN 1	1			
Staphylococcus chromogenes		18.10	0.28	LAC 92	95			
Staphylococcus felis		1.93	0.02		1			
646712	**Low discrimination**							**646712**
Staphylococcus simulans		70.17	0.38	BGA 90				
Staphylococcus hyicus		16.46	0.18	GAL 8				
Staphylococcus piscifermentans		6.81	0.28					

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
646713	**Low discrimination**							**646713**
					YEP			
Staphylococcus hyicus		63.45	0.59	MAN 1	1			
Staphylococcus felis		13.66	0.43		1			
Staphylococcus chromogenes		11.62	0.48		95			
646721	**Good identification**							**646721**
					YEP			
Staphylococcus hyicus		99.17	0.37	XYL 1	1			
Staphylococcus chromogenes		0.83	-.02	LAC 92	95			
646723	**Very good identification**							**646723**
					YEP			
Staphylococcus hyicus		99.35	0.59	XYL 1	1			
Staphylococcus chromogenes		0.65	0.18		95			
646740	**Low discrimination**							**646740**
Staphylococcus warneri		66.17	0.40	PHS 1				
Staphylococcus piscifermentans		24.25	0.32					
Staphylococcus hominis subsp. hominis		3.58	0.08					
646741	**Low discrimination**							**646741**
					YEP	PYR	COA	
Staphylococcus hyicus		54.37	0.37	MLT 1	1	1	24	
Staphylococcus chromogenes		45.11	0.39	LAC 92	95	45	1	
Staphylococcus intermedius		0.52	0.01		1	90	99	
646743	**Low discrimination**							**646743**
					YEP			
Staphylococcus hyicus		59.72	0.59	MLT 1	1			
Staphylococcus chromogenes		38.78	0.59		95			
Staphylococcus simulans		0.97	0.29		1			
646750	**Low discrimination**							**646750**
					NOV	PYR		
Staphylococcus saprophyticus subsp. bovis		52.58	0.31	ARG 1	99	99		
Staphylococcus warneri		35.95	0.37		1	8		
Staphylococcus piscifermentans		7.80	0.25		1	15		
646751	**Low discrimination**							**646751**
					COA	NOV		
Staphylococcus chromogenes		59.43	0.28	GLR 1	1	1		
Staphylococcus saprophyticus subsp. bovis		13.71	0.18	LAC 90	1	99		
Staphylococcus aureus subsp. aureus		11.52	0.01		99	4		
646752	**Low discrimination**							**646752**
					NOV	PYR		
Staphylococcus simulans		49.53	0.31	BGA 90	1	70		
Staphylococcus saprophyticus subsp. bovis		12.02	0.18	GAL 8	99	99		
Staphylococcus warneri		10.96	0.27		1	8		
646753	**Low discrimination**							**646753**
					COA	YEP		
Staphylococcus aureus subsp. aureus		56.41	0.42	VPT 87	99	84		
Staphylococcus chromogenes		26.45	0.48	GLR 1	1	95		
Staphylococcus simulans		9.35	0.42		1	1		
646773	**Low discrimination**							**646773**
Staphylococcus xylosus		70.16	0.35	ARG 1				
Staphylococcus gallinarum		11.74	0.18					

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus equorum		8.60	0.18				
647303	**Good identification**						**647303**
					YEP	COA	
Staphylococcus hyicus		98.62	0.37	ESL 1	1	24	
Staphylococcus schleiferi subsp. coagulans		0.73	0.02	TRE 92	1	99	
Staphylococcus chromogenes		0.65	-.04		95	1	
647400	**Good identification**						**647400**
Staphylococcus piscifermentans		100.0	0.44	GLR 1			
647402	**Good identification**						**647402**
Staphylococcus piscifermentans		100.0	0.48	GLR 1			
647410	**Good identification**						**647410**
Staphylococcus piscifermentans		100.0	0.37	GLR 1			
647412	**Good identification**						**647412**
Staphylococcus piscifermentans		100.0	0.41	GLR 1			
647440	**Good identification**						**647440**
Staphylococcus piscifermentans		100.0	0.44	GLR 1			
647442	**Good identification**						**647442**
Staphylococcus piscifermentans		100.0	0.48	GLR 1			
647450	**Good identification**						**647450**
Staphylococcus piscifermentans		100.0	0.37	GLR 1			
647452	**Good identification**						**647452**
Staphylococcus piscifermentans		100.0	0.41	GLR 1			
647453	**Good identification**						**647453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1			
647500	**Good identification**						**647500**
Staphylococcus piscifermentans		100.0	0.48	GLR 1			
647502	**Good identification**						**647502**
Staphylococcus piscifermentans		100.0	0.52	GLR 1			
647510	**Good identification**						**647510**
Staphylococcus piscifermentans		100.0	0.41	GLR 1			
647512	**Good identification**						**647512**
Staphylococcus piscifermentans		100.0	0.44	GLR 1			
647540	**Good identification**						**647540**
Staphylococcus piscifermentans		100.0	0.48	GLR 1			
647542	**Good identification**						**647542**
Staphylococcus piscifermentans		100.0	0.52	GLR 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
647550	**Good identification**					**647550**
Staphylococcus piscifermentans		100.0	0.41	GLR 1		
647552	**Good identification**					**647552**
Staphylococcus piscifermentans		100.0	0.44	GLR 1		
647600	**Good identification**					**647600**
Staphylococcus piscifermentans		100.0	0.48	GLR 1		
647602	**Good identification**					**647602**
Staphylococcus piscifermentans		96.49	0.52	GLR 1		
Staphylococcus simulans		2.06	0.06			
Staphylococcus hyicus		1.45	-.04			
647603	**Acceptable identification**					**647603**
					YEP	
Staphylococcus hyicus		94.05	0.37	ESL 1	1	
Staphylococcus simulans		4.53	0.17	GAL 92	1	
Staphylococcus chromogenes		0.79	-.02		95	
647610	**Good identification**					**647610**
Staphylococcus piscifermentans		100.0	0.41	GLR 1		
647612	**Low discrimination**					**647612**
Staphylococcus piscifermentans		83.37	0.44	GLR 1		
Staphylococcus simulans		16.63	0.19			
647613	**Acceptable identification**					**647613**
Staphylococcus simulans		91.64	0.30	BGA 90		
Staphylococcus hyicus		4.81	-.04	ESL 1		
Staphylococcus felis		3.56	-.09			
647640	**Good identification**					**647640**
Staphylococcus piscifermentans		98.57	0.48	GLR 1		
Staphylococcus warneri		1.43	0.09			
647642	**Good identification**					**647642**
Staphylococcus piscifermentans		99.05	0.52	GLR 1		
Staphylococcus simulans		0.95	-.01			
647650	**Acceptable identification**					**647650**
					YEP	
Staphylococcus piscifermentans		92.71	0.41	GLR 1	1	
Staphylococcus pasteuri		7.29	0.02		86	
647652	**Acceptable identification**					**647652**
Staphylococcus piscifermentans		91.78	0.44	GLR 1		
Staphylococcus simulans		8.23	0.12			
647671	**Low discrimination**					**647671**
Staphylococcus equorum		87.22	0.30	ARG 1		
Staphylococcus gallinarum		12.78	0.10	GAL 91		
647673	**Low discrimination**					**647673**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus equorum		86.20	0.38	ARG 1		
Staphylococcus gallinarum		12.02	0.18	GAL 91		
Staphylococcus xylosus		1.78	0.02			
647700	**Good identification**					**647700**
Staphylococcus piscifermentans		98.52	0.52	GLR 1		
Staphylococcus hyicus		1.48	-.04			
647701	**Good identification**					**647701**
					YEP	
Staphylococcus hyicus		98.51	0.37	ESL 1	1	
Staphylococcus chromogenes		0.83	-.02	LAC 92	95	
Staphylococcus piscifermentans		0.67	0.11		1	
647702	**Low discrimination**					**647702**
Staphylococcus piscifermentans		89.68	0.55	GLR 1		
Staphylococcus hyicus		10.32	0.18			
647703	**Very good identification**					**647703**
					YEP	
Staphylococcus hyicus		99.26	0.59	ESL 1	1	
Staphylococcus chromogenes		0.65	0.18		95	
Staphylococcus piscifermentans		0.09	0.14		1	
647710	**Good identification**					**647710**
Staphylococcus piscifermentans		100.0	0.44	GLR 1		
647712	**Good identification**					**647712**
Staphylococcus piscifermentans		100.0	0.48	GLR 1		
647731	**Low discrimination**					**647731**
Staphylococcus equorum		87.22	0.30	ARG 1		
Staphylococcus gallinarum		12.78	0.10	MLT 91		
647733	**Low discrimination**					**647733**
Staphylococcus equorum		86.57	0.38	ARG 1		
Staphylococcus gallinarum		12.07	0.18	MLT 91		
Staphylococcus xylosus		1.36	0.00			
647740	**Good identification**					**647740**
Staphylococcus piscifermentans		100.0	0.52	GLR 1		
647742	**Very good identification**					**647742**
Staphylococcus piscifermentans		100.0	0.55	GLR 1		
647750	**Good identification**					**647750**
Staphylococcus piscifermentans		100.0	0.44	GLR 1		
647752	**Good identification**					**647752**
Staphylococcus piscifermentans		100.0	0.48	GLR 1		
647761	**Acceptable identification**					**647761**
Staphylococcus gallinarum		93.35	0.30	ARG 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus equorum		6.65	0.09	MAN 91		
647763	**Acceptable identification**					**647763**
Staphylococcus gallinarum		92.21	0.38	ARG 1		
Staphylococcus equorum		6.90	0.18	MAN 91		
Staphylococcus xylosus		0.90	-.02			
647771	**Low discrimination**					**647771**
Staphylococcus gallinarum		58.47	0.51	ARG 1		
Staphylococcus equorum		40.77	0.51			
Staphylococcus xylosus		0.76	0.14			
647773	**Low discrimination**					**647773**
Staphylococcus gallinarum		57.23	0.59	ARG 1		
Staphylococcus equorum		41.91	0.59			
Staphylococcus xylosus		0.86	0.23			
650240	**Good identification**					**650240**
Staphylococcus warneri		100.0	0.31	ORN 1		
650250	**Good identification**					**650250**
Staphylococcus warneri		100.0	0.28	ORN 1		
650440	**Good identification**					**650440**
Staphylococcus warneri		100.0	0.31	ORN 1		
650450	**Good identification**					**650450**
Staphylococcus warneri		100.0	0.28	ORN 1		
650600	**Low discrimination**					**650600**
Staphylococcus warneri		83.04	0.31	ORN 1		
Staphylococcus pasteuri		16.96	0.02	MLT 95		
650640	**Low discrimination**					**650640**
					PYR	
Staphylococcus warneri		89.48	0.58	ORN 1	8	
Staphylococcus pasteuri		5.91	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		4.61	0.05		90	
650642	**Acceptable identification**					**650642**
Staphylococcus warneri		92.51	0.48	ORN 1		
Staphylococcus pasteuri		7.49	0.11			
650650	**Low discrimination**					**650650**
Staphylococcus warneri		64.10	0.55	ORN 1		
Staphylococcus pasteuri		35.90	0.35			
650652	**Low discrimination**					**650652**
Staphylococcus warneri		59.30	0.45	ORN 1		
Staphylococcus pasteuri		40.70	0.27			
650740	**Good identification**					**650740**
Staphylococcus warneri		100.0	0.47	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
650742	**Good identification**					**650742**
	Staphylococcus warneri	100.0	0.37	ORN 1		
650750	**Good identification**					**650750**
	Staphylococcus warneri	100.0	0.44	ORN 1		
650752	**Good identification**					**650752**
	Staphylococcus warneri	100.0	0.35	ORN 1		
652603	**Low discrimination**					**652603**
	Staphylococcus simulans	62.42	0.27	ORN 1		
	Staphylococcus hyicus	37.58	0.15	BGA 90		
652610	**Acceptable identification**					**652610**
	Staphylococcus pasteuri	90.82	0.27	VPT 86		
	Staphylococcus warneri	9.18	0.21	ORN 1		
652612	**Low discrimination**					**652612**
					YEP	
	Staphylococcus simulans	61.57	0.29	ORN 1	1	
	Staphylococcus pasteuri	35.50	0.19	BGA 90	86	
	Staphylococcus warneri	2.93	0.11		43	
652613	**Good identification**					**652613**
	Staphylococcus simulans	100.0	0.39	ORN 1		
652640	**Low discrimination**					**652640**
					PYR	
	Staphylococcus warneri	53.83	0.50	ORN 1	8	
	Staphylococcus petrasii subsp. croceilyticus	25.87	0.18		90	
	Staphylococcus pasteuri	20.31	0.27		1	
652642	**Low discrimination**					**652642**
					YEP	
	Staphylococcus warneri	64.42	0.40	ORN 1	43	
	Staphylococcus pasteuri	29.78	0.19		86	
	Staphylococcus simulans	5.80	0.09		1	
652650	**Low discrimination**					**652650**
	Staphylococcus pasteuri	76.19	0.43	VPT 86		
	Staphylococcus warneri	23.81	0.47	ORN 1		
652652	**Low discrimination**					**652652**
					YEP	
	Staphylococcus pasteuri	72.37	0.35	VPT 86	86	
	Staphylococcus warneri	18.46	0.37	ORN 1	43	
	Staphylococcus simulans	9.18	0.21		1	
652653	**Good identification**					**652653**
	Staphylococcus simulans	100.0	0.32	ORN 1		
652703	**Good identification**					**652703**
	Staphylococcus hyicus	98.76	0.37	ORN 1		
	Staphylococcus simulans	1.24	0.05	PHS 92		
652740	**Good identification**					**652740**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		100.0	0.40	ORN 1			
652742	**Good identification**						**652742**
Staphylococcus warneri		100.0	0.30	ORN 1			
652750	**Low discrimination**						**652750**
					NOV	PYR	
Staphylococcus warneri		86.10	0.37	ORN 1	1	8	
Staphylococcus pasteuri		8.81	0.02		1	1	
Staphylococcus saprophyticus subsp. bovis		5.09	0.02		99	99	
652752	**Low discrimination**						**652752**
Staphylococcus warneri		87.95	0.27	ORN 1			
Staphylococcus simulans		12.05	-.01				
656303	**Good identification**						**656303**
					YEP		
Staphylococcus hyicus		99.35	0.37	ORN 1	1		
Staphylococcus chromogenes		0.65	-.04	TRE 92	95		
656603	**Acceptable identification**						**656603**
					YEP		
Staphylococcus hyicus		94.65	0.37	ORN 1	1		
Staphylococcus simulans		4.56	0.17	GAL 92	1		
Staphylococcus chromogenes		0.79	-.02		95		
656613	**Acceptable identification**						**656613**
Staphylococcus simulans		91.64	0.30	ORN 1			
Staphylococcus hyicus		4.81	-.04	BGA 90			
Staphylococcus felis		3.56	-.09				
656701	**Good identification**						**656701**
					YEP		
Staphylococcus hyicus		99.17	0.37	ORN 1	1		
Staphylococcus chromogenes		0.83	-.02	LAC 92	95		
656703	**Very good identification**						**656703**
					YEP		
Staphylococcus hyicus		99.35	0.59	ORN 1	1		
Staphylococcus chromogenes		0.65	0.18		95		
660040	**Good identification**						**660040**
Staphylococcus warneri		100.0	0.34	SUC 95			
660050	**Good identification**						**660050**
Staphylococcus warneri		100.0	0.31	SUC 95			
660200	**Good identification**						**660200**
Staphylococcus warneri		100.0	0.34	TRE 95			
660203	**Good identification**						**660203**
Staphylococcus simulans		100.0	0.33	NIT 96			
660210	**Good identification**						**660210**
Staphylococcus warneri		100.0	0.31	TRE 95			
660212	**Acceptable identification**						**660212**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		93.11	0.35	NIT 96		
Staphylococcus warneri		6.89	0.21	TRE 95		
660213	**Good identification**					**660213**
Staphylococcus simulans		100.0	0.45	NIT 96		
660240	**Very good identification**					**660240**
Staphylococcus warneri		100.0	0.60	TRE 95		
660242	**Acceptable identification**					**660242**
Staphylococcus warneri		94.53	0.50	TRE 95		
Staphylococcus simulans		5.47	0.15			
660250	**Very good identification**					**660250**
Staphylococcus warneri		100.0	0.57	TRE 95		
660252	**Low discrimination**					**660252**
Staphylococcus warneri		75.77	0.47	TRE 95		
Staphylococcus simulans		24.23	0.27			
660253	**Acceptable identification**					**660253**
Staphylococcus simulans		92.66	0.38	NIT 96		
Staphylococcus capitis subsp. ureolyticus		5.57	0.00	TRE 95		
Staphylococcus warneri		1.77	0.12			
660340	**Good identification**					**660340**
Staphylococcus warneri		100.0	0.50	TRE 95		
660342	**Good identification**					**660342**
Staphylococcus warneri		100.0	0.40	TRE 95		
660350	**Good identification**					**660350**
Staphylococcus warneri		100.0	0.47	TRE 95		
660352	**Acceptable identification**					**660352**
Staphylococcus warneri		91.91	0.37	TRE 95		
Staphylococcus simulans		8.09	0.06			
660400	**Good identification**					**660400**
Staphylococcus warneri		100.0	0.34	SUC 95		
660403	**Low discrimination**					**660403**
					NOV	
Staphylococcus simulans		87.32	0.31	NIT 96	1	
Staphylococcus cohnii subsp. urealyticum		12.68	0.19	SUC 96	99	
660410	**Good identification**					**660410**
Staphylococcus warneri		100.0	0.31	SUC 95		
660412	**Acceptable identification**					**660412**
Staphylococcus simulans		91.46	0.32	NIT 96		
Staphylococcus warneri		8.54	0.21	SUC 96		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
660413	**Low discrimination**					**660413**
					NOV	
Staphylococcus simulans		73.15	0.43	NIT 96	1	
Staphylococcus cohnii subsp. urealyticum		26.85	0.40	SUC 96	99	
660440	**Very good identification**					**660440**
Staphylococcus warneri		100.0	0.60	SUC 95		
660442	**Good identification**					**660442**
Staphylococcus warneri		95.62	0.50	SUC 95		
Staphylococcus simulans		4.38	0.13			
660443	**Low discrimination**					**660443**
					NOV	
Staphylococcus cohnii subsp. urealyticum		61.79	0.35	ARG 1	99	
Staphylococcus simulans		33.73	0.24	MAN 91	1	
Staphylococcus warneri		4.49	0.15		1	
660450	**Very good identification**					**660450**
					NOV	
Staphylococcus warneri		99.26	0.57	SUC 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		0.74	-.08		99	
660451	**Low discrimination**					**660451**
					NOV	
Staphylococcus cohnii subsp. urealyticum		86.37	0.35	ARG 1	99	
Staphylococcus warneri		13.63	0.22	LAC 91	1	
660452	**Low discrimination**					**660452**
					NOV	
Staphylococcus warneri		70.46	0.47	SUC 95	1	
Staphylococcus simulans		17.83	0.25		1	
Staphylococcus saprophyticus subsp. saprophyticus		8.92	0.08		99	
660453	**Low discrimination**					**660453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		81.89	0.55	ARG 1	99	
Staphylococcus simulans		17.69	0.36		1	
Staphylococcus warneri		0.43	0.12		1	
660513	**Low discrimination**					**660513**
					NOV	
Staphylococcus cohnii subsp. urealyticum		64.41	0.32	ARG 1	99	
Staphylococcus simulans		35.60	0.21		1	
660540	**Good identification**					**660540**
Staphylococcus warneri		100.0	0.50	SUC 95		
660542	**Good identification**					**660542**
Staphylococcus warneri		100.0	0.40	SUC 95		
660543	**Acceptable identification**					**660543**
					NOV	
Staphylococcus cohnii subsp. urealyticum		90.03	0.27	ARG 1	99	
Staphylococcus simulans		9.97	0.02	MAN 91	1	
660550	**Good identification**					**660550**
Staphylococcus warneri		100.0	0.47	SUC 95		
660551	**Low discrimination**					**660551**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					NOV	
Staphylococcus cohnii subsp. urealyticum		89.59	0.27	ARG 1	99	
Staphylococcus warneri		10.42	0.12	LAC 91	1	
660552	**Low discrimination**					**660552**
					NOV	
Staphylococcus warneri		89.01	0.37	SUC 95	1	
Staphylococcus simulans		6.20	0.03		1	
Staphylococcus cohnii subsp. urealyticum		4.79	0.07		99	
660553	**Good identification**					**660553**
					NOV	
Staphylococcus cohnii subsp. urealyticum		95.80	0.48	ARG 1	99	
Staphylococcus simulans		4.20	0.14		1	
660600	**Very good identification**					**660600**
					YEP	
Staphylococcus warneri		98.38	0.60	MLT 95	43	
Staphylococcus pasteuri		0.81	0.02		86	
Staphylococcus simulans		0.81	0.07		1	
660602	**Low discrimination**					**660602**
					YEP	
Staphylococcus simulans		70.79	0.48	NIT 96	1	
Staphylococcus warneri		28.92	0.50		43	
Staphylococcus pasteuri		0.29	-.06		86	
660603	**Very good identification**					**660603**
Staphylococcus simulans		99.57	0.59	NIT 96		
Staphylococcus warneri		0.25	0.15			
Staphylococcus hyicus		0.18	-.04			
660610	**Low discrimination**					**660610**
					YEP	
Staphylococcus warneri		89.64	0.57	MLT 95	43	
Staphylococcus pasteuri		6.28	0.19		86	
Staphylococcus simulans		4.08	0.20		1	
660611	**Low discrimination**					**660611**
Staphylococcus simulans		88.19	0.31	NIT 96		
Staphylococcus warneri		11.81	0.22	LAC 99		
660612	**Good identification**					**660612**
					YEP	
Staphylococcus simulans		92.57	0.61	NIT 96	1	
Staphylococcus warneri		6.85	0.47		43	
Staphylococcus pasteuri		0.59	0.11		86	
660613	**Very good identification**					**660613**
					NOV	
Staphylococcus simulans		99.91	0.72	NIT 96	1	
Staphylococcus cohnii subsp. urealyticum		0.05	0.09		99	
Staphylococcus warneri		0.05	0.12		1	
660633	**Good identification**					**660633**
					NOV	
Staphylococcus simulans		96.81	0.31	NIT 96	1	
Staphylococcus xylosus		3.19	-.04	XYL 1	99	
660640	**Very good identification**					**660640**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		99.51	0.86		8	
Staphylococcus pasteuri		0.27	0.19		1	
Staphylococcus petrasii subsp. croceilyticus		0.21	0.05		90	
660641	**Good identification**					**660641**
Staphylococcus warneri		96.90	0.51	MNS 2		
Staphylococcus simulans		3.10	0.11			
660642	**Good identification**					**660642**
					YEP	
Staphylococcus warneri		94.06	0.76		43	
Staphylococcus simulans		5.44	0.41		1	
Staphylococcus pasteuri		0.31	0.11		86	
660643	**Acceptable identification**					**660643**
Staphylococcus simulans		90.47	0.52	NIT 96		
Staphylococcus warneri		9.53	0.42			
660650	**Very good identification**					**660650**
					NOV	
Staphylococcus warneri		97.31	0.83		1	
Staphylococcus pasteuri		2.20	0.35		1	
Staphylococcus saprophyticus subsp. saprophyticus		0.39	0.13		99	
660651	**Low discrimination**					**660651**
Staphylococcus warneri		84.99	0.49	MNS 2		
Staphylococcus simulans		15.01	0.23			
660652	**Low discrimination**					**660652**
					NOV	
Staphylococcus warneri		70.68	0.74		1	
Staphylococcus simulans		22.60	0.54		1	
Staphylococcus saprophyticus subsp. saprophyticus		4.76	0.29		99	
660653	**Very good identification**					**660653**
					NOV	
Staphylococcus simulans		97.48	0.64	NIT 96	1	
Staphylococcus warneri		1.86	0.39		1	
Staphylococcus cohnii subsp. urealyticum		0.58	0.24		99	
660660	**Good identification**					**660660**
Staphylococcus warneri		100.0	0.45	XYL 1		
660662	**Acceptable identification**					**660662**
Staphylococcus warneri		94.53	0.35	XYL 1		
Staphylococcus simulans		5.47	0.00			
660670	**Good identification**					**660670**
Staphylococcus warneri		100.0	0.42	XYL 1		
660672	**Low discrimination**					**660672**
Staphylococcus warneri		75.77	0.32	XYL 1		
Staphylococcus simulans		24.23	0.13			
660700	**Good identification**					**660700**
Staphylococcus warneri		100.0	0.50	MLT 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
660702	**Low discrimination**						**660702**
Staphylococcus warneri		59.74	0.40	MLT 95			
Staphylococcus simulans		40.26	0.27				
660703	**Low discrimination**						**660703**
					COA	PYR	
Staphylococcus simulans		79.50	0.37	NIT 96	1	70	
Staphylococcus hyicus		18.95	0.17	GAL 8	24	1	
Staphylococcus intermedius		1.55	0.00		99	90	
660710	**Good identification**						**660710**
Staphylococcus warneri		100.0	0.47	MLT 95			
660712	**Low discrimination**						**660712**
Staphylococcus simulans		78.83	0.39	NIT 96			
Staphylococcus warneri		21.17	0.37	GAL 8			
660713	**Good identification**						**660713**
					COA	NOV	
Staphylococcus simulans		98.37	0.50	NIT 96	1	1	
Staphylococcus intermedius		1.30	0.09	GAL 8	99	1	
Staphylococcus xylosus		0.33	-.05		1	99	
660740	**Very good identification**						**660740**
Staphylococcus warneri		100.0	0.76				
660741	**Good identification**						**660741**
Staphylococcus warneri		100.0	0.41	MNS 2			
660742	**Very good identification**						**660742**
Staphylococcus warneri		98.43	0.66				
Staphylococcus simulans		1.57	0.19				
660743	**Low discrimination**						**660743**
					COA	PYR	
Staphylococcus simulans		69.09	0.30	NIT 96	1	70	
Staphylococcus warneri		26.42	0.31	GAL 8	1	8	
Staphylococcus intermedius		4.49	0.04		99	90	
660750	**Very good identification**						**660750**
					NOV	PYR	
Staphylococcus warneri		99.74	0.73		1	8	
Staphylococcus saprophyticus subsp. bovis		0.15	0.05		99	99	
Staphylococcus saprophyticus subsp. saprophyticus		0.11	-.09		99	7	
660751	**Low discrimination**						**660751**
					NOV		
Staphylococcus warneri		82.16	0.38	MNS 2	1		
Staphylococcus xylosus		13.85	0.09		99		
Staphylococcus simulans		4.00	0.02		1		
660752	**Good identification**						**660752**
					NOV		
Staphylococcus warneri		90.37	0.63		1		
Staphylococcus simulans		7.96	0.32		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.68	0.07		99		
660753	**Low discrimination**						**660753**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus simulans		80.50	0.43	NIT 96	1		
Staphylococcus xylosus		8.02	0.18	GAL 8	99		
Staphylococcus warneri		5.57	0.28		1		
660760	**Good identification**						**660760**
Staphylococcus warneri		100.0	0.35	XYL 1			
660770	**Good identification**						**660770**
Staphylococcus warneri		100.0	0.32	XYL 1			
660771	**Good identification**						**660771**
Staphylococcus xylosus		100.0	0.31	ARG 1			
660773	**Good identification**						**660773**
					NOV	YEP	
Staphylococcus xylosus		97.30	0.40	ARG 1	99	40	
Staphylococcus arlettae		1.84	0.14	NIT 89	99	99	
Staphylococcus simulans		0.86	0.02		1	1	
661413	**Acceptable identification**						**661413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		93.39	0.31	ARG 1	99		
Staphylococcus simulans		6.61	0.02		1		
661453	**Good identification**						**661453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.47	ARG 1			
661553	**Good identification**						**661553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ARG 1			
661613	**Good identification**						**661613**
Staphylococcus simulans		100.0	0.31	ESL 1			
661640	**Good identification**						**661640**
Staphylococcus warneri		100.0	0.45	ESL 1			
661642	**Acceptable identification**						**661642**
Staphylococcus warneri		94.53	0.35	ESL 1			
Staphylococcus simulans		5.47	0.00				
661650	**Good identification**						**661650**
Staphylococcus warneri		100.0	0.42	ESL 1			
661652	**Low discrimination**						**661652**
Staphylococcus warneri		75.77	0.32	ESL 1			
Staphylococcus simulans		24.23	0.13				
661740	**Good identification**						**661740**
Staphylococcus warneri		100.0	0.35	ESL 1			
661750	**Good identification**						**661750**
Staphylococcus warneri		100.0	0.32	ESL 1			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
661773	**Low discrimination**					**661773**
					YEP	
Staphylococcus equorum		54.97	0.32	ARG 1	1	
Staphylococcus xylosus		41.86	0.28	NIT 92	40	
Staphylococcus arlettae		3.17	0.14		99	
662003	**Good identification**					**662003**
Staphylococcus simulans		100.0	0.33	SUC 96		
662012	**Good identification**					**662012**
Staphylococcus simulans		100.0	0.35	SUC 96		
662013	**Good identification**					**662013**
Staphylococcus simulans		100.0	0.45	SUC 96		
662052	**Acceptable identification**					**662052**
Staphylococcus simulans		93.41	0.27	SUC 96		
Staphylococcus warneri		6.59	0.13	TRE 95		
662053	**Good identification**					**662053**
Staphylococcus simulans		95.46	0.38	SUC 96		
Staphylococcus capitis subsp. ureolyticus		4.54	-.02	TRE 95		
662202	**Good identification**					**662202**
Staphylococcus simulans		99.28	0.51	TRE 95		
Staphylococcus warneri		0.72	0.16			
662203	**Very good identification**					**662203**
Staphylococcus simulans		98.79	0.61	TRE 95		
Staphylococcus hyicus		1.21	0.15			
662211	**Good identification**					**662211**
Staphylococcus simulans		100.0	0.33	TRE 95		
662212	**Very good identification**					**662212**
Staphylococcus simulans		99.87	0.63	TRE 95		
Staphylococcus warneri		0.13	0.13			
662213	**Very good identification**					**662213**
Staphylococcus simulans		99.91	0.74	TRE 95		
Staphylococcus capitis subsp. ureolyticus		0.09	-.02			
662233	**Good identification**					**662233**
Staphylococcus simulans		100.0	0.33	TRE 95		
662240	**Good identification**					**662240**
Staphylococcus warneri		98.92	0.52	TRE 95		
Staphylococcus simulans		1.08	0.02			
662242	**Low discrimination**					**662242**
Staphylococcus simulans		76.42	0.43	TRE 95		
Staphylococcus warneri		23.58	0.43			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
662243	**Very good identification**						**662243**
Staphylococcus simulans		99.21	0.54	TRE 95			
Staphylococcus capitis subsp. ureolyticus		0.79	-.02				
662250	**Acceptable identification**						**662250**
					YEP		
Staphylococcus warneri		92.16	0.49	TRE 95	43		
Staphylococcus simulans		5.56	0.15		1		
Staphylococcus pasteuri		2.29	0.02		86		
662252	**Good identification**						**662252**
Staphylococcus simulans		94.71	0.56	TRE 95			
Staphylococcus warneri		5.29	0.40				
662253	**Very good identification**						**662253**
Staphylococcus simulans		95.46	0.67	TRE 95			
Staphylococcus capitis subsp. ureolyticus		4.54	0.26				
662302	**Acceptable identification**						**662302**
Staphylococcus simulans		94.79	0.29	GAL 8			
Staphylococcus hyicus		5.21	-.04	TRE 95			
662303	**Low discrimination**						**662303**
					COA	PYR	
Staphylococcus hyicus		61.53	0.37	BGA 10	24	1	
Staphylococcus simulans		37.88	0.40	PHS 92	1	70	
Staphylococcus intermedius		0.58	0.00	TRE 92	99	90	
662312	**Good identification**						**662312**
Staphylococcus simulans		100.0	0.41	GAL 8			
662313	**Good identification**						**662313**
					COA	PYR	
Staphylococcus simulans		98.57	0.52	GAL 8	1	70	
Staphylococcus intermedius		1.03	0.09	TRE 95	99	90	
Staphylococcus hyicus		0.40	-.04		24	1	
662340	**Good identification**						**662340**
Staphylococcus warneri		100.0	0.42	TRE 95			
662342	**Low discrimination**						**662342**
Staphylococcus warneri		52.84	0.32	TRE 95			
Staphylococcus simulans		47.16	0.22				
662343	**Low discrimination**						**662343**
					COA	PYR	
Staphylococcus simulans		89.13	0.32	GAL 8	1	70	
Staphylococcus intermedius		4.59	0.04	TRE 95	99	90	
Staphylococcus hyicus		3.26	-.04		24	1	
662350	**Acceptable identification**						**662350**
					NOV	PYR	
Staphylococcus warneri		93.63	0.39	TRE 95	1	8	
Staphylococcus saprophyticus subsp. bovis		6.37	0.05		99	99	
662352	**Low discrimination**						**662352**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		83.14	0.34	GAL 8			
Staphylococcus warneri		16.86	0.29	TRE 95			
662353	**Acceptable identification**						**662353**
					COA	NOV	
Staphylococcus simulans		94.22	0.45	GAL 8	1	1	
Staphylococcus intermedius		3.28	0.13	TRE 95	99	1	
Staphylococcus xylosus		2.51	0.09		1	99	
662373	**Good identification**						**662373**
					NOV		
Staphylococcus xylosus		96.80	0.31	ARG 1	99		
Staphylococcus simulans		3.20	0.04	TRE 96	1		
662402	**Good identification**						**662402**
Staphylococcus simulans		99.09	0.48	SUC 96			
Staphylococcus warneri		0.91	0.16				
662403	**Very good identification**						**662403**
Staphylococcus simulans		99.82	0.59	SUC 96			
Staphylococcus hyicus		0.18	-.04				
662411	**Good identification**						**662411**
Staphylococcus simulans		100.0	0.31	SUC 96			
662412	**Very good identification**						**662412**
Staphylococcus simulans		99.83	0.61	SUC 96			
Staphylococcus warneri		0.17	0.13				
662413	**Very good identification**						**662413**
					NOV		
Staphylococcus simulans		99.63	0.72	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		0.27	0.24		99		
Staphylococcus felis		0.10	0.01		1		
662433	**Good identification**						**662433**
Staphylococcus simulans		100.0	0.31	SUC 96			
662440	**Good identification**						**662440**
Staphylococcus warneri		96.59	0.52	SUC 95			
Staphylococcus auricularis		2.57	0.12				
Staphylococcus simulans		0.84	0.00				
662442	**Low discrimination**						**662442**
Staphylococcus simulans		71.96	0.41	SUC 96			
Staphylococcus warneri		28.04	0.43				
662443	**Good identification**						**662443**
					NOV		
Staphylococcus simulans		98.67	0.52	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		1.33	0.19		99		
662450	**Low discrimination**						**662450**
					NOV	PYR	
Staphylococcus warneri		82.95	0.49	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		11.03	0.22		99	99	
Staphylococcus simulans		3.96	0.13		1	70	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
662452	**Good identification**						**662452**
					NOV	PYR	
Staphylococcus simulans		92.80	0.54	SUC 96	1	70	
Staphylococcus warneri		6.55	0.40		1	8	
Staphylococcus saprophyticus subsp. bovis		0.65	0.09		99	99	
662453	**Very good identification**						**662453**
					NOV		
Staphylococcus simulans		96.71	0.64	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		3.29	0.40		99		
662502	**Good identification**						**662502**
Staphylococcus simulans		100.0	0.27	GAL 8			
662503	**Low discrimination**						**662503**
Staphylococcus simulans		79.32	0.37	GAL 8			
Staphylococcus hyicus		18.91	0.17	SUC 96			
Staphylococcus felis		1.78	-.06				
662512	**Good identification**						**662512**
Staphylococcus simulans		100.0	0.39	GAL 8			
662513	**Low discrimination**						**662513**
Staphylococcus simulans		88.19	0.50	GAL 8			
Staphylococcus condimenti		7.34	0.14	SUC 96			
Staphylococcus felis		3.30	0.12				
662540	**Good identification**						**662540**
Staphylococcus warneri		100.0	0.42	SUC 95			
662542	**Low discrimination**						**662542**
Staphylococcus warneri		58.60	0.32	SUC 95			
Staphylococcus simulans		41.40	0.19				
662543	**Acceptable identification**						**662543**
					NOV		
Staphylococcus simulans		93.78	0.30	GAL 8	1		
Staphylococcus cohnii subsp. urealyticum		6.23	0.12	SUC 96	99		
662550	**Low discrimination**						**662550**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		62.75	0.34	ARG 1	99	99	
Staphylococcus warneri		37.25	0.39		1	8	
662552	**Low discrimination**						**662552**
					NOV	PYR	
Staphylococcus simulans		63.30	0.32	GAL 8	1	70	
Staphylococcus saprophyticus subsp. bovis		20.49	0.22	SUC 96	99	99	
Staphylococcus warneri		16.22	0.29		1	8	
662553	**Low discrimination**						**662553**
					NOV		
Staphylococcus simulans		81.01	0.43	GAL 8	1		
Staphylococcus cohnii subsp. urealyticum		13.59	0.32	SUC 96	99		
Staphylococcus xylosus		3.44	0.11		99		
662573	**Good identification**						**662573**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		97.97	0.33	ARG 1	99	
Staphylococcus simulans		2.03	0.02	SUC 95	1	
662600	**Low discrimination**					**662600**
					YEP	
Staphylococcus warneri		65.82	0.52	MLT 95	43	
Staphylococcus simulans		30.38	0.36		1	
Staphylococcus pasteuri		3.10	0.10		86	
662601	**Acceptable identification**					**662601**
Staphylococcus simulans		92.83	0.47	LAC 99		
Staphylococcus hyicus		5.94	0.15			
Staphylococcus warneri		1.23	0.18			
662602	**Very good identification**					**662602**
					YEP	
Staphylococcus simulans		99.21	0.77		1	
Staphylococcus warneri		0.72	0.43		43	
Staphylococcus pasteuri		0.04	0.02		86	
662603	**Very good identification**					**662603**
Staphylococcus simulans		99.26	0.88			
Staphylococcus hyicus		0.74	0.37			
Staphylococcus felis		0.01	-.10			
662610	**Low discrimination**					**662610**
					YEP	
Staphylococcus simulans		64.55	0.48	LAC 99	1	
Staphylococcus warneri		25.32	0.49		43	
Staphylococcus pasteuri		10.13	0.27		86	
662611	**Very good identification**					**662611**
Staphylococcus simulans		99.76	0.59	LAC 99		
Staphylococcus warneri		0.24	0.15			
662612	**Very good identification**					**662612**
					YEP	
Staphylococcus simulans		99.80	0.89		1	
Staphylococcus warneri		0.13	0.40		43	
Staphylococcus pasteuri		0.07	0.19		86	
662613	**Excellent identification**					**662613**
Staphylococcus simulans		99.99	1.00			
Staphylococcus felis		0.01	0.07			
Staphylococcus hyicus		0.00	-.04			
662622	**Good identification**					**662622**
Staphylococcus simulans		100.0	0.36	XYL 1		
662623	**Good identification**					**662623**
Staphylococcus simulans		99.26	0.47	XYL 1		
Staphylococcus hyicus		0.74	-.04			
662632	**Good identification**					**662632**
Staphylococcus simulans		100.0	0.48	XYL 1		
662633	**Very good identification**					**662633**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus simulans		98.90	0.59	XYL 1	1		
Staphylococcus xylosus		1.10	0.15		99		
662640	**Very good identification**						**662640**
					PYR		
Staphylococcus warneri		95.29	0.79		8		
Staphylococcus petrasii subsp. croceilyticus		1.85	0.18		90		
Staphylococcus pasteuri		1.45	0.27		1		
662641	**Low discrimination**						**662641**
					PYR		
Staphylococcus simulans		62.57	0.39	LAC 99	70		
Staphylococcus warneri		34.95	0.44		8		
Staphylococcus petrasii subsp. petrasii		2.49	0.05		99		
662642	**Low discrimination**						**662642**
					YEP		
Staphylococcus simulans		76.01	0.70		1		
Staphylococcus warneri		23.45	0.69		43		
Staphylococcus pasteuri		0.44	0.19		86		
662643	**Very good identification**						**662643**
					PYR		
Staphylococcus simulans		99.77	0.81		70		
Staphylococcus warneri		0.19	0.34		8		
Staphylococcus petrasii subsp. petrasii		0.02	0.00		99		
662650	**Low discrimination**						**662650**
					YEP		
Staphylococcus warneri		81.99	0.76		43		
Staphylococcus pasteuri		10.60	0.43		86		
Staphylococcus simulans		4.94	0.41		1		
662651	**Low discrimination**						**662651**
					NOV	PYR	
Staphylococcus simulans		86.78	0.52	LAC 99	1	70	
Staphylococcus warneri		8.78	0.41		1	8	
Staphylococcus saprophyticus subsp. bovis		3.01	0.22		99	99	
662652	**Good identification**						**662652**
					YEP		
Staphylococcus simulans		93.80	0.82		1		
Staphylococcus warneri		5.24	0.66		43		
Staphylococcus pasteuri		0.83	0.35		86		
662653	**Very good identification**						**662653**
					NOV		
Staphylococcus simulans		99.92	0.93		1		
Staphylococcus warneri		0.03	0.31		1		
Staphylococcus xylosus		0.03	0.16		99		
662660	**Good identification**						**662660**
Staphylococcus warneri		100.0	0.38	XYL 1			
662662	**Low discrimination**						**662662**
Staphylococcus simulans		76.42	0.29	XYL 1			
Staphylococcus warneri		23.58	0.28				
662663	**Acceptable identification**						**662663**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		92.25	0.39	XYL 1	1		
Staphylococcus xylosus		7.75	0.14		99		
662670	**Low discrimination**						**662670**
					YEP		
Staphylococcus warneri		84.06	0.35	XYL 1	43		
Staphylococcus pasteuri		10.87	0.02		86		
Staphylococcus simulans		5.07	0.00		1		
662671	**Acceptable identification**						**662671**
					NOV		
Staphylococcus xylosus		91.96	0.29	ARG 1	99		
Staphylococcus simulans		8.04	0.11	GAL 91	1		
662672	**Acceptable identification**						**662672**
					NOV		
Staphylococcus simulans		93.72	0.41	XYL 1	1		
Staphylococcus warneri		5.24	0.25		1		
Staphylococcus xylosus		1.04	-.03		99		
662673	**Low discrimination**						**662673**
					NOV		
Staphylococcus simulans		75.25	0.52	XYL 1	1		
Staphylococcus xylosus		24.75	0.38		99		
662700	**Low discrimination**						**662700**
Staphylococcus warneri		85.95	0.42	MLT 95			
Staphylococcus simulans		10.92	0.14				
Staphylococcus hyicus		3.13	-.04				
662701	**Low discrimination**						**662701**
					COA	PYR	
Staphylococcus hyicus		87.82	0.37	BGA 10	24	1	
Staphylococcus simulans		10.38	0.25	PHS 92	1	70	
Staphylococcus intermedius		1.80	0.07	LAC 92	99	90	
662702	**Good identification**						**662702**
Staphylococcus simulans		94.36	0.55	GAL 8			
Staphylococcus hyicus		3.14	0.17				
Staphylococcus warneri		2.50	0.32				
662703	**Low discrimination**						**662703**
					COA	PYR	
Staphylococcus simulans		50.03	0.66	GAL 8	1	70	
Staphylococcus hyicus		49.19	0.59		24	1	
Staphylococcus intermedius		0.54	0.24		99	90	
662710	**Low discrimination**						**662710**
					NOV	PYR	
Staphylococcus warneri		56.50	0.39	MLT 95	1	8	
Staphylococcus simulans		39.66	0.26		1	70	
Staphylococcus saprophyticus subsp. bovis		3.85	0.05		99	99	
662711	**Low discrimination**						**662711**
					COA	NOV	
Staphylococcus simulans		84.87	0.37	GAL 8	1	1	
Staphylococcus intermedius		9.97	0.16	LAC 99	99	1	
Staphylococcus xylosus		3.35	0.05		1	99	
662712	**Very good identification**						**662712**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		99.52	0.67	GAL 8			
Staphylococcus warneri		0.48	0.29				
662713	**Very good identification**						**662713**
					COA		
Staphylococcus simulans		98.09	0.78	GAL 8	1		
Staphylococcus intermedius		0.72	0.32		99		
Staphylococcus condimenti		0.51	0.18		1		
662731	**Good identification**						**662731**
Staphylococcus xylosus		100.0	0.26	ARG 1			
662733	**Low discrimination**						**662733**
					NOV		
Staphylococcus xylosus		56.39	0.36	ARG 1	99		
Staphylococcus simulans		43.61	0.37	MLT 93	1		
662740	**Very good identification**						**662740**
					PYR		
Staphylococcus warneri		98.56	0.68		8		
Staphylococcus haemolyticus		0.89	0.19		98		
Staphylococcus simulans		0.30	0.07		70		
662741	**Low discrimination**						**662741**
					COA	PYR	
Staphylococcus warneri		46.09	0.34	MNS 2	1	8	
Staphylococcus simulans		22.72	0.18		1	70	
Staphylococcus intermedius		13.18	0.11		99	90	
662742	**Low discrimination**						**662742**
					PYR		
Staphylococcus warneri		52.42	0.59		8		
Staphylococcus simulans		46.79	0.48		70		
Staphylococcus haemolyticus		0.80	0.14		98		
662743	**Good identification**						**662743**
					COA	PYR	
Staphylococcus simulans		92.46	0.59	GAL 8	1	70	
Staphylococcus intermedius		3.33	0.27		99	90	
Staphylococcus hyicus		2.04	0.17		24	1	
662750	**Low discrimination**						**662750**
					NOV	PYR	
Staphylococcus warneri		72.33	0.65		1	8	
Staphylococcus saprophyticus subsp. bovis		25.65	0.47		99	99	
Staphylococcus simulans		1.20	0.19		1	70	
662751	**Low discrimination**						**662751**
					NOV		
Staphylococcus saprophyticus subsp. bovis		35.27	0.34	ARG 1	99		
Staphylococcus xylosus		25.83	0.28		99		
Staphylococcus simulans		22.11	0.30		1		
662752	**Low discrimination**						**662752**
					NOV	PYR	
Staphylococcus simulans		79.34	0.60	GAL 8	1	70	
Staphylococcus warneri		16.09	0.56		1	8	
Staphylococcus saprophyticus subsp. bovis		4.28	0.34		99	99	
662753	**Good identification**						**662753**
					COA	NOV	
Staphylococcus simulans		94.02	0.71	GAL 8	1	1	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus xylosus		3.16	0.37		1	99	
Staphylococcus intermedius		2.29	0.36		99	1	
662760	**Good identification**						**662760**
Staphylococcus warneri		100.0	0.27	XYL 1			
662763	**Acceptable identification**						**662763**
					NOV		
Staphylococcus xylosus		90.71	0.34	ARG 1	99		
Staphylococcus simulans		9.29	0.18	MAN 94	1		
662771	**Good identification**						**662771**
Staphylococcus xylosus		99.48	0.50	ARG 1			
Staphylococcus equorum		0.52	0.04				
662773	**Very good identification**						**662773**
					NOV		
Staphylococcus xylosus		97.01	0.59	ARG 1	99		
Staphylococcus simulans		2.54	0.30		1		
Staphylococcus equorum		0.46	0.12		99		
663213	**Good identification**						**663213**
Staphylococcus simulans		100.0	0.33	ESL 1			
663413	**Acceptable identification**						**663413**
					NOV		
Staphylococcus simulans		90.59	0.31	ESL 1	1		
Staphylococcus cohnii subsp. urealyticum		9.41	0.16	SUC 96	99		
663453	**Low discrimination**						**663453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		56.73	0.31	ARG 1	99		
Staphylococcus simulans		43.27	0.23		1		
663602	**Good identification**						**663602**
Staphylococcus simulans		97.44	0.36	ESL 1			
Staphylococcus piscifermentans		2.56	0.14				
663603	**Good identification**						**663603**
Staphylococcus simulans		99.26	0.47	ESL 1			
Staphylococcus hyicus		0.74	-.04				
663612	**Good identification**						**663612**
Staphylococcus simulans		100.0	0.48	ESL 1			
663613	**Very good identification**						**663613**
Staphylococcus simulans		100.0	0.59	ESL 1			
663640	**Good identification**						**663640**
Staphylococcus warneri		95.96	0.38	ESL 1			
Staphylococcus piscifermentans		4.04	0.11				
663642	**Low discrimination**						**663642**
Staphylococcus simulans		73.16	0.29	ESL 1			
Staphylococcus warneri		22.57	0.28				
Staphylococcus piscifermentans		4.28	0.14				

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
663643	**Good identification**					**663643**
Staphylococcus simulans		100.0	0.39	ESL 1		
663650	**Low discrimination**					**663650**
					YEP	
Staphylococcus warneri		84.06	0.35	ESL 1	43	
Staphylococcus pasteuri		10.87	0.02		86	
Staphylococcus simulans		5.07	0.00		1	
663652	**Acceptable identification**					**663652**
Staphylococcus simulans		94.71	0.41	ESL 1		
Staphylococcus warneri		5.29	0.25			
663653	**Good identification**					**663653**
					NOV	
Staphylococcus simulans		99.29	0.52	ESL 1	1	
Staphylococcus xylosus		0.71	0.04		99	
663673	**Low discrimination**					**663673**
					NOV	
Staphylococcus equorum		62.44	0.33	ARG 1	99	
Staphylococcus xylosus		33.45	0.26	GAL 91	99	
Staphylococcus simulans		4.11	0.11		1	
663713	**Good identification**					**663713**
					NOV	
Staphylococcus simulans		97.27	0.37	ESL 1	1	
Staphylococcus xylosus		2.73	0.02	GAL 8	99	
663733	**Low discrimination**					**663733**
Staphylococcus equorum		71.04	0.33	ARG 1		
Staphylococcus xylosus		28.96	0.23	MLT 91		
663740	**Low discrimination**					**663740**
Staphylococcus warneri		83.34	0.27	ESL 1		
Staphylococcus piscifermentans		16.66	0.14			
663753	**Low discrimination**					**663753**
					NOV	
Staphylococcus simulans		49.71	0.30	ESL 1	1	
Staphylococcus xylosus		41.33	0.25	GAL 8	99	
Staphylococcus equorum		8.96	0.12		99	
663771	**Low discrimination**					**663771**
Staphylococcus equorum		66.49	0.45	ARG 1		
Staphylococcus xylosus		32.18	0.37			
Staphylococcus gallinarum		1.33	0.07			
663773	**Low discrimination**					**663773**
Staphylococcus equorum		64.32	0.54	ARG 1		
Staphylococcus xylosus		34.46	0.47			
Staphylococcus gallinarum		1.23	0.15			
664213	**Good identification**					**664213**
Staphylococcus simulans		100.0	0.35	NIT 96		
664253	**Good identification**					**664253**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		100.0	0.28	NIT 96			
664413	**Low discrimination**						**664413**
					NOV		
Staphylococcus cohnii subsp. urealyticum		60.69	0.43	ARG 1	99		
Staphylococcus simulans		36.74	0.33		1		
Staphylococcus felis		2.57	0.01		1		
664443	**Low discrimination**						**664443**
					NOV		
Staphylococcus cohnii subsp. urealyticum		89.18	0.38	ARG 1	99		
Staphylococcus simulans		10.82	0.14	MAN 91	1		
664450	**Acceptable identification**						**664450**
					NOV		
Staphylococcus kloosii		92.99	0.47	ARG 1	99		
Staphylococcus warneri		7.01	0.16		1		
664451	**Good identification**						**664451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	ARG 1			
664452	**Low discrimination**						**664452**
					NOV	YEP	
Staphylococcus kloosii		75.18	0.48	ARG 1	99	99	
Staphylococcus simulans		14.56	0.15		1	1	
Staphylococcus cohnii subsp. urealyticum		10.26	0.18		99	64	
664453	**Very good identification**						**664453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		95.42	0.59	ARG 1	99		
Staphylococcus simulans		4.58	0.26		1		
664470	**Good identification**						**664470**
Staphylococcus kloosii		100.0	0.34	ARG 1			
664472	**Good identification**						**664472**
Staphylococcus kloosii		100.0	0.35	ARG 1			
664512	**Low discrimination**						**664512**
					NOV	YEP	
Staphylococcus kloosii		73.16	0.27	ARG 1	99	99	
Staphylococcus simulans		26.84	0.01	MLT 95	1	1	
664513	**Low discrimination**						**664513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		44.89	0.36	ARG 1	99		
Staphylococcus condimenti		33.03	0.14		1		
Staphylococcus felis		14.86	0.12		1		
664543	**Good identification**						**664543**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1			
664550	**Good identification**						**664550**
Staphylococcus kloosii		100.0	0.53	ARG 1			
664551	**Good identification**						**664551**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1			
664552	**Very good identification**						**664552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		97.08	0.53	ARG 1			
Staphylococcus cohnii subsp. urealyticum		2.92	0.10				
664553	**Good identification**						**664553**
					COA	NOV	
Staphylococcus cohnii subsp. urealyticum		98.06	0.51	ARG 1	1	99	
Staphylococcus intermedius		0.99	0.03		99	1	
Staphylococcus simulans		0.96	0.04		1	1	
664570	**Good identification**						**664570**
Staphylococcus kloosii		100.0	0.40	ARG 1			
664572	**Good identification**						**664572**
Staphylococcus kloosii		100.0	0.41	ARG 1			
664602	**Good identification**						**664602**
Staphylococcus simulans		98.78	0.39	NIT 96			
Staphylococcus warneri		1.22	0.09				
664603	**Acceptable identification**						**664603**
Staphylococcus simulans		93.83	0.49	NIT 96			
Staphylococcus hyicus		5.83	0.17				
Staphylococcus felis		0.33	-.10				
664612	**Good identification**						**664612**
Staphylococcus simulans		100.0	0.51	NIT 96			
664613	**Very good identification**						**664613**
					NOV		
Staphylococcus simulans		99.09	0.62	NIT 96	1		
Staphylococcus felis		0.59	0.07		1		
Staphylococcus cohnii subsp. urealyticum		0.21	0.12		99		
664640	**Good identification**						**664640**
Staphylococcus warneri		100.0	0.45	PHS 1			
664642	**Low discrimination**						**664642**
Staphylococcus simulans		65.64	0.31	NIT 96			
Staphylococcus warneri		34.36	0.35				
664643	**Good identification**						**664643**
					NOV		
Staphylococcus simulans		98.95	0.42	NIT 96	1		
Staphylococcus cohnii subsp. urealyticum		1.05	0.07		99		
664650	**Acceptable identification**						**664650**
					NOV	YEP	
Staphylococcus warneri		90.53	0.42	PHS 1	1	43	
Staphylococcus kloosii		6.25	0.26		99	99	
Staphylococcus simulans		3.22	0.03		1	1	
664652	**Low discrimination**						**664652**
					NOV	YEP	
Staphylococcus simulans		89.60	0.44	NIT 96	1	1	
Staphylococcus warneri		8.49	0.32		1	43	
Staphylococcus kloosii		1.91	0.27		99	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
664653	**Very good identification**						**664653**
					COA	NOV	
Staphylococcus simulans		97.03	0.55	NIT 96	1	1	
Staphylococcus cohnii subsp. urealyticum		2.61	0.28		1	99	
Staphylococcus intermedius		0.36	0.03		99	1	
664703	**Low discrimination**						**664703**
					COA	PYR	
Staphylococcus hyicus		71.78	0.39	BGA 10	24	1	
Staphylococcus intermedius		16.49	0.32	NIT 99	99	90	
Staphylococcus simulans		8.73	0.28		1	70	
664712	**Good identification**						**664712**
					COA		
Staphylococcus simulans		95.86	0.29	NIT 96	1		
Staphylococcus intermedius		4.14	-.01	GAL 8	99		
664713	**Low discrimination**						**664713**
					COA		
Staphylococcus intermedius		43.65	0.40	GLR 3	99		
Staphylococcus simulans		34.19	0.40	NIT 93	1		
Staphylococcus condimenti		12.80	0.18		1		
664740	**Good identification**						**664740**
Staphylococcus warneri		100.0	0.35	PHS 1			
664743	**Low discrimination**						**664743**
					COA	YEP	
Staphylococcus intermedius		79.29	0.35	GLR 3	99	1	
Staphylococcus simulans		12.57	0.20	NIT 93	1	1	
Staphylococcus chromogenes		5.82	0.10		1	95	
664750	**Low discrimination**						**664750**
					NOV		
Staphylococcus warneri		70.19	0.32	PHS 1	1		
Staphylococcus kloosii		29.81	0.32		99		
664751	**Good identification**						**664751**
					COA	NOV	
Staphylococcus intermedius		95.91	0.28	GLR 3	99	1	
Staphylococcus xylosus		4.09	-.02	NIT 93	1	99	
664753	**Low discrimination**						**664753**
					COA	NOV	
Staphylococcus intermedius		77.34	0.44	GLR 3	99	1	
Staphylococcus simulans		18.14	0.33	NIT 93	1	1	
Staphylococcus cohnii subsp. urealyticum		2.40	0.20		1	99	
664773	**Low discrimination**						**664773**
					COA	NOV	
Staphylococcus xylosus		87.33	0.29	ARG 1	1	99	
Staphylococcus arlettae		8.80	0.18	NIT 89	1	99	
Staphylococcus intermedius		3.88	0.03		99	1	
665413	**Good identification**						**665413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ARG 1			
665443	**Good identification**						**665443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ARG 1			
665450	**Good identification**						**665450**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus kloosii		100.0	0.47	ARG 1			
665451	**Good identification**						**665451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ARG 1			
665452	**Acceptable identification**						**665452**
Staphylococcus kloosii		94.96	0.48	ARG 1			
Staphylococcus cohnii subsp. urealyticum		5.04	0.09				
665453	**Good identification**						**665453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.51	ARG 1			
665470	**Good identification**						**665470**
Staphylococcus kloosii		100.0	0.34	ARG 1			
665472	**Good identification**						**665472**
Staphylococcus kloosii		100.0	0.35	ARG 1			
665512	**Good identification**						**665512**
Staphylococcus kloosii		100.0	0.27	ARG 1			
665513	**Good identification**						**665513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1			
665550	**Good identification**						**665550**
Staphylococcus kloosii		100.0	0.53	ARG 1			
665552	**Very good identification**						**665552**
Staphylococcus kloosii		100.0	0.53	ARG 1			
665553	**Good identification**						**665553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ARG 1			
665570	**Good identification**						**665570**
Staphylococcus kloosii		100.0	0.40	ARG 1			
665572	**Good identification**						**665572**
Staphylococcus kloosii		100.0	0.41	ARG 1			
665652	**Low discrimination**						**665652**
					NOV	YEP	
Staphylococcus kloosii		67.81	0.27	ARG 1	99	99	
Staphylococcus simulans		32.19	0.03	SUC 9	1	1	
665750	**Good identification**						**665750**
Staphylococcus kloosii		100.0	0.32	ARG 1			
665752	**Good identification**						**665752**
Staphylococcus kloosii		100.0	0.33	ARG 1			
665771	**Low discrimination**						**665771**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus equorum		77.94	0.27	ARG 1				
Staphylococcus gallinarum		11.97	0.07	NIT 92				
Staphylococcus xylosus		10.08	0.07					
665773	**Low discrimination**							**665773**
Staphylococcus equorum		74.23	0.35	ARG 1				
Staphylococcus gallinarum		10.86	0.15	NIT 92				
Staphylococcus xylosus		10.63	0.17					
666013	**Low discrimination**							**666013**
Staphylococcus simulans		84.56	0.35	SUC 96				
Staphylococcus carnosus subsp. carnosus		10.75	0.14	TRE 95				
Staphylococcus felis		4.69	0.01					
666053	**Good identification**							**666053**
Staphylococcus simulans		100.0	0.28	SUC 96				
666202	**Good identification**							**666202**
Staphylococcus simulans		98.59	0.41	TRE 95				
Staphylococcus hyicus		1.41	-.04					
666203	**Low discrimination**							**666203**
					YEP			
Staphylococcus simulans		69.43	0.52	TRE 95	1			
Staphylococcus hyicus		29.42	0.37		1			
Staphylococcus chromogenes		0.95	0.10		95			
666212	**Very good identification**							**666212**
Staphylococcus simulans		100.0	0.53	TRE 95				
666213	**Very good identification**							**666213**
Staphylococcus simulans		99.25	0.64	TRE 95				
Staphylococcus felis		0.47	0.07					
Staphylococcus hyicus		0.11	-.04					
666242	**Good identification**							**666242**
Staphylococcus simulans		100.0	0.34	TRE 95				
666243	**Good identification**							**666243**
					YEP			
Staphylococcus simulans		95.34	0.44	TRE 95	1			
Staphylococcus chromogenes		2.91	0.10		95			
Staphylococcus hyicus		0.91	-.04		1			
666252	**Good identification**							**666252**
Staphylococcus simulans		100.0	0.46	TRE 95				
666253	**Good identification**							**666253**
					COA			
Staphylococcus simulans		94.97	0.57	TRE 95	1			
Staphylococcus delphini		4.54	0.18		99			
Staphylococcus intermedius		0.28	0.03		99			
666301	**Acceptable identification**							**666301**
					YEP	PYR	COA	
Staphylococcus hyicus		92.40	0.37	BGA 10	1	1	24	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus intermedius		4.62	0.15	TRE 92	1	90	99	
Staphylococcus chromogenes		2.99	0.10	LAC 92	95	45	1	
666303	**Good identification**							**666303**
					YEP	PYR	COA	
Staphylococcus hyicus		93.19	0.59	BGA 10	1	1	24	
Staphylococcus intermedius		2.49	0.32	TRE 92	1	90	99	
Staphylococcus chromogenes		2.36	0.30		95	45	1	
666312	**Good identification**							**666312**
					COA			
Staphylococcus simulans		96.70	0.31	GAL 8	1			
Staphylococcus intermedius		3.31	-.01	TRE 95	99			
666313	**Low discrimination**							**666313**
					COA			
Staphylococcus intermedius		36.81	0.40	GLR 3	99			
Staphylococcus simulans		36.42	0.42	TRE 93	1			
Staphylococcus condimenti		10.79	0.18		1			
666343	**Low discrimination**							**666343**
					YEP	PYR	COA	
Staphylococcus intermedius		45.45	0.35	GLR 3	1	90	99	
Staphylococcus chromogenes		28.72	0.30	TRE 93	95	45	1	
Staphylococcus hyicus		11.47	0.17		1	1	24	
666351	**Good identification**							**666351**
					COA	YEP		
Staphylococcus intermedius		95.69	0.28	GLR 3	99	1		
Staphylococcus chromogenes		4.31	-.01	TRE 93	1	95		
666353	**Low discrimination**							**666353**
					COA			
Staphylococcus intermedius		65.28	0.44	GLR 3	99			
Staphylococcus simulans		19.34	0.35	TRE 93	1			
Staphylococcus delphini		10.63	0.18		99			
666402	**Good identification**							**666402**
Staphylococcus simulans		100.0	0.39	SUC 96				
666403	**Low discrimination**							**666403**
Staphylococcus simulans		80.67	0.49	SUC 96				
Staphylococcus felis		13.92	0.25					
Staphylococcus hyicus		5.02	0.17					
666412	**Good identification**							**666412**
Staphylococcus simulans		99.03	0.51	SUC 96				
Staphylococcus felis		0.97	0.01					
666413	**Low discrimination**							**666413**
Staphylococcus simulans		75.45	0.62	SUC 96				
Staphylococcus felis		21.79	0.42					
Staphylococcus condimenti		1.64	0.14					
666442	**Good identification**							**666442**
Staphylococcus simulans		100.0	0.31	SUC 96				
666443	**Acceptable identification**							**666443**
					NOV			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		94.29	0.42	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		5.72	0.23		99		
666452	**Good identification**						**666452**
Staphylococcus simulans		100.0	0.44	SUC 96			
666453	**Low discrimination**						**666453**
					NOV		
Staphylococcus simulans		86.23	0.55	SUC 96	1		
Staphylococcus cohnii subsp. urealyticum		13.21	0.43		99		
Staphylococcus felis		0.56	0.01		1		
666503	**Low discrimination**						**666503**
Staphylococcus hyicus		47.46	0.39	BGA 10			
Staphylococcus felis		38.35	0.36	SUC 99			
Staphylococcus simulans		5.77	0.28				
666512	**Low discrimination**						**666512**
Staphylococcus simulans		45.22	0.29	GAL 8			
Staphylococcus condimenti		37.78	0.14	SUC 96			
Staphylococcus felis		17.00	0.12				
666513	**Low discrimination**						**666513**
Staphylococcus condimenti		66.61	0.55	GLR 1			
Staphylococcus felis		29.97	0.53				
Staphylococcus simulans		2.70	0.40				
666553	**Low discrimination**						**666553**
					COA	NOV	
Staphylococcus simulans		31.24	0.33	GAL 8	1	1	
Staphylococcus cohnii subsp. urealyticum		23.59	0.36	SUC 96	1	99	
Staphylococcus intermedius		17.87	0.26		99	1	
666601	**Low discrimination**						**666601**
					YEP		
Staphylococcus hyicus		66.33	0.37	BGA 10	1		
Staphylococcus simulans		30.04	0.37	GAL 92	1		
Staphylococcus chromogenes		2.74	0.12	LAC 92	95		
666602	**Very good identification**						**666602**
Staphylococcus simulans		99.09	0.67				
Staphylococcus hyicus		0.86	0.17				
Staphylococcus felis		0.05	-.10				
666603	**Low discrimination**						**666603**
Staphylococcus simulans		78.11	0.78				
Staphylococcus hyicus		20.03	0.59				
Staphylococcus felis		1.14	0.31				
666610	**Good identification**						**666610**
Staphylococcus simulans		100.0	0.38	LAC 99			
666611	**Good identification**						**666611**
					COA		
Staphylococcus simulans		95.56	0.49	LAC 99	1		
Staphylococcus felis		2.33	0.07		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus intermedius		0.96	0.06		99		
666612	**Very good identification**						**666612**
Staphylococcus simulans		99.92	0.79				
Staphylococcus felis		0.08	0.07				
666613	**Very good identification**						**666613**
Staphylococcus simulans		97.31	0.90				
Staphylococcus felis		2.38	0.48				
Staphylococcus condimenti		0.13	0.18				
666623	**Low discrimination**						**666623**
Staphylococcus simulans		78.67	0.37	XYL 1			
Staphylococcus hyicus		20.18	0.17				
Staphylococcus felis		1.15	-.10				
666632	**Good identification**						**666632**
Staphylococcus simulans		100.0	0.38	XYL 1			
666633	**Good identification**						**666633**
					NOV		
Staphylococcus simulans		96.73	0.49	XYL 1	1		
Staphylococcus felis		2.36	0.07		1		
Staphylococcus xylosus		0.91	0.04		99		
666640	**Low discrimination**						**666640**
Staphylococcus warneri		73.52	0.38	PHS 1			
Staphylococcus simulans		26.48	0.19				
666641	**Low discrimination**						**666641**
					YEP		
Staphylococcus simulans		76.80	0.30	LAC 99	1		
Staphylococcus chromogenes		15.59	0.12		95		
Staphylococcus hyicus		3.81	-.04		1		
666642	**Very good identification**						**666642**
Staphylococcus simulans		99.07	0.60				
Staphylococcus warneri		0.93	0.28				
666643	**Very good identification**						**666643**
					YEP		
Staphylococcus simulans		97.31	0.71		1		
Staphylococcus chromogenes		1.80	0.32		95		
Staphylococcus hyicus		0.56	0.17		1		
666650	**Low discrimination**						**666650**
					NOV	PYR	
Staphylococcus simulans		52.19	0.31	LAC 99	1	70	
Staphylococcus warneri		26.24	0.35		1	8	
Staphylococcus saprophyticus subsp. bovis		18.18	0.22		99	99	
666651	**Acceptable identification**						**666651**
					COA	NOV	
Staphylococcus simulans		92.40	0.42	LAC 99	1	1	
Staphylococcus intermedius		3.09	0.10		99	1	
Staphylococcus saprophyticus subsp. bovis		2.40	0.09		1	99	
666652	**Very good identification**						**666652**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	PYR		
Staphylococcus simulans		99.74	0.72		1	70		
Staphylococcus warneri		0.17	0.25		1	8		
Staphylococcus saprophyticus subsp. bovis		0.09	0.09		99	99		
666653	**Very good identification**							**666653**
					COA	YEP		
Staphylococcus simulans		99.55	0.83		1	1		
Staphylococcus intermedius		0.21	0.26		99	1		
Staphylococcus chromogenes		0.13	0.20		1	95		
666663	**Acceptable identification**							**666663**
					NOV			
Staphylococcus simulans		93.37	0.30	XYL 1	1			
Staphylococcus xylosus		6.63	0.02		99			
666672	**Good identification**							**666672**
Staphylococcus simulans		100.0	0.31	XYL 1				
666673	**Low discrimination**							**666673**
					NOV			
Staphylococcus simulans		78.23	0.42	XYL 1	1			
Staphylococcus xylosus		21.77	0.27		99			
666701	**Good identification**							**666701**
					YEP	PYR	COA	
Staphylococcus hyicus		91.29	0.59	BGA 10	1	1	24	
Staphylococcus intermedius		5.27	0.38	LAC 92	1	90	99	
Staphylococcus chromogenes		2.95	0.32		95	45	1	
666702	**Low discrimination**							**666702**
					COA	PYR		
Staphylococcus hyicus		51.43	0.39	BGA 10	24	1		
Staphylococcus simulans		44.84	0.45	MNS 99	1	70		
Staphylococcus intermedius		1.59	0.14		99	90		
666703	**Good identification**							**666703**
					COA	PYR		
Staphylococcus hyicus		90.58	0.80	BGA 10	24	1		
Staphylococcus intermedius		2.79	0.55		99	90		
Staphylococcus simulans		2.67	0.56		1	70		
666711	**Low discrimination**							**666711**
					COA			
Staphylococcus intermedius		69.98	0.47	GLR 3	99			
Staphylococcus condimenti		9.49	0.18	LAC 86	1			
Staphylococcus simulans		6.14	0.27		1			
666712	**Low discrimination**							**666712**
Staphylococcus simulans		89.86	0.58	GAL 8				
Staphylococcus condimenti		4.69	0.18					
Staphylococcus felis		2.86	0.18					
666713	**Low discrimination**							**666713**
Staphylococcus condimenti		35.59	0.59	GLR 1				
Staphylococcus simulans		23.05	0.68					
Staphylococcus felis		21.68	0.59					
666723	**Acceptable identification**							**666723**
					COA	PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus hyicus		90.06	0.39	BGA 10	24	1		
Staphylococcus intermedius		2.78	0.14	XYL 1	99	90		
Staphylococcus simulans		2.66	0.15		1	70		
666740	**Low discrimination**							**666740**
					COA	PYR		
Staphylococcus warneri		84.13	0.27	PHS 1	1	8		
Staphylococcus intermedius		15.88	0.01		99	90		
666741	**Low discrimination**							**666741**
					YEP	PYR	COA	
Staphylococcus intermedius		66.32	0.42	GLR 3	1	90	99	
Staphylococcus chromogenes		24.76	0.32	LAC 86	95	45	1	
Staphylococcus hyicus		7.74	0.17		1	1	24	
666742	**Low discrimination**							**666742**
					COA	YEP		
Staphylococcus simulans		80.49	0.38	GAL 8	1	1		
Staphylococcus intermedius		9.50	0.17		99	1		
Staphylococcus chromogenes		5.20	0.10		1	95		
666743	**Low discrimination**							**666743**
					COA	YEP		
Staphylococcus intermedius		48.60	0.58	GLR 3	99	1		
Staphylococcus chromogenes		26.59	0.51		1	95		
Staphylococcus simulans		13.92	0.49		1	1		
666750	**Low discrimination**							**666750**
					COA	NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		79.58	0.34	ARG 1	1	99	99	
Staphylococcus warneri		9.07	0.24		1	1	8	
Staphylococcus intermedius		6.39	0.10		99	1	90	
666751	**Low discrimination**							**666751**
					COA	NOV	YEP	
Staphylococcus intermedius		89.07	0.51	GLR 3	99	1	1	
Staphylococcus chromogenes		3.47	0.20	LAC 86	1	1	95	
Staphylococcus saprophyticus subsp. bovis		2.80	0.22		1	99	80	
666752	**Low discrimination**							**666752**
					COA	NOV		
Staphylococcus simulans		88.42	0.50	GAL 8	1	1		
Staphylococcus intermedius		7.06	0.26		99	1		
Staphylococcus saprophyticus subsp. bovis		3.58	0.22		1	99		
666753	**Low discrimination**							**666753**
					COA	YEP		
Staphylococcus intermedius		64.49	0.67	GLR 3	99	1		
Staphylococcus simulans		27.33	0.61		1	1		
Staphylococcus chromogenes		3.68	0.40		1	95		
666771	**Acceptable identification**							**666771**
					COA	NOV		
Staphylococcus xylosus		91.06	0.38	ARG 1	1	99		
Staphylococcus gallinarum		3.34	0.07		1	99		
Staphylococcus intermedius		3.08	0.10		99	1		
666773	**Low discrimination**							**666773**
					COA	PYR	NOV	
Staphylococcus xylosus		81.72	0.48	ARG 1	1	65	99	
Staphylococcus intermedius		5.96	0.26		99	90	1	
Staphylococcus lutrae		4.82	0.18		99	10	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
667453	**Low discrimination**						**667453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		85.51	0.35	ARG 1	99		
Staphylococcus simulans		14.50	0.14		1		
667553	**Good identification**						**667553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1			
667603	**Low discrimination**						**667603**
Staphylococcus simulans		78.67	0.37	ESL 1			
Staphylococcus hyicus		20.18	0.17				
Staphylococcus felis		1.15	-.10				
667612	**Good identification**						**667612**
Staphylococcus simulans		100.0	0.38	ESL 1			
667613	**Good identification**						**667613**
Staphylococcus simulans		97.62	0.49	ESL 1			
Staphylococcus felis		2.38	0.07				
667643	**Good identification**						**667643**
Staphylococcus simulans		100.0	0.30	ESL 1			
667652	**Good identification**						**667652**
Staphylococcus simulans		100.0	0.31	ESL 1			
667653	**Good identification**						**667653**
Staphylococcus simulans		100.0	0.42	ESL 1			
667671	**Low discrimination**						**667671**
Staphylococcus equorum		81.56	0.28	ARG 1			
Staphylococcus gallinarum		11.02	0.07	GAL 91			
Staphylococcus xylosus		7.42	0.05				
667673	**Low discrimination**						**667673**
Staphylococcus equorum		80.39	0.36	ARG 1			
Staphylococcus gallinarum		10.34	0.15	GAL 91			
Staphylococcus xylosus		8.10	0.15				
667703	**Acceptable identification**						**667703**
					COA	PYR	
Staphylococcus hyicus		90.73	0.39	BGA 10	24	1	
Staphylococcus intermedius		2.80	0.14	ESL 1	99	90	
Staphylococcus simulans		2.68	0.15		1	70	
667731	**Low discrimination**						**667731**
Staphylococcus equorum		83.03	0.28	ARG 1			
Staphylococcus gallinarum		11.22	0.07	MLT 91			
Staphylococcus xylosus		5.75	0.03				
667733	**Low discrimination**						**667733**
Staphylococcus equorum		82.97	0.36	ARG 1			
Staphylococcus gallinarum		10.67	0.15	MLT 91			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus xylosus		6.36	0.12			
667761	**Low discrimination**					**667761**
Staphylococcus gallinarum		89.16	0.28	ARG 1		
Staphylococcus equorum		6.89	0.08	MAN 91		
Staphylococcus xylosus		3.96	0.01			
667763	**Low discrimination**					**667763**
Staphylococcus gallinarum		88.28	0.36	ARG 1		
Staphylococcus equorum		7.16	0.16	MAN 91		
Staphylococcus xylosus		4.56	0.11			
667771	**Low discrimination**					**667771**
Staphylococcus gallinarum		54.80	0.48	ARG 1		
Staphylococcus equorum		41.43	0.49			
Staphylococcus xylosus		3.77	0.26			
667773	**Low discrimination**					**667773**
Staphylococcus gallinarum		53.36	0.56	ARG 1		
Staphylococcus equorum		42.37	0.57			
Staphylococcus xylosus		4.27	0.35			
670613	**Good identification**					**670613**
Staphylococcus simulans		100.0	0.31	ORN 1		
670640	**Good identification**					**670640**
Staphylococcus warneri		100.0	0.45	ORN 1		
670642	**Acceptable identification**					**670642**
Staphylococcus warneri		94.53	0.35	ORN 1		
Staphylococcus simulans		5.47	0.00			
670650	**Good identification**					**670650**
Staphylococcus warneri		100.0	0.42	ORN 1		
670652	**Low discrimination**					**670652**
Staphylococcus warneri		75.77	0.32	ORN 1		
Staphylococcus simulans		24.23	0.13			
670740	**Good identification**					**670740**
Staphylococcus warneri		100.0	0.35	ORN 1		
670750	**Good identification**					**670750**
Staphylococcus warneri		100.0	0.32	ORN 1		
672213	**Good identification**					**672213**
Staphylococcus simulans		100.0	0.33	ORN 1		
672413	**Good identification**					**672413**
Staphylococcus simulans		100.0	0.31	ORN 1		
672602	**Good identification**					**672602**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		100.0	0.36	ORN 1		
672603	**Good identification**					**672603**
Staphylococcus simulans		99.26	0.47	ORN 1		
Staphylococcus hyicus		0.74	-.04			
672612	**Good identification**					**672612**
Staphylococcus simulans		100.0	0.48	ORN 1		
672613	**Very good identification**					**672613**
Staphylococcus simulans		100.0	0.59	ORN 1		
672640	**Good identification**					**672640**
Staphylococcus warneri		100.0	0.38	ORN 1		
672642	**Low discrimination**					**672642**
Staphylococcus simulans		76.42	0.29	ORN 1		
Staphylococcus warneri		23.58	0.28			
672643	**Good identification**					**672643**
Staphylococcus simulans		100.0	0.39	ORN 1		
672650	**Low discrimination**					**672650**
					YEP	
Staphylococcus warneri		84.06	0.35	ORN 1	43	
Staphylococcus pasteuri		10.87	0.02		86	
Staphylococcus simulans		5.07	0.00		1	
672652	**Acceptable identification**					**672652**
Staphylococcus simulans		94.71	0.41	ORN 1		
Staphylococcus warneri		5.29	0.25			
672653	**Good identification**					**672653**
Staphylococcus simulans		100.0	0.52	ORN 1		
672713	**Good identification**					**672713**
Staphylococcus simulans		100.0	0.37	ORN 1		
672740	**Good identification**					**672740**
Staphylococcus warneri		100.0	0.27	ORN 1		
672753	**Good identification**					**672753**
					NOV	
Staphylococcus simulans		96.75	0.30	ORN 1	1	
Staphylococcus xylosus		3.25	-.04	GAL 8	99	
676603	**Low discrimination**					**676603**
Staphylococcus simulans		78.67	0.37	ORN 1		
Staphylococcus hyicus		20.18	0.17			
Staphylococcus felis		1.15	-.10			
676612	**Good identification**					**676612**
Staphylococcus simulans		100.0	0.38	ORN 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
676613	**Good identification**						**676613**
Staphylococcus simulans		97.62	0.49	ORN 1			
Staphylococcus felis		2.38	0.07				
676643	**Good identification**						**676643**
Staphylococcus simulans		100.0	0.30	ORN 1			
676652	**Good identification**						**676652**
Staphylococcus simulans		100.0	0.31	ORN 1			
676653	**Good identification**						**676653**
Staphylococcus simulans		100.0	0.42	ORN 1			
676703	**Acceptable identification**						**676703**
					COA	PYR	
Staphylococcus hyicus		90.73	0.39	ORN 1	24	1	
Staphylococcus intermedius		2.80	0.14	BGA 10	99	90	
Staphylococcus simulans		2.68	0.15		1	70	
700000	**Low discrimination**						**700000**
					OXI		
Dermacoccus nishinomiyaensis		65.47	0.33	VPT 5	93		
Micrococcus sp.		20.63	0.13	ARG 1	89		
Kocuria varians		11.68	0.25		13		
700011	**Good identification**						**700011**
Staphylococcus capitis subsp. capitis		95.13	0.35	URE 1			
Staphylococcus capitis subsp. ureolyticus		4.87	-.01	NIT 89			
700040	**Low discrimination**						**700040**
					OXI		
Staphylococcus warneri		79.92	0.35	SUC 95	1		
Kocuria varians		12.50	0.11	TRE 95	13		
Micrococcus sp.		7.58	-.10		89		
700050	**Low discrimination**						**700050**
					OXI	NOV	
Staphylococcus warneri		67.56	0.32	SUC 95	1	1	
Macrococcus equipersicus		19.27	0.11	TRE 95	99	99	
Macrococcus bovicus		8.05	0.02		99	99	
700051	**Good identification**						**700051**
Staphylococcus capitis subsp. ureolyticus		100.0	0.28	NIT 95			
700053	**Good identification**						**700053**
Staphylococcus capitis subsp. ureolyticus		100.0	0.45	NIT 95			
700100	**Low discrimination**						**700100**
					OXI		
Kocuria varians		58.41	0.29	VPT 14	13		
Dermacoccus nishinomiyaensis		41.59	0.18	ARG 1	93		
700200	**Low discrimination**						**700200**
					OXI		
Staphylococcus warneri		67.71	0.35	TRE 95	1		
Dermacoccus nishinomiyaensis		17.27	0.08	MLT 95	93		
Micrococcus sp.		8.43	-.08		89		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
700210	**Low discrimination**						**700210**
Staphylococcus warneri		83.99	0.32	TRE 95			
Staphylococcus capitis subsp. capitis		9.64	-.01	MLT 95			
Staphylococcus capitis subsp. ureolyticus		6.38	-.14				
700211	**Low discrimination**						**700211**
Staphylococcus capitis subsp. capitis		60.18	0.40	URE 1			
Staphylococcus capitis subsp. ureolyticus		39.83	0.28	NIT 89			
700213	**Good identification**						**700213**
Staphylococcus capitis subsp. ureolyticus		99.58	0.45	NIT 95			
Staphylococcus capitis subsp. capitis		0.42	0.05	MLT 96			
700240	**Low discrimination**						**700240**
					NOV	PYR	
Staphylococcus warneri		70.64	0.61	TRE 95	1	8	
Staphylococcus hominis subsp. novobiosepticus		13.32	0.30		99	1	
Staphylococcus petrasii subsp. petrasii		3.48	0.20		1	99	
700241	**Low discrimination**						**700241**
					PYR		
Staphylococcus petrasii subsp. petrasii		50.07	0.39	NIT 90	99		
Staphylococcus capitis subsp. ureolyticus		32.39	0.28	TRE 99	1		
Staphylococcus epidermidis		7.22	0.28		1		
700242	**Low discrimination**						**700242**
					NOV		
Staphylococcus warneri		50.38	0.52	TRE 95	1		
Staphylococcus epidermidis		16.46	0.32		1		
Staphylococcus hominis subsp. novobiosepticus		15.35	0.24		99		
700243	**Low discrimination**						**700243**
					PYR		
Staphylococcus capitis subsp. ureolyticus		81.44	0.45	NIT 95	1		
Staphylococcus petrasii subsp. petrasii		9.24	0.34	MAN 96	99		
Staphylococcus epidermidis		7.84	0.38		1		
700250	**Low discrimination**						**700250**
					NOV		
Staphylococcus warneri		85.79	0.58	TRE 95	1		
Staphylococcus capitis subsp. ureolyticus		8.23	0.15		1		
Staphylococcus saprophyticus subsp. saprophyticus		2.52	0.06		99		
700251	**Very good identification**						**700251**
					PYR		
Staphylococcus capitis subsp. ureolyticus		99.02	0.56	NIT 95	1		
Staphylococcus petrasii subsp. petrasii		0.71	0.20	LAC 88	99		
Staphylococcus warneri		0.21	0.24		8		
700252	**Low discrimination**						**700252**
					NOV		
Staphylococcus capitis subsp. ureolyticus		57.25	0.33	NIT 95	1		
Staphylococcus warneri		27.13	0.49	MNS 99	1		
Staphylococcus saprophyticus subsp. saprophyticus		13.56	0.22		99		
700253	**Very good identification**						**700253**
					PYR		
Staphylococcus capitis subsp. ureolyticus		99.94	0.74	NIT 95	1		
Staphylococcus petrasii subsp. petrasii		0.05	0.14		99		

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
Staphylococcus warneri		0.01	0.14		8			
700273	**Good identification**							**700273**
Staphylococcus capitis subsp. ureolyticus		100.0	0.33	NIT 95				
700340	**Low discrimination**							**700340**
					NOV			
Staphylococcus hominis subsp. novobiosepticus		48.69	0.39	ARG 1	99			
Staphylococcus warneri		27.17	0.51	NIT 90	1			
Staphylococcus epidermidis		16.76	0.38		1			
700341	**Low discrimination**							**700341**
					PYR	NOV		
Staphylococcus epidermidis		44.35	0.43	NIT 97	1	1		
Staphylococcus petrasii subsp. petrasii		36.19	0.36		99	1		
Staphylococcus hominis subsp. novobiosepticus		17.34	0.27		1	99		
700342	**Low discrimination**							**700342**
					NOV			
Staphylococcus epidermidis		55.74	0.48	NIT 97	1			
Staphylococcus hominis subsp. novobiosepticus		27.53	0.34		99			
Staphylococcus warneri		9.51	0.41		1			
700343	**Low discrimination**							**700343**
					PYR	NOV		
Staphylococcus epidermidis		81.72	0.53	NIT 97	1	1		
Staphylococcus petrasii subsp. petrasii		11.34	0.30		99	1		
Staphylococcus hominis subsp. novobiosepticus		5.43	0.21		1	99		
700350	**Acceptable identification**							**700350**
					NOV			
Staphylococcus warneri		92.98	0.48	TRE 95	1			
Staphylococcus hominis subsp. hominis		3.22	0.12		1			
Staphylococcus hominis subsp. novobiosepticus		2.32	-.02		99			
700352	**Low discrimination**							**700352**
					NOV			
Staphylococcus warneri		66.21	0.38	TRE 95	1			
Staphylococcus saprophyticus subsp. saprophyticus		9.11	0.00		99			
Staphylococcus hominis subsp. hominis		8.07	0.14		1			
700353	**Low discrimination**							**700353**
					PYR			
Staphylococcus capitis subsp. ureolyticus		88.93	0.33	NIT 95	1			
Staphylococcus caprae		5.96	0.11	GAL 1	75			
Staphylococcus petrasii subsp. petrasii		3.08	0.11		99			
700400	**Low discrimination**							**700400**
Staphylococcus warneri		56.19	0.35	SUC 95				
Staphylococcus carnosus subsp. utilis		24.40	0.03	MLT 95				
Staphylococcus auricularis		13.39	0.14					
700410	**Low discrimination**							**700410**
					OXI	NOV	YEP	
Staphylococcus warneri		78.37	0.32	SUC 95	1	1	43	
Micrococcus equipersicus		8.27	0.02	MLT 95	99	99	99	
Staphylococcus cohnii subsp. cohnii		7.36	0.06		1	99	12	
700440	**Low discrimination**							**700440**
					PYR			
Staphylococcus warneri		84.19	0.61	SUC 95	8			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		4.14	0.20		99		
Staphylococcus petrasii subsp. jettensis		3.65	0.10		99		
700441	**Acceptable identification**						**700441**
					NOV	PYR	
Staphylococcus petrasii subsp. petrasii		92.12	0.39	NIT 90	1	99	
Staphylococcus warneri		4.25	0.27	SUC 99	1	8	
Staphylococcus cohnii subsp. cohnii		2.01	0.15		99	1	
700442	**Low discrimination**						**700442**
					NOV	PYR	
Staphylococcus warneri		79.37	0.52	SUC 95	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		11.72	0.14		99	7	
Staphylococcus petrasii subsp. petrasii		6.31	0.14		1	99	
700443	**Good identification**						**700443**
					PYR		
Staphylococcus petrasii subsp. petrasii		97.23	0.34	NIT 90	99		
Staphylococcus warneri		2.77	0.17	SUC 99	8		
700450	**Low discrimination**						**700450**
					NOV		
Staphylococcus warneri		73.62	0.58	SUC 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		21.20	0.27		99		
Staphylococcus pasteuri		1.67	0.10		1		
700452	**Low discrimination**						**700452**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		82.48	0.42	ARG 8	99		
Staphylococcus warneri		16.85	0.49	SUC 91	1		
Staphylococcus pasteuri		0.47	0.02		1		
700540	**Low discrimination**						**700540**
					PYR		
Staphylococcus warneri		89.75	0.51	SUC 95	8		
Staphylococcus petrasii subsp. petrasii		9.32	0.16		99		
Staphylococcus auricularis		0.92	0.02		40		
700541	**Good identification**						**700541**
					PYR		
Staphylococcus petrasii subsp. petrasii		97.86	0.36	NIT 90	99		
Staphylococcus warneri		2.14	0.16	SUC 99	8		
700542	**Low discrimination**						**700542**
					PYR		
Staphylococcus warneri		79.04	0.41	SUC 95	8		
Staphylococcus petrasii subsp. petrasii		13.26	0.11		99		
Staphylococcus caprae		4.48	-.05		75		
700543	**Low discrimination**						**700543**
Staphylococcus petrasii subsp. petrasii		78.42	0.30	NIT 90			
Staphylococcus caprae		21.58	0.13	SUC 99			
700550	**Low discrimination**						**700550**
					NOV		
Staphylococcus warneri		86.62	0.48	SUC 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		6.87	0.05		99		
Staphylococcus caprae		2.26	-.05		1		
700553	**Good identification**						**700553**
Staphylococcus caprae		97.92	0.39	NIT 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		2.08	0.11	PHS 99			
700600	**Low discrimination**						**700600**
					PYR		
Staphylococcus warneri		83.63	0.61	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		4.12	0.20		99		
Staphylococcus petrasii subsp. jettensis		3.63	0.10		99		
700601	**Low discrimination**						**700601**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		86.26	0.39	NIT 90	1	99	
Kocuria kristinae		9.77	0.20	MLT 99	99	93	
Staphylococcus warneri		3.98	0.27		1	8	
700602	**Low discrimination**						**700602**
					PYR	YEP	
Staphylococcus warneri		85.44	0.52	MLT 95	8	43	
Staphylococcus petrasii subsp. petrasii		6.79	0.14		99	1	
Staphylococcus pasteuri		4.52	0.11		1	86	
700603	**Low discrimination**						**700603**
					OXI		
Staphylococcus petrasii subsp. petrasii		89.99	0.34	NIT 90	1		
Kocuria kristinae		4.73	0.07	MLT 99	99		
Staphylococcus simulans		2.71	0.08		1		
700610	**Low discrimination**						**700610**
					NOV		
Staphylococcus warneri		71.26	0.58	MLT 95	1		
Staphylococcus pasteuri		26.10	0.35		1		
Staphylococcus saprophyticus subsp. saprophyticus		2.10	0.06		99		
700612	**Low discrimination**						**700612**
					NOV		
Staphylococcus warneri		50.09	0.49	MLT 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		25.04	0.22		99		
Staphylococcus pasteuri		22.48	0.27		1		
700640	**Low discrimination**						**700640**
					PYR		
Staphylococcus warneri		56.71	0.88		8		
Staphylococcus petrasii subsp. petrasii		14.54	0.61		99		
Staphylococcus petrasii subsp. jettensis		12.81	0.51		99		
700641	**Very good identification**						**700641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		97.90	0.80	NIT 90	1	99	
Kocuria kristinae		1.01	0.39		99	93	
Staphylococcus warneri		0.87	0.53		1	8	
700642	**Low discrimination**						**700642**
					PYR		
Staphylococcus warneri		59.96	0.78		8		
Staphylococcus petrasii subsp. petrasii		24.84	0.55		99		
Staphylococcus petrasii subsp. jettensis		7.17	0.36		99		
700643	**Very good identification**						**700643**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		98.82	0.75	NIT 90	1	99	
Staphylococcus warneri		0.54	0.43		1	8	
Kocuria kristinae		0.47	0.27		99	93	
700650	**Low discrimination**						**700650**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus warneri		75.76	0.85		1		
Staphylococcus saprophyticus subsp. saprophyticus		11.61	0.47		99		
Staphylococcus pasteuri		8.97	0.51		1		
700651	**Good identification**						**700651**
					PYR		
Staphylococcus petrasii subsp. petrasii		92.29	0.61	NIT 90	99		
Staphylococcus warneri		5.32	0.50	MAN 10	8		
Staphylococcus capitis subsp. ureolyticus		1.32	0.15		1		
700652	**Low discrimination**						**700652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		68.11	0.63	ARG 8	99		
Staphylococcus warneri		26.15	0.75		1		
Staphylococcus pasteuri		3.80	0.43		1		
700653	**Low discrimination**						**700653**
					NOV	PYR	
Staphylococcus petrasii subsp. petrasii		77.68	0.55	NIT 90	1	99	
Staphylococcus capitis subsp. ureolyticus		15.08	0.33	MAN 10	1	1	
Staphylococcus saprophyticus subsp. saprophyticus		3.57	0.22		99	7	
700660	**Low discrimination**						**700660**
					PYR		
Staphylococcus warneri		57.66	0.47	XYL 1	8		
Staphylococcus petrasii subsp. petrasii		14.78	0.20		99		
Staphylococcus petrasii subsp. jettensis		13.03	0.10		99		
700661	**Good identification**						**700661**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		98.12	0.39	NIT 90	1	99	
Kocuria kristinae		1.01	-.02	XYL 1	99	93	
Staphylococcus warneri		0.87	0.12		1	8	
700662	**Low discrimination**						**700662**
					PYR		
Staphylococcus warneri		62.02	0.37	XYL 1	8		
Staphylococcus petrasii subsp. petrasii		25.69	0.14		99		
Staphylococcus petrasii subsp. jettensis		7.42	-.05		99		
700663	**Good identification**						**700663**
Staphylococcus petrasii subsp. petrasii		100.0	0.34	NIT 90			
700670	**Low discrimination**						**700670**
					NOV		
Staphylococcus warneri		76.28	0.44	XYL 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		11.69	0.06		99		
Staphylococcus pasteuri		9.03	0.10		1		
700700	**Acceptable identification**						**700700**
					PYR		
Staphylococcus warneri		90.59	0.51	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		9.41	0.16		99		
700701	**Acceptable identification**						**700701**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		92.73	0.36	NIT 90	1	99	
Kocuria kristinae		5.25	0.10	MLT 99	99	93	
Staphylococcus warneri		2.02	0.16		1	8	
700702	**Low discrimination**						**700702**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR		
Staphylococcus warneri		85.63	0.41	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		14.37	0.11		99		
700703	**Good identification**						**700703**
					OXI		
Staphylococcus petrasii subsp. petrasii		97.44	0.30	NIT 90	1		
Kocuria kristinae		2.56	-.03	MLT 99	99		
700710	**Good identification**						**700710**
					PYR		
Staphylococcus warneri		98.43	0.48	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		1.57	-.04		99		
700712	**Low discrimination**						**700712**
					NOV		
Staphylococcus warneri		87.90	0.38	MLT 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		12.10	0.00		99		
700740	**Low discrimination**						**700740**
					PYR		
Staphylococcus warneri		62.89	0.77		8		
Staphylococcus petrasii subsp. petrasii		34.04	0.57		99		
Staphylococcus hominis subsp. hominis		1.76	0.40		1		
700741	**Very good identification**						**700741**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		99.05	0.77	NIT 90	1	99	
Kocuria kristinae		0.51	0.30		99	93	
Staphylococcus warneri		0.42	0.43		1	8	
700742	**Low discrimination**						**700742**
					PYR		
Staphylococcus warneri		49.81	0.68		8		
Staphylococcus petrasii subsp. petrasii		43.55	0.52		99		
Staphylococcus hominis subsp. hominis		4.90	0.41		1		
700743	**Very good identification**						**700743**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		99.31	0.71	NIT 90	1	99	
Staphylococcus warneri		0.26	0.33		1	8	
Kocuria kristinae		0.24	0.17		99	93	
700750	**Low discrimination**						**700750**
					NOV	PYR	
Staphylococcus warneri		87.87	0.74		1	8	
Staphylococcus petrasii subsp. petrasii		7.30	0.37		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		3.71	0.25		99	7	
700751	**Very good identification**						**700751**
					PYR		
Staphylococcus petrasii subsp. petrasii		96.69	0.57	NIT 90	99		
Staphylococcus warneri		2.64	0.40	MAN 10	8		
Staphylococcus caprae		0.48	0.04		75		
700752	**Low discrimination**						**700752**
					NOV	PYR	
Staphylococcus warneri		52.38	0.65		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		37.57	0.41		99	7	
Staphylococcus petrasii subsp. petrasii		7.03	0.32		1	99	
700753	**Low discrimination**						**700753**
					COA	PYR	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		78.31	0.52	NIT 90	1	99	
Staphylococcus caprae		13.77	0.30	MAN 10	1	75	
Staphylococcus aureus subsp. aureus		5.90	0.18		99	10	
700760	**Low discrimination**						**700760**
					PYR		
Staphylococcus warneri		64.88	0.36	XYL 1	8		
Staphylococcus petrasii subsp. petrasii		35.12	0.16		99		
700761	**Good identification**						**700761**
Staphylococcus petrasii subsp. petrasii		100.0	0.36	NIT 90			
700763	**Good identification**						**700763**
Staphylococcus petrasii subsp. petrasii		100.0	0.30	NIT 90			
700770	**Acceptable identification**						**700770**
					PYR		
Staphylococcus warneri		92.33	0.33	XYL 1	8		
Staphylococcus petrasii subsp. petrasii		7.67	-.04		99		
701253	**Good identification**						**701253**
Staphylococcus capitis subsp. ureolyticus		100.0	0.33	ESL 1			
701640	**Low discrimination**						**701640**
					PYR		
Staphylococcus warneri		57.16	0.47	ESL 1	8		
Staphylococcus petrasii subsp. petrasii		14.65	0.20		99		
Staphylococcus petrasii subsp. jettensis		12.91	0.10		99		
701641	**Low discrimination**						**701641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		79.14	0.39	ESL 1	1	99	
Kocuria kristinae		20.16	0.27	NIT 90	99	93	
Staphylococcus warneri		0.70	0.12		1	8	
701642	**Low discrimination**						**701642**
					PYR		
Staphylococcus warneri		59.47	0.37	ESL 1	8		
Staphylococcus petrasii subsp. petrasii		24.63	0.14		99		
Staphylococcus petrasii subsp. jettensis		7.11	-.05		99		
701643	**Low discrimination**						**701643**
					OXI		
Staphylococcus petrasii subsp. petrasii		89.43	0.34	ESL 1	1		
Kocuria kristinae		10.58	0.15	NIT 90	99		
701650	**Low discrimination**						**701650**
					NOV		
Staphylococcus warneri		76.28	0.44	ESL 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		11.69	0.06		99		
Staphylococcus pasteuri		9.03	0.10		1		
701740	**Low discrimination**						**701740**
					PYR		
Staphylococcus warneri		61.95	0.36	ESL 1	8		
Staphylococcus petrasii subsp. petrasii		33.53	0.16		99		
Staphylococcus piscifermentans		4.52	0.14		15		
701741	**Low discrimination**						**701741**
					OXI		
Staphylococcus petrasii subsp. petrasii		88.70	0.36	ESL 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Kocuria kristinae		11.30	0.17	NIT 90	99	
701743	**Acceptable identification**					**701743**
					OXI	
Staphylococcus petrasii subsp. petrasii		94.42	0.30	ESL 1	1	
Kocuria kristinae		5.58	0.05	NIT 90	99	
701750	**Acceptable identification**					**701750**
					PYR	
Staphylococcus warneri		92.33	0.33	ESL 1	8	
Staphylococcus petrasii subsp. petrasii		7.67	-.04		99	
702000	**Low discrimination**					**702000**
					OXI	
Kocuria varians		52.44	0.39	VPT 14	13	
Dermacoccus nishinomiyaensis		33.88	0.28	ARG 1	93	
Staphylococcus carnosus subsp. utilis		6.54	0.03		1	
702001	**Low discrimination**					**702001**
Staphylococcus capitis subsp. capitis		81.53	0.30	URE 1		
Kocuria varians		9.15	0.17	MAN 93		
Staphylococcus capitis subsp. ureolyticus		5.43	-.03			
702002	**Good identification**					**702002**
Kocuria varians		100.0	0.28	VPT 14		
702011	**Low discrimination**					**702011**
Staphylococcus capitis subsp. capitis		89.27	0.53	URE 1		
Staphylococcus capitis subsp. ureolyticus		10.74	0.25			
702013	**Good identification**					**702013**
					COA	
Staphylococcus capitis subsp. ureolyticus		96.98	0.43	SUC 96	1	
Staphylococcus capitis subsp. capitis		2.24	0.19	MLT 96	1	
Staphylococcus schleiferi subsp. coagulans		0.57	0.12		99	
702020	**Acceptable identification**					**702020**
Kocuria varians		92.82	0.30	VPT 14		
Kocuria rosea		7.18	-.02	ARG 1		
702040	**Low discrimination**					**702040**
Staphylococcus hominis subsp. hominis		29.21	0.29	ARG 12		
Kocuria varians		24.01	0.26	SUC 99		
Staphylococcus auricularis		13.68	0.19			
702042	**Low discrimination**					**702042**
Staphylococcus hominis subsp. hominis		41.64	0.30	ARG 12		
Staphylococcus epidermidis		25.80	0.22	SUC 99		
Staphylococcus capitis subsp. ureolyticus		12.19	0.02			
702043	**Acceptable identification**					**702043**
					PYR	
Staphylococcus capitis subsp. ureolyticus		94.99	0.43	SUC 96	1	
Staphylococcus epidermidis		3.77	0.28	MAN 96	1	
Staphylococcus petrasii subsp. petrasii		1.24	0.12		99	
702051	**Very good identification**					**702051**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. ureolyticus		99.57	0.54	SUC 96	1	
Staphylococcus capitis subsp. capitis		0.35	0.12	LAC 88	1	
Staphylococcus petrasii subsp. petrasii		0.08	-.02		99	
702052	**Good identification**					**702052**
Staphylococcus capitis subsp. ureolyticus		96.64	0.31	SUC 96		
Staphylococcus hominis subsp. hominis		2.06	0.13	MNS 99		
Staphylococcus warneri		1.31	0.15			
702053	**Very good identification**					**702053**
Staphylococcus capitis subsp. ureolyticus		99.98	0.72	SUC 96		
Staphylococcus caprae		0.02	-.02			
702073	**Good identification**					**702073**
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	SUC 96		
702100	**Acceptable identification**					**702100**
					OXI	
Kocuria varians		92.42	0.43	VPT 14	13	
Dermacoccus nishinomiyaensis		7.58	0.13	ARG 1	93	
702101	**Low discrimination**					**702101**
					COA	
Staphylococcus schleiferi subsp. coagulans		84.82	0.37	PHS 99	99	
Kocuria varians		11.83	0.21	LAC 86	1	
Staphylococcus schleiferi subsp. schleiferi		3.36	0.04		1	
702102	**Low discrimination**					**702102**
					COA	
Kocuria varians		87.93	0.31	VPT 14	1	
Staphylococcus schleiferi subsp. coagulans		12.07	0.12	ARG 1	99	
702103	**Very good identification**					**702103**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.01	0.53	PHS 99	99	
Kocuria varians		0.63	0.10		1	
Staphylococcus epidermidis		0.36	0.02		1	
702111	**Low discrimination**					**702111**
					COA	
Staphylococcus schleiferi subsp. coagulans		89.25	0.36	PHS 99	99	
Staphylococcus capitis subsp. capitis		10.75	0.12	LAC 86	1	
702113	**Good identification**					**702113**
					COA	
Staphylococcus schleiferi subsp. coagulans		97.47	0.53	PHS 99	99	
Staphylococcus capitis subsp. ureolyticus		1.69	0.02		1	
Staphylococcus caprae		0.84	-.02		1	
702120	**Good identification**					**702120**
Kocuria varians		100.0	0.33	VPT 14		
702140	**Low discrimination**					**702140**
Staphylococcus epidermidis		26.69	0.28	SUC 99		
Kocuria varians		25.29	0.29			
Staphylococcus hominis subsp. hominis		22.21	0.29			
702141	**Low discrimination**					**702141**
					PYR NOV	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		72.45	0.33	SUC 99	1	1	
Staphylococcus petrasii subsp. petrasii		16.45	0.14		99	1	
Staphylococcus hominis subsp. novobiosepticus		7.88	0.05		1	99	
702142	**Low discrimination**						**702142**
					NOV		
Staphylococcus epidermidis		64.10	0.38	SUC 99	1		
Staphylococcus hominis subsp. hominis		19.78	0.31		1		
Staphylococcus hominis subsp. novobiosepticus		8.81	0.12		99		
702143	**Low discrimination**						**702143**
Staphylococcus epidermidis		80.17	0.43	SUC 99			
Staphylococcus caprae		9.37	0.13				
Staphylococcus capitis subsp. ureolyticus		3.60	0.02				
702153	**Low discrimination**						**702153**
					COA		
Staphylococcus caprae		66.60	0.39	PHS 99	1		
Staphylococcus capitis subsp. ureolyticus		32.31	0.31	TRE 90	1		
Staphylococcus schleiferi subsp. coagulans		0.79	0.12		99		
702200	**Low discrimination**						**702200**
Staphylococcus hominis subsp. hominis		33.52	0.29	ARG 12			
Kocuria varians		17.12	0.21	MLT 99			
Staphylococcus warneri		15.46	0.27				
702201	**Low discrimination**						**702201**
					PYR		
Staphylococcus capitis subsp. capitis		47.43	0.36	URE 1	1		
Staphylococcus capitis subsp. ureolyticus		40.81	0.25	MAN 93	1		
Staphylococcus petrasii subsp. petrasii		7.24	0.18		99		
702202	**Low discrimination**						**702202**
Staphylococcus hominis subsp. hominis		42.12	0.30	ARG 12			
Staphylococcus epidermidis		26.10	0.22	MLT 99			
Staphylococcus capitis subsp. ureolyticus		12.33	0.02				
702203	**Acceptable identification**						**702203**
					PYR		
Staphylococcus capitis subsp. ureolyticus		94.03	0.43	MAN 96	1		
Staphylococcus epidermidis		3.73	0.28	MLT 96	1		
Staphylococcus petrasii subsp. petrasii		1.23	0.12		99		
702211	**Low discrimination**						**702211**
					PYR		
Staphylococcus capitis subsp. ureolyticus		60.82	0.54	MLT 96	1		
Staphylococcus capitis subsp. capitis		39.13	0.59	LAC 88	1		
Staphylococcus petrasii subsp. petrasii		0.05	-.02		99		
702212	**Acceptable identification**						**702212**
Staphylococcus capitis subsp. ureolyticus		94.19	0.31	MLT 96			
Staphylococcus simulans		2.53	0.11	MNS 99			
Staphylococcus hominis subsp. hominis		2.00	0.13				
702213	**Very good identification**						**702213**
Staphylococcus capitis subsp. ureolyticus		99.72	0.72	MLT 96			
Staphylococcus capitis subsp. capitis		0.18	0.24				

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		0.09	0.22			
702233	**Good identification**					**702233**
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	XYL 1		
702240	**Low discrimination**					**702240**
					NOV	
Staphylococcus hominis subsp. hominis		54.02	0.70	ARG 12	1	
Staphylococcus hominis subsp. novobiosepticus		18.95	0.49		99	
Staphylococcus epidermidis		12.41	0.53		1	
702241	**Low discrimination**					**702241**
					PYR	
Staphylococcus capitis subsp. ureolyticus		45.25	0.54	MAN 96	1	
Staphylococcus petrasii subsp. petrasii		33.13	0.59	LAC 88	99	
Staphylococcus epidermidis		17.16	0.59		1	
702242	**Low discrimination**					**702242**
					NOV	
Staphylococcus hominis subsp. hominis		51.74	0.71	ARG 12	1	
Staphylococcus epidermidis		32.05	0.64		1	
Staphylococcus hominis subsp. novobiosepticus		8.32	0.44		99	
702243	**Low discrimination**					**702243**
					PYR	
Staphylococcus capitis subsp. ureolyticus		81.54	0.72	MAN 96	1	
Staphylococcus epidermidis		13.35	0.69		1	
Staphylococcus petrasii subsp. petrasii		4.38	0.53		99	
702250	**Low discrimination**					**702250**
Staphylococcus capitis subsp. ureolyticus		51.99	0.41	MNS 99		
Staphylococcus hominis subsp. hominis		28.49	0.53	LAC 88		
Staphylococcus warneri		12.23	0.51			
702251	**Very good identification**					**702251**
					PYR	
Staphylococcus capitis subsp. ureolyticus		99.60	0.82	LAC 88	1	
Staphylococcus petrasii subsp. petrasii		0.34	0.39		99	
Staphylococcus capitis subsp. capitis		0.03	0.18		1	
702252	**Low discrimination**					**702252**
Staphylococcus capitis subsp. ureolyticus		89.94	0.59	MNS 99		
Staphylococcus hominis subsp. hominis		7.89	0.54			
Staphylococcus warneri		0.96	0.41			
702253	**Excellent identification**					**702253**
					PYR	
Staphylococcus capitis subsp. ureolyticus		99.96	1.00		1	
Staphylococcus petrasii subsp. petrasii		0.03	0.34		99	
Staphylococcus epidermidis		0.01	0.28		1	
702260	**Low discrimination**					**702260**
					NOV	
Staphylococcus hominis subsp. hominis		56.81	0.29	ARG 12	1	
Staphylococcus hominis subsp. novobiosepticus		19.92	0.08	XYL 1	99	
Staphylococcus epidermidis		13.05	0.12		1	
702262	**Low discrimination**					**702262**
					NOV	
Staphylococcus hominis subsp. hominis		54.02	0.30	ARG 12	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		33.46	0.22	XYL 1	1		
Staphylococcus hominis subsp. novobiosepticus		8.69	0.03		99		
702263	**Low discrimination**						**702263**
					PYR		
Staphylococcus capitis subsp. ureolyticus		82.13	0.31	MAN 96	1		
Staphylococcus epidermidis		13.45	0.28	XYL 1	1		
Staphylococcus petrasii subsp. petrasii		4.42	0.12		99		
702271	**Good identification**						**702271**
					PYR		
Staphylococcus capitis subsp. ureolyticus		99.66	0.41	XYL 1	1		
Staphylococcus petrasii subsp. petrasii		0.34	-.02	LAC 88	99		
702273	**Very good identification**						**702273**
Staphylococcus capitis subsp. ureolyticus		100.0	0.59	XYL 1			
702300	**Low discrimination**						**702300**
					NOV		
Staphylococcus epidermidis		29.51	0.28	MLT 99	1		
Staphylococcus hominis subsp. hominis		24.57	0.29		1		
Staphylococcus hominis subsp. novobiosepticus		23.86	0.18		99		
702301	**Low discrimination**						**702301**
					COA	PYR	
Staphylococcus epidermidis		54.90	0.33	MLT 99	1	1	
Staphylococcus schleiferi subsp. coagulans		25.14	0.27		99	20	
Staphylococcus petrasii subsp. petrasii		12.47	0.14		1	99	
702302	**Low discrimination**						**702302**
					NOV		
Staphylococcus epidermidis		66.00	0.38	MLT 99	1		
Staphylococcus hominis subsp. hominis		20.37	0.31		1		
Staphylococcus hominis subsp. novobiosepticus		9.07	0.12		99		
702303	**Low discrimination**						**702303**
					COA		
Staphylococcus epidermidis		49.51	0.43	MLT 99	1		
Staphylococcus schleiferi subsp. coagulans		43.97	0.43		99		
Staphylococcus capitis subsp. ureolyticus		2.22	0.02		1		
702313	**Low discrimination**						**702313**
					COA		
Staphylococcus capitis subsp. ureolyticus		56.00	0.31	GAL 1	1		
Staphylococcus schleiferi subsp. coagulans		42.61	0.42	MLT 96	99		
Staphylococcus epidermidis		0.53	0.02		1		
702340	**Low discrimination**						**702340**
					NOV		
Staphylococcus epidermidis		36.62	0.69		1		
Staphylococcus hominis subsp. hominis		30.48	0.70		1		
Staphylococcus hominis subsp. novobiosepticus		29.60	0.59		99		
702341	**Low discrimination**						**702341**
					PYR	NOV	
Staphylococcus epidermidis		74.05	0.74		1	1	
Staphylococcus petrasii subsp. petrasii		16.82	0.55		99	1	
Staphylococcus hominis subsp. novobiosepticus		8.06	0.47		1	99	
702342	**Low discrimination**						**702342**
					NOV		
Staphylococcus epidermidis		68.49	0.79		1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		21.14	0.72		1		
Staphylococcus hominis subsp. novobiosepticus		9.42	0.53		99		
702343	**Good identification**						**702343**
					PYR	NOV	
Staphylococcus epidermidis		93.28	0.85		1	1	
Staphylococcus petrasii subsp. petrasii		3.60	0.50		99	1	
Staphylococcus hominis subsp. novobiosepticus		1.73	0.41		1	99	
702350	**Low discrimination**						**702350**
					PYR		
Staphylococcus hominis subsp. hominis		69.80	0.54	ARG 12	1		
Staphylococcus warneri		8.73	0.41	MAN 13	8		
Staphylococcus haemolyticus		6.77	0.31		98		
702351	**Low discrimination**						**702351**
					PYR		
Staphylococcus capitis subsp. ureolyticus		74.22	0.41	GAL 1	1		
Staphylococcus petrasii subsp. petrasii		16.60	0.36	LAC 88	99		
Staphylococcus epidermidis		6.65	0.33		1		
702352	**Low discrimination**						**702352**
Staphylococcus hominis subsp. hominis		66.57	0.55	ARG 12			
Staphylococcus epidermidis		14.58	0.38	MAN 13			
Staphylococcus capitis subsp. ureolyticus		7.07	0.18				
702353	**Low discrimination**						**702353**
Staphylococcus capitis subsp. ureolyticus		89.92	0.59	GAL 1			
Staphylococcus epidermidis		3.48	0.43				
Staphylococcus caprae		2.86	0.30				
702360	**Low discrimination**						**702360**
					NOV		
Staphylococcus epidermidis		36.59	0.28	XYL 1	1		
Staphylococcus hominis subsp. hominis		30.46	0.29		1		
Staphylococcus hominis subsp. novobiosepticus		29.58	0.18		99		
702361	**Low discrimination**						**702361**
					PYR	NOV	
Staphylococcus epidermidis		74.85	0.33	XYL 1	1	1	
Staphylococcus petrasii subsp. petrasii		17.00	0.14		99	1	
Staphylococcus hominis subsp. novobiosepticus		8.15	0.05		1	99	
702362	**Low discrimination**						**702362**
					NOV		
Staphylococcus epidermidis		69.15	0.38	XYL 1	1		
Staphylococcus hominis subsp. hominis		21.34	0.31		1		
Staphylococcus hominis subsp. novobiosepticus		9.51	0.12		99		
702363	**Acceptable identification**						**702363**
					PYR	NOV	
Staphylococcus epidermidis		93.63	0.43	XYL 1	1	1	
Staphylococcus petrasii subsp. petrasii		3.62	0.09		99	1	
Staphylococcus hominis subsp. novobiosepticus		1.73	0.00		1	99	
702402	**Low discrimination**						**702402**
Staphylococcus piscifermentans		57.53	0.32	VPT 1			
Staphylococcus auricularis		22.25	0.14	ESL 90			
Staphylococcus warneri		13.33	0.18				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
702440	**Low discrimination**						**702440**
					PYR		
Staphylococcus petrasii subsp. petrasii		28.17	0.39	SUC 99	99		
Staphylococcus warneri		27.26	0.54	MNS 90	8		
Staphylococcus auricularis		16.76	0.41		40		
702441	**Very good identification**						**702441**
					PYR		
Staphylococcus petrasii subsp. petrasii		98.98	0.59	SUC 99	99		
Staphylococcus auricularis		0.73	0.22		40		
Staphylococcus warneri		0.22	0.19		8		
702442	**Low discrimination**						**702442**
					PYR		
Staphylococcus petrasii subsp. petrasii		36.61	0.34	SUC 99	99		
Staphylococcus hominis subsp. hominis		27.91	0.40	MNS 90	1		
Staphylococcus warneri		21.93	0.44		8		
702443	**Very good identification**						**702443**
					PYR		
Staphylococcus petrasii subsp. petrasii		98.45	0.53	SUC 99	99		
Staphylococcus capitis subsp. ureolyticus		0.77	0.02		1		
Staphylococcus caprae		0.18	-.09		75		
702450	**Low discrimination**						**702450**
					PYR	YEP	
Staphylococcus warneri		66.78	0.51	SUC 95	8	43	
Staphylococcus petrasii subsp. petrasii		10.59	0.20		99	1	
Staphylococcus pasteuri		8.67	0.18		1	86	
702451	**Low discrimination**						**702451**
					PYR		
Staphylococcus petrasii subsp. petrasii		86.16	0.39	SUC 99	99		
Staphylococcus capitis subsp. ureolyticus		10.70	0.13	MAN 10	1		
Staphylococcus warneri		1.23	0.16		8		
702452	**Low discrimination**						**702452**
					PYR		
Staphylococcus warneri		45.26	0.41	SUC 95	8		
Staphylococcus hominis subsp. hominis		11.88	0.23		1		
Staphylococcus petrasii subsp. petrasii		11.59	0.14		99		
702453	**Low discrimination**						**702453**
					PYR		
Staphylococcus capitis subsp. ureolyticus		55.02	0.31	SUC 96	1		
Staphylococcus petrasii subsp. petrasii		32.54	0.34	TRE 1	99		
Staphylococcus caprae		10.31	0.18		75		
702500	**Low discrimination**						**702500**
Staphylococcus piscifermentans		42.82	0.32	VPT 1			
Kocuria varians		27.29	0.20	ESL 90			
Staphylococcus auricularis		16.56	0.14				
702502	**Low discrimination**						**702502**
Staphylococcus piscifermentans		89.30	0.36	VPT 1			
Kocuria varians		10.70	0.08	ESL 90			
702512	**Low discrimination**						**702512**
Staphylococcus piscifermentans		66.93	0.28	VPT 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests			Code
Staphylococcus caprae		33.07	0.00	ESL 90				
702540	**Low discrimination**							**702540**
					PYR			
Staphylococcus petrasii subsp. petrasii		40.69	0.36	SUC 99	99			
Staphylococcus hominis subsp. hominis		23.12	0.40	MNS 90	1			
Staphylococcus warneri		18.65	0.43		8			
702541	**Very good identification**							**702541**
					PYR			
Staphylococcus petrasii subsp. petrasii		98.88	0.55	SUC 99	99			
Staphylococcus caprae		0.77	0.06		75			
Staphylococcus hominis subsp. hominis		0.13	0.05		1			
702542	**Low discrimination**							**702542**
					OXI	PYR	NOV	
Staphylococcus hominis subsp. hominis		34.09	0.41	ARG 12	1	1	1	
Staphylococcus petrasii subsp. petrasii		27.52	0.30	SUC 99	1	99	1	
Macrococcus caseolyticus		10.60	0.18		99	80	99	
702543	**Low discrimination**							**702543**
					PYR			
Staphylococcus petrasii subsp. petrasii		78.09	0.50	SUC 99	99			
Staphylococcus caprae		21.49	0.33		75			
Staphylococcus hominis subsp. hominis		0.22	0.06		1			
702550	**Low discrimination**							**702550**
					NOV	PYR		
Staphylococcus warneri		29.93	0.41	SUC 95	1	8		
Staphylococcus saprophyticus subsp. bovis		28.63	0.31		99	99		
Staphylococcus caprae		16.40	0.15		1	75		
702551	**Low discrimination**							**702551**
					NOV			
Staphylococcus caprae		55.27	0.33	PHS 99	1			
Staphylococcus petrasii subsp. petrasii		41.44	0.36	LAC 95	1			
Staphylococcus saprophyticus subsp. bovis		3.29	0.18		99			
702552	**Low discrimination**							**702552**
Staphylococcus caprae		88.71	0.41	PHS 99				
Staphylococcus warneri		2.84	0.31	MNS 88				
Staphylococcus hominis subsp. hominis		2.56	0.24					
702553	**Very good identification**							**702553**
					COA	PYR		
Staphylococcus caprae		96.14	0.59	PHS 99	1	75		
Staphylococcus petrasii subsp. petrasii		2.04	0.30		1	99		
Staphylococcus aureus subsp. aureus		1.69	0.18		99	10		
702600	**Low discrimination**							**702600**
					PYR			
Staphylococcus petrasii subsp. petrasii		29.14	0.39	MLT 99	99			
Staphylococcus warneri		28.20	0.54	MNS 90	8			
Staphylococcus hominis subsp. hominis		10.19	0.39		1			
702601	**Very good identification**							**702601**
					PYR			
Staphylococcus petrasii subsp. petrasii		99.19	0.59	MLT 99	99			
Staphylococcus auricularis		0.29	0.13		40			
Staphylococcus warneri		0.22	0.19		8			
702602	**Low discrimination**							**702602**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR		
Staphylococcus petrasii subsp. petrasii		32.03	0.34	MLT 99	99		
Staphylococcus hominis subsp. hominis		24.41	0.40	MNS 90	1		
Staphylococcus warneri		19.19	0.44		8		
702603	**Good identification**						**702603**
					PYR		
Staphylococcus petrasii subsp. petrasii		91.18	0.53	MLT 99	99		
Staphylococcus simulans		7.32	0.36		70		
Staphylococcus capitis subsp. ureolyticus		0.71	0.02		1		
702610	**Low discrimination**						**702610**
					PYR	YEP	
Staphylococcus pasteuri		61.55	0.43	GLR 99	1	86	
Staphylococcus warneri		29.42	0.51	MLT 86	8	43	
Staphylococcus petrasii subsp. petrasii		4.66	0.20		99	1	
702611	**Low discrimination**						**702611**
					PYR		
Staphylococcus petrasii subsp. petrasii		75.17	0.39	MAN 10	99		
Staphylococcus capitis subsp. capitis		12.13	0.24	MLT 99	1		
Staphylococcus capitis subsp. ureolyticus		9.33	0.13		1		
702612	**Low discrimination**						**702612**
					YEP		
Staphylococcus pasteuri		40.29	0.35	GLR 99	86		
Staphylococcus simulans		31.28	0.38	MLT 86	1		
Staphylococcus warneri		15.71	0.41		43		
702613	**Low discrimination**						**702613**
					PYR		
Staphylococcus simulans		51.56	0.48	VPT 4	70		
Staphylococcus capitis subsp. ureolyticus		30.16	0.31	BGA 90	1		
Staphylococcus petrasii subsp. petrasii		17.84	0.34		99		
702640	**Low discrimination**						**702640**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		46.99	0.80	MNS 90	99	1	
Staphylococcus hominis subsp. hominis		16.43	0.80		1	49	
Staphylococcus petrasii subsp. croceilyticus		14.29	0.59		90	99	
702641	**Excellent identification**						**702641**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.77	1.00		99		
Staphylococcus hominis subsp. hominis		0.08	0.45		1		
Staphylococcus warneri		0.04	0.45		8		
702642	**Low discrimination**						**702642**
					PYR		
Staphylococcus petrasii subsp. petrasii		49.71	0.75	MNS 90	99		
Staphylococcus hominis subsp. hominis		37.89	0.82		1		
Staphylococcus warneri		5.72	0.70		8		
702643	**Excellent identification**						**702643**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.43	0.95		99		
Staphylococcus capitis subsp. ureolyticus		0.19	0.31		1		
Staphylococcus hominis subsp. hominis		0.17	0.47		1		
702650	**Low discrimination**						**702650**
					PYR	YEP	
Staphylococcus warneri		32.18	0.77		8	43	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		26.59	0.61		99	1	
Staphylococcus pasteuri		21.77	0.59		1	86	
702651	**Very good identification**						**702651**
					PYR		
Staphylococcus petrasii subsp. petrasii		96.51	0.80	MAN 10	99		
Staphylococcus capitis subsp. ureolyticus		2.91	0.41		1		
Staphylococcus warneri		0.27	0.42		8		
702652	**Low discrimination**						**702652**
					PYR		
Staphylococcus hominis subsp. hominis		27.47	0.65	ARG 12	1		
Staphylococcus petrasii subsp. petrasii		26.79	0.55	MAN 13	99		
Staphylococcus warneri		20.08	0.67		8		
702653	**Low discrimination**						**702653**
					PYR		
Staphylococcus petrasii subsp. petrasii		69.39	0.75	MAN 10	99		
Staphylococcus capitis subsp. ureolyticus		28.45	0.59		1		
Staphylococcus simulans		0.91	0.41		70		
702660	**Low discrimination**						**702660**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		47.23	0.39	XYL 1	99	1	
Staphylococcus hominis subsp. hominis		16.51	0.39	MNS 90	1	49	
Staphylococcus petrasii subsp. croceilyticus		14.36	0.18		90	99	
702661	**Very good identification**						**702661**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.92	0.59	XYL 1	99		
Staphylococcus hominis subsp. hominis		0.08	0.04		1		
702662	**Low discrimination**						**702662**
					PYR		
Staphylococcus petrasii subsp. petrasii		50.29	0.34	XYL 1	99		
Staphylococcus hominis subsp. hominis		38.33	0.40	MNS 90	1		
Staphylococcus warneri		5.78	0.29		8		
702663	**Very good identification**						**702663**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.64	0.53	XYL 1	99		
Staphylococcus capitis subsp. ureolyticus		0.19	-.11		1		
Staphylococcus hominis subsp. hominis		0.17	0.06		1		
702670	**Low discrimination**						**702670**
					PYR	YEP	
Staphylococcus warneri		32.86	0.36	XYL 1	8	43	
Staphylococcus petrasii subsp. petrasii		27.15	0.20		99	1	
Staphylococcus pasteuri		22.23	0.18		1	86	
702671	**Good identification**						**702671**
					PYR		
Staphylococcus petrasii subsp. petrasii		97.08	0.39	MAN 10	99		
Staphylococcus capitis subsp. ureolyticus		2.92	0.00	XYL 1	1		
702673	**Low discrimination**						**702673**
					PYR		
Staphylococcus petrasii subsp. petrasii		69.72	0.34	MAN 10	99		
Staphylococcus capitis subsp. ureolyticus		28.58	0.18	XYL 1	1		
Staphylococcus simulans		0.92	0.00		70		
702700	**Low discrimination**						**702700**
					PYR		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		43.50	0.36	MLT 99	99		
Staphylococcus hominis subsp. hominis		24.71	0.40	MNS 90	1		
Staphylococcus warneri		19.94	0.43		8		
702701	**Very good identification**						**702701**
					YEP	PYR	
Staphylococcus petrasii subsp. petrasii		99.21	0.55	MLT 99	1	99	
Staphylococcus hyicus		0.31	0.02		1	1	
Staphylococcus chromogenes		0.14	-.02		95	45	
702702	**Low discrimination**						**702702**
					PYR		
Staphylococcus hominis subsp. hominis		40.33	0.41	ARG 12	1		
Staphylococcus petrasii subsp. petrasii		32.56	0.30	MLT 99	99		
Staphylococcus piscifermentans		14.95	0.39		15		
702703	**Acceptable identification**						**702703**
					YEP	PYR	
Staphylococcus petrasii subsp. petrasii		90.27	0.50	MLT 99	1	99	
Staphylococcus hyicus		6.01	0.24		1	1	
Staphylococcus chromogenes		2.07	0.18		95	45	
702710	**Low discrimination**						**702710**
					PYR		
Staphylococcus warneri		49.68	0.41	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		16.63	0.16		99		
Staphylococcus hominis subsp. hominis		12.71	0.23		1		
702711	**Good identification**						**702711**
Staphylococcus petrasii subsp. petrasii		100.0	0.36	MAN 10			
702712	**Low discrimination**						**702712**
Staphylococcus warneri		23.59	0.31	MLT 95			
Staphylococcus piscifermentans		22.59	0.32				
Staphylococcus hominis subsp. hominis		21.25	0.24				
702713	**Low discrimination**						**702713**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		39.16	0.30	MAN 10	1	99	
Staphylococcus aureus subsp. aureus		32.47	0.18	MLT 99	99	10	
Staphylococcus simulans		14.77	0.27		1	70	
702740	**Low discrimination**						**702740**
					PYR		
Staphylococcus petrasii subsp. petrasii		57.78	0.77	MNS 90	99		
Staphylococcus hominis subsp. hominis		32.83	0.81		1		
Staphylococcus warneri		5.08	0.70		8		
702741	**Excellent identification**						**702741**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.76	0.96		99		
Staphylococcus hominis subsp. hominis		0.13	0.46		1		
Staphylococcus epidermidis		0.05	0.33		1		
702742	**Low discrimination**						**702742**
					PYR		
Staphylococcus hominis subsp. hominis		52.07	0.82	ARG 12	1		
Staphylococcus petrasii subsp. petrasii		42.04	0.71		99		
Staphylococcus haemolyticus		2.58	0.54		98		
702743	**Very good identification**						**702743**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR		
Staphylococcus petrasii subsp. petrasii		99.07	0.91		99		
Staphylococcus hominis subsp. hominis		0.28	0.47		1		
Staphylococcus epidermidis		0.26	0.43		1		
702750	**Low discrimination**						**702750**
					PYR		
Staphylococcus petrasii subsp. petrasii		35.86	0.57	MAN 10	99		
Staphylococcus hominis subsp. hominis		27.40	0.64	MNS 90	1		
Staphylococcus warneri		20.56	0.67		8		
702751	**Very good identification**						**702751**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		98.30	0.77	MAN 10	1	99	
Staphylococcus caprae		0.49	0.24		1	75	
Staphylococcus aureus subsp. aureus		0.44	0.18		99	10	
702752	**Low discrimination**						**702752**
					PYR		
Staphylococcus hominis subsp. hominis		44.68	0.65	ARG 12	1		
Staphylococcus petrasii subsp. petrasii		26.82	0.52	MAN 13	99		
Staphylococcus warneri		9.52	0.57		8		
702753	**Low discrimination**						**702753**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		49.49	0.71	MAN 10	1	99	
Staphylococcus aureus subsp. aureus		41.02	0.59		99	10	
Staphylococcus caprae		8.70	0.50		1	75	
702760	**Low discrimination**						**702760**
					PYR		
Staphylococcus petrasii subsp. petrasii		58.31	0.36	XYL 1	99		
Staphylococcus hominis subsp. hominis		33.13	0.40	MNS 90	1		
Staphylococcus warneri		5.13	0.29		8		
702761	**Very good identification**						**702761**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.87	0.55	XYL 1	99		
Staphylococcus hominis subsp. hominis		0.13	0.05		1		
702762	**Low discrimination**						**702762**
					PYR		
Staphylococcus hominis subsp. hominis		52.61	0.41	ARG 12	1		
Staphylococcus petrasii subsp. petrasii		42.47	0.30	XYL 1	99		
Staphylococcus haemolyticus		2.61	0.13		98		
702763	**Good identification**						**702763**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.46	0.50	XYL 1	99		
Staphylococcus hominis subsp. hominis		0.28	0.06		1		
Staphylococcus epidermidis		0.26	0.02		1		
702771	**Good identification**						**702771**
Staphylococcus petrasii subsp. petrasii		100.0	0.36	MAN 10			
702773	**Low discrimination**						**702773**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		48.78	0.30	MAN 10	1	99	
Staphylococcus aureus subsp. aureus		40.44	0.18	XYL 1	99	10	
Staphylococcus caprae		8.58	0.09		1	75	
703053	**Good identification**						**703053**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	ESL 1		
703101	**Good identification**					**703101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	PHS 99		
703103	**Very good identification**					**703103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	PHS 99		
703111	**Good identification**					**703111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	PHS 99		
703113	**Good identification**					**703113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	PHS 99		
703213	**Good identification**					**703213**
					COA	
Staphylococcus capitis subsp. ureolyticus		99.27	0.31	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		0.73	0.01	MLT 96	99	
703240	**Low discrimination**					**703240**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		67.25	0.27	ARG 1	99	
Staphylococcus hominis subsp. hominis		22.28	0.29	ESL 8	1	
Staphylococcus epidermidis		5.12	0.12		1	
703243	**Low discrimination**					**703243**
					PYR	
Staphylococcus capitis subsp. ureolyticus		78.94	0.31	ESL 1	1	
Staphylococcus epidermidis		12.93	0.28	MAN 96	1	
Staphylococcus petrasii subsp. petrasii		4.24	0.12		99	
703251	**Good identification**					**703251**
					PYR	
Staphylococcus capitis subsp. ureolyticus		99.66	0.41	ESL 1	1	
Staphylococcus petrasii subsp. petrasii		0.34	-.02	LAC 88	99	
703253	**Very good identification**					**703253**
Staphylococcus capitis subsp. ureolyticus		100.0	0.59	ESL 1		
703301	**Good identification**					**703301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.27	PHS 99		
703303	**Good identification**					**703303**
					COA	
Staphylococcus schleiferi subsp. coagulans		98.83	0.43	PHS 99	99	
Staphylococcus epidermidis		1.17	0.02		1	
703313	**Good identification**					**703313**
					COA	
Staphylococcus schleiferi subsp. coagulans		98.64	0.42	PHS 99	99	
Staphylococcus capitis subsp. ureolyticus		1.36	-.11		1	
703340	**Low discrimination**					**703340**
					NOV	
Staphylococcus hominis subsp. novobiosepticus		78.12	0.37	ARG 1	99	
Staphylococcus epidermidis		11.23	0.28	ESL 8	1	
Staphylococcus hominis subsp. hominis		9.34	0.29		1	
703341	**Low discrimination**					**703341**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR	NOV	
Staphylococcus epidermidis		46.21	0.33	ESL 1	1	1	
Staphylococcus hominis subsp. novobiosepticus		43.29	0.25		1	99	
Staphylococcus petrasii subsp. petrasii		10.50	0.14		99	1	
703342	**Low discrimination**						**703342**
					NOV		
Staphylococcus hominis subsp. novobiosepticus		46.87	0.32	ARG 1	99		
Staphylococcus epidermidis		39.60	0.38	ESL 8	1		
Staphylococcus hominis subsp. hominis		12.22	0.31		1		
703343	**Low discrimination**						**703343**
					PYR	NOV	
Staphylococcus epidermidis		82.14	0.43	ESL 1	1	1	
Staphylococcus hominis subsp. novobiosepticus		13.09	0.19		1	99	
Staphylococcus petrasii subsp. petrasii		3.17	0.09		99	1	
703400	**Good identification**						**703400**
Staphylococcus piscifermentans		100.0	0.48	VPT 1			
703402	**Good identification**						**703402**
Staphylococcus piscifermentans		100.0	0.52	VPT 1			
703410	**Good identification**						**703410**
Staphylococcus piscifermentans		100.0	0.41	VPT 1			
703412	**Good identification**						**703412**
Staphylococcus piscifermentans		100.0	0.44	VPT 1			
703440	**Good identification**						**703440**
					PYR		
Staphylococcus piscifermentans		95.67	0.48	VPT 1	15		
Staphylococcus petrasii subsp. petrasii		2.20	-.02		99		
Staphylococcus warneri		2.13	0.13		8		
703442	**Good identification**						**703442**
Staphylococcus piscifermentans		100.0	0.52	VPT 1			
703450	**Good identification**						**703450**
Staphylococcus piscifermentans		96.38	0.41	VPT 1			
Staphylococcus warneri		3.62	0.10				
703452	**Good identification**						**703452**
Staphylococcus piscifermentans		100.0	0.44	VPT 1			
703500	**Good identification**						**703500**
Staphylococcus piscifermentans		100.0	0.52	VPT 1			
703502	**Very good identification**						**703502**
Staphylococcus piscifermentans		100.0	0.55	VPT 1			
703510	**Good identification**						**703510**
Staphylococcus piscifermentans		100.0	0.44	VPT 1			
703512	**Good identification**						**703512**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus piscifermentans		100.0	0.48	VPT 1			
703540	**Good identification**						**703540**
Staphylococcus piscifermentans		98.99	0.52	VPT 1			
Staphylococcus petrasii subsp. petrasii		1.01	-.05				
703542	**Very good identification**						**703542**
Staphylococcus piscifermentans		100.0	0.55	VPT 1			
703550	**Good identification**						**703550**
Staphylococcus piscifermentans		100.0	0.44	VPT 1			
703552	**Acceptable identification**						**703552**
Staphylococcus piscifermentans		94.80	0.48	VPT 1			
Staphylococcus caprae		5.20	0.00				
703600	**Good identification**						**703600**
					PYR		
Staphylococcus piscifermentans		97.07	0.52	VPT 1	15		
Staphylococcus petrasii subsp. petrasii		1.49	-.02		99		
Staphylococcus warneri		1.44	0.13		8		
703602	**Very good identification**						**703602**
Staphylococcus piscifermentans		100.0	0.55	VPT 1			
703610	**Acceptable identification**						**703610**
					YEP		
Staphylococcus piscifermentans		92.81	0.44	VPT 1	1		
Staphylococcus pasteuri		4.87	0.02		86		
Staphylococcus warneri		2.33	0.10		43		
703612	**Good identification**						**703612**
Staphylococcus piscifermentans		100.0	0.48	VPT 1			
703640	**Low discrimination**						**703640**
					PYR		
Staphylococcus petrasii subsp. petrasii		36.01	0.39	ESL 1	99		
Staphylococcus piscifermentans		23.74	0.52	MNS 90	15		
Staphylococcus hominis subsp. hominis		12.59	0.39		1		
703641	**Very good identification**						**703641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		99.64	0.59	ESL 1	1	99	
Kocuria kristinae		0.21	0.04		99	93	
Staphylococcus hominis subsp. hominis		0.08	0.04		1	1	
703642	**Low discrimination**						**703642**
					PYR		
Staphylococcus piscifermentans		48.02	0.55	VPT 1	15		
Staphylococcus petrasii subsp. petrasii		26.14	0.34		99		
Staphylococcus hominis subsp. hominis		19.93	0.40		1		
703643	**Very good identification**						**703643**
					PYR		
Staphylococcus petrasii subsp. petrasii		99.43	0.53	ESL 1	99		
Staphylococcus piscifermentans		0.21	0.14		15		
Staphylococcus capitis subsp. ureolyticus		0.19	-.11		1		
703650	**Low discrimination**						**703650**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus piscifermentans		40.84	0.44	VPT 1	15	
Staphylococcus warneri		19.44	0.36		8	
Staphylococcus petrasii subsp. petrasii		16.06	0.20		99	
703651	**Good identification**					**703651**
					PYR	
Staphylococcus petrasii subsp. petrasii		97.08	0.39	ESL 1	99	
Staphylococcus capitis subsp. ureolyticus		2.92	0.00	MAN 10	1	
703652	**Low discrimination**					**703652**
					PYR	
Staphylococcus piscifermentans		66.38	0.48	VPT 1	15	
Staphylococcus hominis subsp. hominis		9.61	0.23		1	
Staphylococcus petrasii subsp. petrasii		9.37	0.14		99	
703653	**Low discrimination**					**703653**
					PYR	
Staphylococcus petrasii subsp. petrasii		69.72	0.34	ESL 1	99	
Staphylococcus capitis subsp. ureolyticus		28.58	0.18	MAN 10	1	
Staphylococcus simulans		0.92	0.00		70	
703700	**Very good identification**					**703700**
Staphylococcus piscifermentans		97.34	0.55	VPT 1		
Rothia mucilaginosa		2.00	0.00			
Staphylococcus petrasii subsp. petrasii		0.66	-.05			
703702	**Very good identification**					**703702**
Staphylococcus piscifermentans		100.0	0.59	VPT 1		
703710	**Good identification**					**703710**
Staphylococcus piscifermentans		100.0	0.48	VPT 1		
703712	**Good identification**					**703712**
Staphylococcus piscifermentans		100.0	0.51	VPT 1		
703740	**Low discrimination**					**703740**
					PYR	
Staphylococcus piscifermentans		45.16	0.55	VPT 1	15	
Staphylococcus petrasii subsp. petrasii		30.45	0.36		99	
Staphylococcus hominis subsp. hominis		17.30	0.40		1	
703741	**Very good identification**					**703741**
Staphylococcus petrasii subsp. petrasii		97.31	0.55	ESL 1		
Rothia mucilaginosa		2.30	0.18			
Staphylococcus piscifermentans		0.16	0.14			
703742	**Low discrimination**					**703742**
					PYR	
Staphylococcus piscifermentans		63.70	0.59	VPT 1	15	
Staphylococcus hominis subsp. hominis		19.10	0.41		1	
Staphylococcus petrasii subsp. petrasii		15.42	0.30		99	
703743	**Good identification**					**703743**
					PYR	
Staphylococcus petrasii subsp. petrasii		99.00	0.50	ESL 1	99	
Staphylococcus piscifermentans		0.46	0.18		15	
Staphylococcus hominis subsp. hominis		0.28	0.06		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
703750	**Low discrimination**						**703750**
					PYR		
Staphylococcus piscifermentans		67.43	0.48	VPT 1	15		
Staphylococcus petrasii subsp. petrasii		11.79	0.16		99		
Staphylococcus hominis subsp. hominis		9.01	0.23		1		
703751	**Good identification**						**703751**
Staphylococcus petrasii subsp. petrasii		100.0	0.36	ESL 1			
703752	**Low discrimination**						**703752**
					PYR		
Staphylococcus piscifermentans		81.74	0.51	VPT 1	15		
Staphylococcus hominis subsp. hominis		8.54	0.24		1		
Staphylococcus petrasii subsp. petrasii		5.13	0.11		99		
703753	**Low discrimination**						**703753**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		49.44	0.30	ESL 1	1	99	
Staphylococcus aureus subsp. aureus		40.99	0.18	MAN 10	99	10	
Staphylococcus caprae		8.69	0.09		1	75	
704101	**Low discrimination**						**704101**
					COA		
Staphylococcus schleiferi subsp. coagulans		69.67	0.37	NIT 99	99		
Staphylococcus schleiferi subsp. schleiferi		30.33	0.26	LAC 86	1		
704103	**Very good identification**						**704103**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.54	0.53	NIT 99	99		
Staphylococcus caprae		0.46	-.07		1		
704111	**Good identification**						**704111**
					COA		
Staphylococcus schleiferi subsp. coagulans		97.02	0.36	NIT 99	99		
Staphylococcus caprae		2.98	-.07	LAC 86	1		
704113	**Acceptable identification**						**704113**
					COA		
Staphylococcus schleiferi subsp. coagulans		91.32	0.53	NIT 99	99		
Staphylococcus caprae		8.68	0.20		1		
704143	**Low discrimination**						**704143**
					COA		
Staphylococcus caprae		86.33	0.34	NIT 90	1		
Staphylococcus epidermidis		11.77	0.28	TRE 90	1		
Staphylococcus schleiferi subsp. coagulans		1.91	0.12	MAN 95	99		
704151	**Good identification**						**704151**
Staphylococcus caprae		100.0	0.34	NIT 90			
704152	**Good identification**						**704152**
Staphylococcus caprae		100.0	0.43	NIT 90			
704153	**Very good identification**						**704153**
					COA		
Staphylococcus caprae		99.89	0.61	NIT 90	1		
Staphylococcus schleiferi subsp. coagulans		0.11	0.12	TRE 90	99		
704240	**Low discrimination**						**704240**
					COA		
Staphylococcus epidermidis		65.49	0.38	NIT 97	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus aureus subsp. anaerobius		31.12	0.18		99	
Staphylococcus warneri		3.40	0.20		1	
704241	**Good identification**					**704241**
					PYR	
Staphylococcus epidermidis		98.00	0.43	NIT 97	1	
Staphylococcus petrasii subsp. petrasii		1.21	-.02		99	
Staphylococcus capitis subsp. ureolyticus		0.78	-.14		1	
704242	**Good identification**					**704242**
Staphylococcus epidermidis		99.46	0.48	NIT 97		
Staphylococcus warneri		0.54	0.10			
704243	**Very good identification**					**704243**
Staphylococcus epidermidis		98.18	0.53	NIT 97		
Staphylococcus capitis subsp. ureolyticus		1.82	0.04			
704253	**Low discrimination**					**704253**
					COA	
Staphylococcus capitis subsp. ureolyticus		79.35	0.33	NIT 95	1	
Staphylococcus delphini		13.11	0.18	PHS 1	99	
Staphylococcus caprae		5.32	0.11		1	
704301	**Low discrimination**					**704301**
					COA	
Staphylococcus schleiferi subsp. coagulans		72.32	0.27	NIT 99	99	
Staphylococcus epidermidis		27.68	0.18	LAC 86	1	
704303	**Low discrimination**					**704303**
					COA YEP	
Staphylococcus schleiferi subsp. coagulans		82.93	0.43	NIT 99	99 1	
Staphylococcus epidermidis		16.36	0.28		1 1	
Staphylococcus chromogenes		0.71	-.04		1 95	
704313	**Low discrimination**					**704313**
					COA	
Staphylococcus schleiferi subsp. coagulans		89.98	0.42	NIT 99	99	
Staphylococcus caprae		10.02	0.11		1	
704340	**Very good identification**					**704340**
					NOV	
Staphylococcus epidermidis		99.20	0.53	NIT 97	1	
Staphylococcus hominis subsp. novobiosepticus		0.51	-.02		99	
Staphylococcus warneri		0.29	0.10		1	
704341	**Very good identification**					**704341**
					PYR	
Staphylococcus epidermidis		99.40	0.59	NIT 97	1	
Staphylococcus caprae		0.45	-.01		75	
Staphylococcus petrasii subsp. petrasii		0.15	-.05		99	
704342	**Very good identification**					**704342**
					NOV	
Staphylococcus epidermidis		99.22	0.63	NIT 97	1	
Staphylococcus caprae		0.69	0.08		1	
Staphylococcus hominis subsp. novobiosepticus		0.09	-.07		99	
704343	**Very good identification**					**704343**
					COA	
Staphylococcus epidermidis		97.24	0.69	NIT 97	1	
Staphylococcus caprae		2.67	0.26		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. coagulans		0.05	0.02		99		
704352	**Acceptable identification**						**704352**
Staphylococcus caprae		92.89	0.34	NIT 90			
Staphylococcus epidermidis		7.11	0.22	TRE 90			
704353	**Acceptable identification**						**704353**
					COA		
Staphylococcus caprae		92.11	0.52	NIT 90	1		
Staphylococcus aureus subsp. aureus		3.59	0.18	TRE 90	99		
Staphylococcus delphini		2.29	0.18		99		
704363	**Good identification**						**704363**
Staphylococcus epidermidis		100.0	0.28	NIT 97			
704401	**Good identification**						**704401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	URE 1			
704450	**Low discrimination**						**704450**
					NOV	YEP	
Staphylococcus kloosii		39.93	0.27	VPT 10	99	99	
Staphylococcus warneri		33.08	0.17	ARG 1	1	43	
Staphylococcus caprae		26.98	-.05		1	1	
704453	**Good identification**						**704453**
					COA	NOV	
Staphylococcus caprae		99.11	0.39	NIT 90	1	1	
Staphylococcus cohnii subsp. urealyticum		0.57	0.10	GAL 99	1	99	
Staphylococcus aureus subsp. aureus		0.32	-.17		99	4	
704501	**Good identification**						**704501**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.36	URE 1			
704513	**Good identification**						**704513**
					COA		
Staphylococcus caprae		98.21	0.39	NIT 90	1		
Staphylococcus schleiferi subsp. coagulans		1.16	0.12	MLT 99	99		
Staphylococcus felis		0.63	-.04		1		
704541	**Good identification**						**704541**
Staphylococcus caprae		98.71	0.28	NIT 90			
Staphylococcus petrasii subsp. petrasii		1.29	-.05	MAN 95			
704542	**Good identification**						**704542**
Staphylococcus caprae		100.0	0.36	NIT 90			
704543	**Very good identification**						**704543**
Staphylococcus caprae		100.0	0.54	NIT 90			
704550	**Good identification**						**704550**
					NOV	YEP	
Staphylococcus caprae		97.18	0.36	NIT 90	1	1	
Staphylococcus kloosii		2.82	0.33	MNS 88	99	99	
704551	**Very good identification**						**704551**
Staphylococcus caprae		100.0	0.54	NIT 90			
704552	**Very good identification**						**704552**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV	YEP	
Staphylococcus caprae		99.84	0.63	NIT 90	1	1	
Staphylococcus kloosii		0.17	0.33	MNS 88	99	99	
704553	**Very good identification**						**704553**
					COA		
Staphylococcus caprae		99.84	0.80	NIT 90	1		
Staphylococcus aureus subsp. aureus		0.16	0.18		99		
704573	**Good identification**						**704573**
Staphylococcus caprae		100.0	0.39	NIT 90			
704640	**Low discrimination**						**704640**
					PYR		
Staphylococcus warneri		56.84	0.47	PHS 1	8		
Staphylococcus petrasii subsp. petrasii		14.58	0.20		99		
Staphylococcus petrasii subsp. jettensis		12.84	0.10		99		
704641	**Good identification**						**704641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		97.33	0.39	NIT 90	1	99	
Kocuria kristinae		1.00	-.02	PHS 1	99	93	
Staphylococcus warneri		0.86	0.12		1	8	
704642	**Low discrimination**						**704642**
					PYR		
Staphylococcus warneri		55.63	0.37	PHS 1	8		
Staphylococcus petrasii subsp. petrasii		23.04	0.14		99		
Staphylococcus petrasii subsp. jettensis		6.65	-.05		99		
704643	**Low discrimination**						**704643**
					PYR		
Staphylococcus petrasii subsp. petrasii		87.50	0.34	NIT 90	99		
Staphylococcus caprae		5.94	0.04	PHS 1	75		
Staphylococcus epidermidis		4.25	0.12		1		
704650	**Low discrimination**						**704650**
					NOV		
Staphylococcus warneri		76.28	0.44	PHS 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		11.69	0.06		99		
Staphylococcus pasteuri		9.03	0.10		1		
704653	**Low discrimination**						**704653**
					COA	PYR	
Staphylococcus caprae		50.43	0.30	NIT 90	1	75	
Staphylococcus aureus subsp. aureus		43.68	0.24	GAL 99	99	10	
Staphylococcus petrasii subsp. petrasii		4.35	0.14		1	99	
704713	**Low discrimination**						**704713**
					COA		
Staphylococcus caprae		66.49	0.30	NIT 90	1		
Staphylococcus aureus subsp. aureus		28.50	0.18	MLT 99	99		
Staphylococcus felis		2.35	0.02		1		
704740	**Low discrimination**						**704740**
					PYR		
Staphylococcus warneri		49.99	0.36	PHS 1	8		
Staphylococcus petrasii subsp. petrasii		27.06	0.16		99		
Staphylococcus caprae		9.48	0.01		75		
704741	**Low discrimination**						**704741**
					PYR		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		73.10	0.36	NIT 90	99		
Staphylococcus caprae		20.88	0.19	PHS 1	75		
Staphylococcus epidermidis		5.13	0.18		1		
704742	**Low discrimination**						**704742**
Staphylococcus caprae		73.37	0.27	NIT 90			
Staphylococcus epidermidis		11.86	0.22	MAN 95			
Staphylococcus warneri		6.79	0.26	MNS 88			
704743	**Low discrimination**						**704743**
					COA	PYR	
Staphylococcus caprae		80.46	0.45	NIT 90	1	75	
Staphylococcus petrasii subsp. petrasii		7.99	0.30	MAN 95	1	99	
Staphylococcus aureus subsp. aureus		6.62	0.18		99	10	
704750	**Low discrimination**						**704750**
					PYR		
Staphylococcus caprae		82.13	0.27	NIT 90	75		
Staphylococcus warneri		16.50	0.33	MNS 88	8		
Staphylococcus petrasii subsp. petrasii		1.37	-.04	LAC 95	99		
704751	**Acceptable identification**						**704751**
					COA	PYR	
Staphylococcus caprae		90.68	0.45	NIT 90	1	75	
Staphylococcus aureus subsp. aureus		7.46	0.18	LAC 95	99	10	
Staphylococcus petrasii subsp. petrasii		1.86	0.16		1	99	
704752	**Very good identification**						**704752**
					COA		
Staphylococcus caprae		96.36	0.54	NIT 90	1		
Staphylococcus aureus subsp. aureus		3.06	0.18	MNS 88	99		
Staphylococcus warneri		0.34	0.24		1		
704753	**Low discrimination**						**704753**
					COA	PYR	
Staphylococcus caprae		69.95	0.72	NIT 90	1	75	
Staphylococcus aureus subsp. aureus		29.99	0.59		99	10	
Staphylococcus petrasii subsp. petrasii		0.04	0.11		1	99	
704773	**Low discrimination**						**704773**
					COA		
Staphylococcus caprae		70.00	0.30	NIT 90	1		
Staphylococcus aureus subsp. aureus		30.00	0.18	XYL 1	99		
705101	**Good identification**						**705101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	NIT 99			
705103	**Very good identification**						**705103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	NIT 99			
705111	**Good identification**						**705111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	NIT 99			
705113	**Good identification**						**705113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	NIT 99			
705301	**Good identification**						**705301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.27	NIT 99			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
705303	**Good identification**						**705303**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.43	NIT 99			
705313	**Good identification**						**705313**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.42	NIT 99			
705343	**Good identification**						**705343**
					COA		
	Staphylococcus epidermidis	95.31	0.28	ESL 1	1		
	Staphylococcus schleiferi subsp. coagulans	4.69	0.02	NIT 97	99		
705450	**Good identification**						**705450**
	Staphylococcus kloosii	100.0	0.27	VPT 10			
705452	**Good identification**						**705452**
	Staphylococcus kloosii	100.0	0.28	VPT 10			
705550	**Low discrimination**						**705550**
					NOV	YEP	
	Staphylococcus kloosii	74.18	0.33	VPT 10	99	99	
	Staphylococcus caprae	25.82	-.05	ARG 1	1	1	
705553	**Good identification**						**705553**
	Staphylococcus caprae	100.0	0.39	ESL 1			
705753	**Low discrimination**						**705753**
					COA		
	Staphylococcus caprae	70.00	0.30	ESL 1	1		
	Staphylococcus aureus subsp. aureus	30.00	0.18	NIT 90	99		
706001	**Low discrimination**						**706001**
					COA		
	Staphylococcus schleiferi subsp. schleiferi	76.15	0.46	URE 1	1		
	Staphylococcus schleiferi subsp. coagulans	19.43	0.37		99		
	Staphylococcus carnosus subsp. carnosus	2.43	0.15		1		
706003	**Good identification**						**706003**
					COA		
	Staphylococcus schleiferi subsp. coagulans	91.63	0.53	GAL 99	99		
	Staphylococcus carnosus subsp. carnosus	7.46	0.27		1		
	Staphylococcus schleiferi subsp. schleiferi	0.59	0.04		1		
706011	**Low discrimination**						**706011**
					COA		
	Staphylococcus capitis subsp. capitis	48.19	0.35	URE 1	1		
	Staphylococcus schleiferi subsp. coagulans	32.70	0.36	PHS 11	99		
	Staphylococcus carnosus subsp. carnosus	17.71	0.27		1		
706013	**Low discrimination**						**706013**
					COA		
	Staphylococcus schleiferi subsp. coagulans	72.28	0.53	GAL 99	99		
	Staphylococcus carnosus subsp. carnosus	25.49	0.39		1		
	Staphylococcus capitis subsp. ureolyticus	1.25	0.02		1		
706040	**Good identification**						**706040**
	Staphylococcus epidermidis	100.0	0.28	SUC 99			
706041	**Acceptable identification**						**706041**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		94.57	0.33	SUC 99		
Staphylococcus schleiferi subsp. schleiferi		5.43	0.04			
706042	**Good identification**					**706042**
Staphylococcus epidermidis		97.12	0.38	SUC 99		
Staphylococcus caprae		2.88	-.05			
706043	**Low discrimination**					**706043**
Staphylococcus epidermidis		84.02	0.43	SUC 99		
Staphylococcus caprae		9.82	0.13			
Staphylococcus capitis subsp. ureolyticus		3.77	0.02			
706053	**Low discrimination**					**706053**
					COA	
Staphylococcus caprae		62.11	0.39	GAL 99	1	
Staphylococcus capitis subsp. ureolyticus		30.13	0.31	TRE 90	1	
Staphylococcus delphini		6.29	0.18		99	
706100	**Acceptable identification**					**706100**
					COA	
Staphylococcus schleiferi subsp. coagulans		92.85	0.37	MNS 99	99	
Staphylococcus schleiferi subsp. schleiferi		3.68	0.04	LAC 86	1	
Staphylococcus epidermidis		1.97	0.02		1	
706101	**Very good identification**					**706101**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.13	0.78	LAC 86	99	
Staphylococcus schleiferi subsp. schleiferi		3.81	0.46		1	
Staphylococcus epidermidis		0.04	0.08		1	
706102	**Very good identification**					**706102**
					COA	
Staphylococcus schleiferi subsp. coagulans		98.37	0.53	MNS 99	99	
Staphylococcus epidermidis		1.08	0.12		1	
Staphylococcus caprae		0.56	-.05		1	
706103	**Excellent identification**					**706103**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.91	0.95		99	
Staphylococcus caprae		0.04	0.13		1	
Staphylococcus epidermidis		0.02	0.18		1	
706110	**Good identification**					**706110**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.37	0.36	MNS 99	99	
Staphylococcus caprae		3.64	-.05	LAC 86	1	
706111	**Very good identification**					**706111**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.54	0.78	LAC 86	99	
Staphylococcus caprae		0.28	0.13		1	
Staphylococcus carnosus subsp. carnosus		0.14	0.15		1	
706112	**Low discrimination**					**706112**
					COA	
Staphylococcus schleiferi subsp. coagulans		89.55	0.53	MNS 99	99	
Staphylococcus caprae		10.45	0.22		1	
706113	**Excellent identification**					**706113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.05	0.94		99	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus caprae		0.86	0.39		1	
Staphylococcus carnosus subsp. carnosus		0.09	0.27		1	
706121	**Good identification**					**706121**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.19	0.37	XYL 1	99	
Staphylococcus schleiferi subsp. schleiferi		3.81	0.04	LAC 86	1	
706123	**Very good identification**					**706123**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	XYL 1		
706131	**Good identification**					**706131**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	XYL 1		
706133	**Good identification**					**706133**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.14	0.53	XYL 1	99	
Staphylococcus caprae		0.86	-.02		1	
706140	**Acceptable identification**					**706140**
Staphylococcus epidermidis		92.06	0.43	SUC 99		
Staphylococcus caprae		7.94	0.10			
706141	**Low discrimination**					**706141**
					COA	
Staphylococcus epidermidis		62.25	0.49	SUC 99	1	
Staphylococcus caprae		21.19	0.28		1	
Staphylococcus schleiferi subsp. coagulans		15.93	0.37		99	
706142	**Low discrimination**					**706142**
					COA	
Staphylococcus epidermidis		65.51	0.53	SUC 99	1	
Staphylococcus caprae		33.88	0.36		1	
Staphylococcus schleiferi subsp. coagulans		0.61	0.12		99	
706143	**Low discrimination**					**706143**
					COA	
Staphylococcus caprae		57.70	0.54	TRE 90	1	
Staphylococcus epidermidis		28.25	0.59	MAN 95	1	
Staphylococcus schleiferi subsp. coagulans		14.02	0.53		99	
706150	**Good identification**					**706150**
Staphylococcus caprae		99.39	0.36	TRE 90		
Staphylococcus epidermidis		0.61	0.02	MNS 88		
706151	**Very good identification**					**706151**
					COA	
Staphylococcus caprae		96.33	0.54	TRE 90	1	
Staphylococcus schleiferi subsp. coagulans		3.52	0.36	LAC 95	99	
Staphylococcus epidermidis		0.15	0.08		1	
706152	**Very good identification**					**706152**
					COA	
Staphylococcus caprae		99.81	0.63	TRE 90	1	
Staphylococcus epidermidis		0.10	0.12	MNS 88	1	
Staphylococcus schleiferi subsp. coagulans		0.09	0.12		99	
706153	**Very good identification**					**706153**
					COA	
Staphylococcus caprae		98.55	0.80	TRE 90	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. coagulans		1.16	0.53		99		
Staphylococcus aureus subsp. aureus		0.16	0.18		99		
706173	**Good identification**						**706173**
					COA		
Staphylococcus caprae		98.83	0.39	TRE 90	1		
Staphylococcus schleiferi subsp. coagulans		1.17	0.12	XYL 1	99		
706200	**Good identification**						**706200**
Staphylococcus epidermidis		100.0	0.28	MLT 99			
706201	**Low discrimination**						**706201**
					COA		
Staphylococcus epidermidis		54.93	0.33	MLT 99	1		
Staphylococcus schleiferi subsp. coagulans		25.15	0.27		99		
Staphylococcus capitis subsp. capitis		15.15	0.17		1		
706202	**Good identification**						**706202**
					COA		
Staphylococcus epidermidis		98.36	0.38	MLT 99	1		
Staphylococcus schleiferi subsp. coagulans		1.64	0.02		99		
706203	**Low discrimination**						**706203**
					COA	YEP	
Staphylococcus epidermidis		48.85	0.43	MLT 99	1	1	
Staphylococcus schleiferi subsp. coagulans		43.39	0.43		99	1	
Staphylococcus chromogenes		4.09	0.17		1	95	
706211	**Low discrimination**						**706211**
					COA		
Staphylococcus capitis subsp. capitis		80.49	0.40	URE 1	1		
Staphylococcus capitis subsp. ureolyticus		10.22	0.13	PHS 11	1		
Staphylococcus schleiferi subsp. coagulans		9.29	0.26		99		
706213	**Low discrimination**						**706213**
					COA		
Staphylococcus capitis subsp. ureolyticus		47.99	0.31	PHS 1	1		
Staphylococcus schleiferi subsp. coagulans		36.51	0.42	MLT 96	99		
Staphylococcus delphini		10.02	0.18		99		
706240	**Very good identification**						**706240**
					NOV		
Staphylococcus epidermidis		97.29	0.69		1		
Staphylococcus hominis subsp. hominis		2.31	0.39		1		
Staphylococcus hominis subsp. novobiosepticus		0.27	0.08		99		
706241	**Very good identification**						**706241**
					PYR		
Staphylococcus epidermidis		99.10	0.74		1		
Staphylococcus capitis subsp. ureolyticus		0.47	0.13		1		
Staphylococcus petrasii subsp. petrasii		0.34	0.18		99		
706242	**Very good identification**						**706242**
					NOV		
Staphylococcus epidermidis		99.06	0.79		1		
Staphylococcus hominis subsp. hominis		0.87	0.40		1		
Staphylococcus hominis subsp. novobiosepticus		0.05	0.03		99		
706243	**Very good identification**						**706243**
					COA		
Staphylococcus epidermidis		98.50	0.85		1		
Staphylococcus capitis subsp. ureolyticus		1.07	0.31		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus delphini		0.22	0.18		99		
706250	**Low discrimination**						**706250**
Staphylococcus epidermidis		62.10	0.28	MAN 1			
Staphylococcus hominis subsp. hominis		21.83	0.22				
Staphylococcus capitis subsp. ureolyticus		13.01	0.00				
706251	**Low discrimination**						**706251**
					COA		
Staphylococcus capitis subsp. ureolyticus		85.16	0.41	PHS 1	1		
Staphylococcus epidermidis		7.63	0.33	LAC 88	1		
Staphylococcus delphini		5.43	0.18		99		
706252	**Low discrimination**						**706252**
					COA		
Staphylococcus epidermidis		44.35	0.38	MAN 1	1		
Staphylococcus capitis subsp. ureolyticus		21.52	0.18		1		
Staphylococcus delphini		18.53	0.18		99		
706253	**Low discrimination**						**706253**
					COA		
Staphylococcus capitis subsp. ureolyticus		51.00	0.59	PHS 1	1		
Staphylococcus delphini		43.91	0.59		99		
Staphylococcus epidermidis		1.97	0.43		1		
706260	**Good identification**						**706260**
Staphylococcus epidermidis		100.0	0.28	XYL 1			
706261	**Good identification**						**706261**
Staphylococcus epidermidis		100.0	0.33	XYL 1			
706262	**Good identification**						**706262**
Staphylococcus epidermidis		100.0	0.38	XYL 1			
706263	**Good identification**						**706263**
Staphylococcus epidermidis		98.92	0.43	XYL 1			
Staphylococcus capitis subsp. ureolyticus		1.08	-.11				
706300	**Low discrimination**						**706300**
					COA		
Staphylococcus epidermidis		86.95	0.43	MLT 99	1		
Staphylococcus schleiferi subsp. coagulans		13.05	0.27		99		
706301	**Low discrimination**						**706301**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		88.21	0.68	LAC 86	99	1	
Staphylococcus epidermidis		11.03	0.49		1	1	
Staphylococcus chromogenes		0.52	0.17		1	95	
706302	**Low discrimination**						**706302**
					COA	YEP	
Staphylococcus epidermidis		77.31	0.53	MLT 99	1	1	
Staphylococcus schleiferi subsp. coagulans		22.50	0.43		99	1	
Staphylococcus chromogenes		0.19	-.04		1	95	
706303	**Good identification**						**706303**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		92.90	0.84		99	1	
Staphylococcus epidermidis		5.99	0.59		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus chromogenes		0.80	0.37		1	95	
706311	**Very good identification**						**706311**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		99.27	0.67	LAC 86	99	1	
Staphylococcus caprae		0.33	0.04		1	1	
Staphylococcus chromogenes		0.18	0.06		1	95	
706312	**Low discrimination**						**706312**
					COA		
Staphylococcus schleiferi subsp. coagulans		85.16	0.42	MNS 99	99		
Staphylococcus caprae		11.64	0.13		1		
Staphylococcus epidermidis		3.20	0.12		1		
706313	**Very good identification**						**706313**
					COA		
Staphylococcus schleiferi subsp. coagulans		97.94	0.83		99		
Staphylococcus caprae		0.99	0.30		1		
Staphylococcus aureus subsp. aureus		0.43	0.18		99		
706321	**Low discrimination**						**706321**
					COA		
Staphylococcus schleiferi subsp. coagulans		88.89	0.27	XYL 1	99		
Staphylococcus epidermidis		11.11	0.08	LAC 86	1		
706323	**Acceptable identification**						**706323**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		93.20	0.43	XYL 1	99	1	
Staphylococcus epidermidis		6.01	0.18		1	1	
Staphylococcus chromogenes		0.80	-.04		1	95	
706333	**Good identification**						**706333**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.00	0.42	XYL 1	99		
Staphylococcus caprae		1.00	-.11		1		
706340	**Very good identification**						**706340**
					NOV		
Staphylococcus epidermidis		99.37	0.84		1		
Staphylococcus hominis subsp. hominis		0.45	0.39		1		
Staphylococcus hominis subsp. novobiosepticus		0.14	0.18		99		
706341	**Very good identification**						**706341**
					COA		
Staphylococcus epidermidis		99.68	0.90		1		
Staphylococcus caprae		0.13	0.19		1		
Staphylococcus schleiferi subsp. coagulans		0.08	0.27		99		
706342	**Excellent identification**						**706342**
Staphylococcus epidermidis		99.61	0.95				
Staphylococcus caprae		0.19	0.27				
Staphylococcus hominis subsp. hominis		0.17	0.41				
706343	**Very good identification**						**706343**
					COA		
Staphylococcus epidermidis		98.84	1.00		1		
Staphylococcus caprae		0.75	0.45		1		
Staphylococcus schleiferi subsp. coagulans		0.16	0.43		99		
706350	**Low discrimination**						**706350**
Staphylococcus epidermidis		59.77	0.43	MAN 1			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus caprae		36.21	0.27				
Staphylococcus hominis subsp. hominis		4.02	0.22				
706351	**Low discrimination**						**706351**
					COA		
Staphylococcus caprae		62.51	0.45	TRE 90	1		
Staphylococcus epidermidis		26.13	0.49	LAC 95	1		
Staphylococcus aureus subsp. aureus		5.14	0.18		99		
706352	**Low discrimination**						**706352**
					COA		
Staphylococcus caprae		74.92	0.54	TRE 90	1		
Staphylococcus epidermidis		20.61	0.53	MNS 88	1		
Staphylococcus aureus subsp. aureus		2.38	0.18		99		
706353	**Low discrimination**						**706353**
					COA		
Staphylococcus caprae		55.93	0.72	TRE 90	1		
Staphylococcus aureus subsp. aureus		23.97	0.59		99		
Staphylococcus delphini		15.31	0.59		99		
706360	**Good identification**						**706360**
Staphylococcus epidermidis		100.0	0.43	XYL 1			
706361	**Good identification**						**706361**
Staphylococcus epidermidis		100.0	0.49	XYL 1			
706362	**Very good identification**						**706362**
Staphylococcus epidermidis		100.0	0.53	XYL 1			
706363	**Very good identification**						**706363**
					COA		
Staphylococcus epidermidis		98.96	0.59	XYL 1	1		
Staphylococcus caprae		0.76	0.04		1		
Staphylococcus schleiferi subsp. coagulans		0.16	0.02		99		
706373	**Low discrimination**						**706373**
					COA		
Staphylococcus caprae		56.11	0.30	TRE 90	1		
Staphylococcus aureus subsp. aureus		24.05	0.18	XYL 1	99		
Staphylococcus delphini		15.36	0.18		99		
706401	**Very good identification**						**706401**
Staphylococcus schleiferi subsp. schleiferi		99.55	0.55	URE 1			
Staphylococcus carnosus subsp. carnosus		0.45	0.07				
706402	**Good identification**						**706402**
Staphylococcus piscifermentans		100.0	0.28	VPT 1			
706443	**Acceptable identification**						**706443**
					PYR		
Staphylococcus caprae		92.53	0.33	GAL 99	75		
Staphylococcus petrasii subsp. petrasii		5.09	0.12	MAN 95	99		
Staphylococcus epidermidis		0.89	0.02		1		
706451	**Good identification**						**706451**
					COA	PYR	
Staphylococcus caprae		98.21	0.33	GAL 99	1	75	
Staphylococcus petrasii subsp. petrasii		1.12	-.02	LAC 95	1	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. aureus		0.67	-.17		99	10	
706452	**Good identification**						**706452**
					COA		
Staphylococcus caprae		98.84	0.41	GAL 99	1		
Staphylococcus piscifermentans		0.90	0.21	MNS 88	1		
Staphylococcus aureus subsp. aureus		0.26	-.17		99		
706453	**Very good identification**						**706453**
					COA		
Staphylococcus caprae		96.46	0.59	GAL 99	1		
Staphylococcus aureus subsp. aureus		3.43	0.24		99		
Staphylococcus simulans		0.06	0.03		1		
706500	**Low discrimination**						**706500**
Staphylococcus piscifermentans		48.28	0.28	VPT 1			
Staphylococcus schleiferi subsp. schleiferi		45.42	0.14	ESL 90			
Staphylococcus caprae		6.31	-.12				
706501	**Good identification**						**706501**
					COA		
Staphylococcus schleiferi subsp. schleiferi		91.14	0.55	URE 1	1		
Staphylococcus schleiferi subsp. coagulans		7.75	0.37		99		
Staphylococcus caprae		0.94	0.06		1		
706503	**Low discrimination**						**706503**
					COA		
Staphylococcus schleiferi subsp. coagulans		65.84	0.53	TRE 1	99		
Staphylococcus caprae		24.63	0.33		1		
Staphylococcus felis		4.95	0.20		1		
706511	**Low discrimination**						**706511**
					COA		
Staphylococcus caprae		67.10	0.33	MLT 99	1		
Staphylococcus schleiferi subsp. coagulans		26.95	0.36	LAC 95	99		
Staphylococcus schleiferi subsp. schleiferi		3.47	0.14		1		
706512	**Good identification**						**706512**
					COA		
Staphylococcus caprae		97.23	0.41	MLT 99	1		
Staphylococcus piscifermentans		1.33	0.25	MNS 88	1		
Staphylococcus schleiferi subsp. coagulans		0.94	0.12		99		
706513	**Low discrimination**						**706513**
					COA		
Staphylococcus caprae		81.08	0.59	MLT 99	1		
Staphylococcus schleiferi subsp. coagulans		10.53	0.53		99		
Staphylococcus felis		5.74	0.37		1		
706540	**Low discrimination**						**706540**
Staphylococcus caprae		89.32	0.30	MAN 95			
Staphylococcus piscifermentans		6.91	0.28	MNS 88			
Staphylococcus hominis subsp. hominis		1.66	0.09	LAC 95			
706541	**Good identification**						**706541**
Staphylococcus caprae		97.27	0.47	MAN 95			
Staphylococcus petrasii subsp. petrasii		1.27	0.14	LAC 95			
Staphylococcus schleiferi subsp. schleiferi		0.97	0.14				
706542	**Very good identification**						**706542**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus caprae		99.07	0.56	MAN 95			
Staphylococcus piscifermentans		0.61	0.32	MNS 88			
Staphylococcus epidermidis		0.22	0.12				
706543	**Very good identification**						**706543**
					COA	YEP	
Staphylococcus caprae		99.46	0.74	MAN 95	1	1	
Staphylococcus aureus subsp. aureus		0.34	0.18		99	84	
Staphylococcus chromogenes		0.08	0.18		1	95	
706550	**Very good identification**						**706550**
					NOV		
Staphylococcus caprae		99.39	0.56	MNS 88	1		
Staphylococcus saprophyticus subsp. bovis		0.44	0.18	LAC 95	99		
Staphylococcus piscifermentans		0.17	0.21		1		
706551	**Very good identification**						**706551**
					COA	NOV	
Staphylococcus caprae		99.64	0.74	LAC 95	1	1	
Staphylococcus aureus subsp. aureus		0.34	0.18		99	4	
Staphylococcus saprophyticus subsp. bovis		0.02	0.06		1	99	
706552	**Very good identification**						**706552**
					COA		
Staphylococcus caprae		99.85	0.82	MNS 88	1		
Staphylococcus aureus subsp. aureus		0.13	0.18		99		
Staphylococcus piscifermentans		0.01	0.25		1		
706553	**Very good identification**						**706553**
					COA		
Staphylococcus caprae		98.27	1.00		1		
Staphylococcus aureus subsp. aureus		1.73	0.59		99		
Staphylococcus schleiferi subsp. coagulans		0.00	0.12		99		
706563	**Good identification**						**706563**
Staphylococcus caprae		100.0	0.33	MAN 95			
706571	**Good identification**						**706571**
Staphylococcus caprae		100.0	0.33	XYL 1			
706572	**Good identification**						**706572**
Staphylococcus caprae		100.0	0.41	XYL 1			
706573	**Very good identification**						**706573**
					COA		
Staphylococcus caprae		98.27	0.59	XYL 1	1		
Staphylococcus aureus subsp. aureus		1.73	0.18		99		
706600	**Low discrimination**						**706600**
					PYR		
Staphylococcus piscifermentans		61.41	0.28	VPT 1	15		
Staphylococcus hominis subsp. hominis		13.60	0.08	ESL 90	1		
Staphylococcus petrasii subsp. petrasii		12.70	-.02		99		
706602	**Low discrimination**						**706602**
Staphylococcus piscifermentans		55.11	0.32	VPT 1			
Staphylococcus simulans		29.16	0.15	ESL 90			
Staphylococcus hominis subsp. hominis		9.55	0.09				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
706603	**Low discrimination**						**706603**
					YEP		
Staphylococcus chromogenes		60.80	0.39	VPT 1	95		
Staphylococcus hyicus		16.09	0.24	GAL 90	1		
Staphylococcus simulans		9.70	0.26		1		
706612	**Low discrimination**						**706612**
Staphylococcus simulans		74.38	0.28	VPT 4			
Staphylococcus piscifermentans		15.06	0.25	BGA 90			
Staphylococcus caprae		4.13	-.09				
706613	**Low discrimination**						**706613**
					COA		
Staphylococcus felis		33.65	0.33	VPT 1	1		
Staphylococcus aureus subsp. aureus		27.92	0.24		99		
Staphylococcus simulans		24.10	0.39		1		
706640	**Low discrimination**						**706640**
					PYR	YEP	
Staphylococcus hominis subsp. hominis		35.32	0.49	ARG 12	1	49	
Staphylococcus petrasii subsp. petrasii		32.99	0.39	PHS 3	99	1	
Staphylococcus petrasii subsp. croceilyticus		10.03	0.18		90	99	
706641	**Very good identification**						**706641**
					YEP	PYR	
Staphylococcus petrasii subsp. petrasii		95.72	0.59	PHS 1	1	99	
Staphylococcus epidermidis		2.84	0.33		1	1	
Staphylococcus chromogenes		0.96	0.20		95	45	
706642	**Low discrimination**						**706642**
					PYR		
Staphylococcus hominis subsp. hominis		49.83	0.50	ARG 12	1		
Staphylococcus petrasii subsp. petrasii		21.35	0.34	PHS 3	99		
Staphylococcus epidermidis		18.04	0.38		1		
706643	**Low discrimination**						**706643**
					YEP	PYR	
Staphylococcus petrasii subsp. petrasii		64.38	0.53	PHS 1	1	99	
Staphylococcus epidermidis		11.23	0.43		1	1	
Staphylococcus chromogenes		10.80	0.39		95	45	
706650	**Low discrimination**						**706650**
					PYR		
Staphylococcus hominis subsp. hominis		24.93	0.32	ARG 12	1		
Staphylococcus warneri		20.96	0.36	PHS 3	8		
Staphylococcus petrasii subsp. petrasii		17.31	0.20	MAN 13	99		
706651	**Low discrimination**						**706651**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		50.03	0.39	PHS 1	1	99	
Staphylococcus aureus subsp. aureus		30.08	0.24	MAN 10	99	10	
Staphylococcus caprae		16.45	0.24		1	75	
706652	**Low discrimination**						**706652**
					COA		
Staphylococcus caprae		44.64	0.32	GAL 99	1		
Staphylococcus aureus subsp. aureus		31.50	0.24	MNS 88	99		
Staphylococcus hominis subsp. hominis		9.84	0.33		1		
706653	**Low discrimination**						**706653**
					COA	PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. aureus		88.58	0.65	GAL 98	99	10	
Staphylococcus caprae		9.30	0.50		1	75	
Staphylococcus petrasii subsp. petrasii		0.80	0.34		1	99	
706700	**Low discrimination**						**706700**
Staphylococcus piscifermentans		67.74	0.32	VPT 1			
Staphylococcus hominis subsp. hominis		10.83	0.09	ESL 90			
Staphylococcus epidermidis		7.61	0.02				
706701	**Low discrimination**						**706701**
					YEP	COA	
Staphylococcus chromogenes		63.06	0.39	VPT 1	95	1	
Staphylococcus hyicus		16.68	0.24	LAC 90	1	24	
Staphylococcus schleiferi subsp. coagulans		9.47	0.27		1	99	
706702	**Low discrimination**						**706702**
					YEP		
Staphylococcus piscifermentans		36.50	0.36	VPT 1	1		
Staphylococcus chromogenes		24.56	0.18	ESL 90	95		
Staphylococcus caprae		14.90	0.06		1		
706703	**Low discrimination**						**706703**
					YEP	COA	
Staphylococcus chromogenes		62.89	0.59	VPT 1	95	1	
Staphylococcus hyicus		21.26	0.45		1	24	
Staphylococcus schleiferi subsp. coagulans		6.45	0.43		1	99	
706712	**Low discrimination**						**706712**
					COA		
Staphylococcus caprae		67.49	0.32	MLT 99	1		
Staphylococcus aureus subsp. aureus		23.57	0.18	MNS 88	99		
Staphylococcus piscifermentans		3.73	0.28		1		
706713	**Low discrimination**						**706713**
					COA		
Staphylococcus aureus subsp. aureus		71.47	0.59	MLT 99	99		
Staphylococcus caprae		15.16	0.50		1		
Staphylococcus felis		5.89	0.43		1		
706740	**Low discrimination**						**706740**
					PYR		
Staphylococcus hominis subsp. hominis		37.98	0.50	ARG 12	1		
Staphylococcus epidermidis		26.66	0.43	PHS 3	1		
Staphylococcus petrasii subsp. petrasii		21.83	0.36		99		
706741	**Low discrimination**						**706741**
					PYR		
Staphylococcus petrasii subsp. petrasii		57.92	0.55	PHS 1	99		
Staphylococcus caprae		16.54	0.39		75		
Staphylococcus epidermidis		14.60	0.49		1		
706742	**Low discrimination**						**706742**
Staphylococcus caprae		47.94	0.47	MAN 95			
Staphylococcus epidermidis		27.84	0.53	MNS 88			
Staphylococcus hominis subsp. hominis		14.70	0.51				
706743	**Low discrimination**						**706743**
					COA	YEP	
Staphylococcus caprae		42.04	0.65	MAN 95	1	1	
Staphylococcus aureus subsp. aureus		38.05	0.59		99	84	

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus chromogenes		9.45	0.59		1	95	
706750	**Low discrimination**						**706750**
					COA	NOV	
Staphylococcus caprae		83.74	0.47	MNS 88	1	1	
Staphylococcus aureus subsp. aureus		5.61	0.18	LAC 95	99	4	
Staphylococcus saprophyticus subsp. bovis		3.99	0.31		1	99	
706751	**Low discrimination**						**706751**
					COA	PYR	
Staphylococcus caprae		51.68	0.65	LAC 95	1	75	
Staphylococcus aureus subsp. aureus		46.77	0.59		99	10	
Staphylococcus petrasii subsp. petrasii		1.06	0.36		1	99	
706752	**Low discrimination**						**706752**
					COA		
Staphylococcus caprae		73.84	0.73	MNS 88	1		
Staphylococcus aureus subsp. aureus		25.79	0.59		99		
Staphylococcus hominis subsp. hominis		0.18	0.34		1		
706753	**Low discrimination**						**706753**
					COA	YEP	
Staphylococcus aureus subsp. aureus		82.44	1.00		99	84	
Staphylococcus caprae		17.48	0.91		1	1	
Staphylococcus chromogenes		0.06	0.48		1	95	
706772	**Low discrimination**						**706772**
					COA		
Staphylococcus caprae		74.11	0.32	XYL 1	1		
Staphylococcus aureus subsp. aureus		25.89	0.18	MNS 88	99		
706773	**Low discrimination**						**706773**
					COA	YEP	
Staphylococcus aureus subsp. aureus		82.46	0.59	XYL 1	99	84	
Staphylococcus caprae		17.49	0.50		1	1	
Staphylococcus chromogenes		0.06	0.07		1	95	
707001	**Good identification**						**707001**
					COA		
Staphylococcus schleiferi subsp. coagulans		96.04	0.37	GAL 99	99		
Staphylococcus schleiferi subsp. schleiferi		3.96	0.04	LAC 86	1		
707003	**Very good identification**						**707003**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.07	0.53	GAL 99	99		
Staphylococcus carnosus subsp. carnosus		0.93	0.07		1		
707011	**Acceptable identification**						**707011**
					COA		
Staphylococcus schleiferi subsp. coagulans		94.11	0.36	GAL 99	99		
Staphylococcus carnosus subsp. carnosus		5.89	0.07	LAC 86	1		
707013	**Good identification**						**707013**
					COA		
Staphylococcus schleiferi subsp. coagulans		96.08	0.52	GAL 99	99		
Staphylococcus carnosus subsp. carnosus		3.92	0.20		1		
707100	**Good identification**						**707100**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	MNS 99			
707101	**Very good identification**						**707101**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.96	0.78	LAC 86	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus schleiferi subsp. schleiferi		0.04	0.04		1	
707102	**Very good identification**					**707102**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.71	0.53	MNS 99	99	
Staphylococcus piscifermentans		0.29	0.11		1	
707103	**Excellent identification**					**707103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.94			
707110	**Good identification**					**707110**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	MNS 99		
707111	**Very good identification**					**707111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.77	LAC 86		
707112	**Good identification**					**707112**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	MNS 99		
707113	**Very good identification**					**707113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.98	0.93		99	
Staphylococcus carnosus subsp. carnosus		0.01	0.07		1	
Staphylococcus caprae		0.01	-.02		1	
707121	**Good identification**					**707121**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	XYL 1		
707123	**Very good identification**					**707123**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	XYL 1		
707131	**Good identification**					**707131**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	XYL 1		
707133	**Good identification**					**707133**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	XYL 1		
707141	**Good identification**					**707141**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.05	0.37	MLT 1	99	
Staphylococcus epidermidis		3.95	0.08	LAC 86	1	
707143	**Good identification**					**707143**
					COA	
Staphylococcus schleiferi subsp. coagulans		93.95	0.53	MLT 1	99	
Staphylococcus caprae		4.07	0.13		1	
Staphylococcus epidermidis		1.99	0.18		1	
707151	**Low discrimination**					**707151**
					COA	
Staphylococcus schleiferi subsp. coagulans		77.65	0.36	MLT 1	99	
Staphylococcus caprae		22.36	0.13	LAC 86	1	
707153	**Low discrimination**					**707153**
					COA	
Staphylococcus schleiferi subsp. coagulans		52.90	0.52	MLT 1	99	
Staphylococcus caprae		47.11	0.39		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
707201	**Good identification**						**707201**
Staphylococcus schleiferi subsp. coagulans		100.0	0.27	GAL 99			
707203	**Good identification**						**707203**
					COA		
Staphylococcus schleiferi subsp. coagulans		98.83	0.43	GAL 99	99		
Staphylococcus epidermidis		1.17	0.02		1		
707213	**Good identification**						**707213**
					COA		
Staphylococcus schleiferi subsp. coagulans		98.64	0.42	GAL 99	99		
Staphylococcus capitis subsp. ureolyticus		1.36	-.11		1		
707240	**Good identification**						**707240**
Staphylococcus epidermidis		100.0	0.28	ESL 1			
707241	**Good identification**						**707241**
Staphylococcus epidermidis		100.0	0.33	ESL 1			
707242	**Good identification**						**707242**
Staphylococcus epidermidis		98.56	0.38	ESL 1			
Staphylococcus piscifermentans		1.44	0.11				
707243	**Good identification**						**707243**
					COA		
Staphylococcus epidermidis		98.09	0.43	ESL 1	1		
Staphylococcus capitis subsp. ureolyticus		1.07	-.11		1		
Staphylococcus schleiferi subsp. coagulans		0.85	0.02		99		
707300	**Low discrimination**						**707300**
					COA		
Staphylococcus schleiferi subsp. coagulans		88.75	0.27	MNS 99	99		
Staphylococcus epidermidis		6.22	0.02	LAC 86	1		
Staphylococcus piscifermentans		5.04	0.11		1		
707301	**Very good identification**						**707301**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.87	0.68	LAC 86	99		
Staphylococcus epidermidis		0.13	0.08		1		
707302	**Good identification**						**707302**
					COA		
Staphylococcus schleiferi subsp. coagulans		95.24	0.43	MNS 99	99		
Staphylococcus epidermidis		3.44	0.12		1		
Staphylococcus piscifermentans		1.32	0.14		1		
707303	**Very good identification**						**707303**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		99.92	0.84		99	1	
Staphylococcus epidermidis		0.07	0.18		1	1	
Staphylococcus chromogenes		0.01	-.04		1	95	
707311	**Very good identification**						**707311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.67	LAC 86			
707312	**Good identification**						**707312**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	MNS 99			
707313	**Very good identification**						**707313**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COA	
Staphylococcus schleiferi subsp. coagulans		99.99	0.83		99	
Staphylococcus caprae		0.01	-.11		1	
707321	**Good identification**					**707321**
Staphylococcus schleiferi subsp. coagulans		100.0	0.27	XYL 1		
707323	**Good identification**					**707323**
Staphylococcus schleiferi subsp. coagulans		100.0	0.43	XYL 1		
707333	**Good identification**					**707333**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	XYL 1		
707340	**Good identification**					**707340**
					NOV	
Staphylococcus epidermidis		97.98	0.43	ESL 1	1	
Staphylococcus hominis subsp. novobiosepticus		1.22	-.04		99	
Staphylococcus piscifermentans		0.80	0.11		1	
707341	**Acceptable identification**					**707341**
					COA	
Staphylococcus epidermidis		92.80	0.49	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		7.20	0.27		99	
707342	**Very good identification**					**707342**
					COA	
Staphylococcus epidermidis		99.34	0.53	ESL 1	1	
Staphylococcus piscifermentans		0.39	0.14		1	
Staphylococcus schleiferi subsp. coagulans		0.28	0.02		99	
707343	**Low discrimination**					**707343**
					COA	
Staphylococcus epidermidis		86.24	0.59	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		12.99	0.43		99	
Staphylococcus caprae		0.66	0.04		1	
707353	**Low discrimination**					**707353**
					COA	
Staphylococcus caprae		36.62	0.30	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		35.11	0.42	TRE 90	99	
Staphylococcus aureus subsp. aureus		15.70	0.18		99	
707400	**Good identification**					**707400**
Staphylococcus piscifermentans		100.0	0.44	VPT 1		
707402	**Good identification**					**707402**
Staphylococcus piscifermentans		100.0	0.48	VPT 1		
707410	**Good identification**					**707410**
Staphylococcus piscifermentans		100.0	0.37	VPT 1		
707412	**Good identification**					**707412**
Staphylococcus piscifermentans		100.0	0.41	VPT 1		
707440	**Good identification**					**707440**
Staphylococcus piscifermentans		100.0	0.44	VPT 1		
707442	**Good identification**					**707442**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus piscifermentans		100.0	0.48	VPT 1		
707450	**Good identification**					**707450**
Staphylococcus piscifermentans		100.0	0.37	VPT 1		
707452	**Low discrimination**					**707452**
Staphylococcus piscifermentans		89.01	0.41	VPT 1		
Staphylococcus caprae		11.00	0.00			
707500	**Good identification**					**707500**
Staphylococcus piscifermentans		100.0	0.48	VPT 1		
707501	**Low discrimination**					**707501**
					COA	
Staphylococcus schleiferi subsp. coagulans		89.00	0.37	TRE 1	99	
Staphylococcus schleiferi subsp. schleiferi		11.00	0.14	LAC 86	1	
707502	**Good identification**					**707502**
					COA	
Staphylococcus piscifermentans		96.62	0.52	VPT 1	1	
Staphylococcus schleiferi subsp. coagulans		3.38	0.12		99	
707503	**Very good identification**					**707503**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.32	0.53	TRE 1	99	
Staphylococcus caprae		0.39	-.09		1	
Staphylococcus piscifermentans		0.29	0.11		1	
707510	**Good identification**					**707510**
Staphylococcus piscifermentans		100.0	0.41	VPT 1		
707511	**Good identification**					**707511**
					COA	
Staphylococcus schleiferi subsp. coagulans		97.45	0.36	TRE 1	99	
Staphylococcus caprae		2.55	-.09	LAC 86	1	
707512	**Low discrimination**					**707512**
					COA	
Staphylococcus piscifermentans		86.38	0.44	VPT 1	1	
Staphylococcus caprae		7.11	0.00		1	
Staphylococcus schleiferi subsp. coagulans		6.51	0.11		99	
707513	**Acceptable identification**					**707513**
					COA	
Staphylococcus schleiferi subsp. coagulans		91.65	0.52	TRE 1	99	
Staphylococcus caprae		7.42	0.18		1	
Staphylococcus felis		0.53	-.04		1	
707540	**Good identification**					**707540**
Staphylococcus piscifermentans		98.57	0.48	VPT 1		
Staphylococcus caprae		1.43	-.12			
707542	**Low discrimination**					**707542**
Staphylococcus piscifermentans		84.47	0.52	VPT 1		
Staphylococcus caprae		15.53	0.15			
707543	**Good identification**					**707543**
					COA	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus caprae		96.77	0.33	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		2.51	0.12	MAN 95	99	
Staphylococcus piscifermentans		0.73	0.11		1	
707550	**Low discrimination**					**707550**
Staphylococcus piscifermentans		60.84	0.41	VPT 1		
Staphylococcus caprae		39.16	0.15			
707551	**Good identification**					**707551**
Staphylococcus caprae		100.0	0.33	ESL 1		
707552	**Low discrimination**					**707552**
Staphylococcus caprae		89.08	0.41	ESL 1		
Staphylococcus piscifermentans		10.93	0.44	MNS 88		
707553	**Very good identification**					**707553**
					COA	
Staphylococcus caprae		98.15	0.59	ESL 1	1	
Staphylococcus aureus subsp. aureus		1.73	0.18		99	
Staphylococcus schleiferi subsp. coagulans		0.12	0.11		99	
707600	**Good identification**					**707600**
Staphylococcus piscifermentans		100.0	0.48	VPT 1		
707602	**Good identification**					**707602**
Staphylococcus piscifermentans		100.0	0.52	VPT 1		
707610	**Good identification**					**707610**
Staphylococcus piscifermentans		100.0	0.41	VPT 1		
707612	**Good identification**					**707612**
Staphylococcus piscifermentans		100.0	0.44	VPT 1		
707640	**Good identification**					**707640**
					PYR	
Staphylococcus piscifermentans		95.46	0.48	VPT 1	15	
Staphylococcus hominis subsp. hominis		2.35	0.08		1	
Staphylococcus petrasii subsp. petrasii		2.19	-.02		99	
707642	**Good identification**					**707642**
Staphylococcus piscifermentans		98.11	0.52	VPT 1		
Staphylococcus hominis subsp. hominis		1.89	0.09			
707650	**Good identification**					**707650**
Staphylococcus piscifermentans		100.0	0.41	VPT 1		
707652	**Good identification**					**707652**
					COA	
Staphylococcus piscifermentans		95.06	0.44	VPT 1	1	
Staphylococcus caprae		2.90	-.09		1	
Staphylococcus aureus subsp. aureus		2.04	-.17		99	
707700	**Good identification**					**707700**
Staphylococcus piscifermentans		100.0	0.52	VPT 1		
707701	**Low discrimination**					**707701**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		88.76	0.27	TRE 1	99	1	
Staphylococcus chromogenes		6.21	-.02	LAC 86	1	95	
Staphylococcus piscifermentans		5.04	0.11		1	1	
707702	**Very good identification**						**707702**
					COA		
Staphylococcus piscifermentans		99.27	0.55	VPT 1	1		
Staphylococcus schleiferi subsp. coagulans		0.73	0.02		99		
707703	**Low discrimination**						**707703**
					YEP	COA	
Staphylococcus schleiferi subsp. coagulans		86.88	0.43	TRE 1	1	99	
Staphylococcus chromogenes		8.91	0.18		95	1	
Staphylococcus hyicus		3.01	0.04		1	24	
707710	**Good identification**						**707710**
Staphylococcus piscifermentans		100.0	0.44	VPT 1			
707712	**Good identification**						**707712**
					COA		
Staphylococcus piscifermentans		96.51	0.48	VPT 1	1		
Staphylococcus caprae		1.96	-.09		1		
Staphylococcus schleiferi subsp. coagulans		1.53	0.01		99		
707713	**Low discrimination**						**707713**
					COA		
Staphylococcus schleiferi subsp. coagulans		62.11	0.42	TRE 1	99		
Staphylococcus aureus subsp. aureus		27.77	0.18		99		
Staphylococcus caprae		5.89	0.09		1		
707740	**Good identification**						**707740**
Staphylococcus piscifermentans		96.11	0.52	VPT 1			
Staphylococcus hominis subsp. hominis		1.71	0.09				
Staphylococcus epidermidis		1.20	0.02				
707742	**Good identification**						**707742**
Staphylococcus piscifermentans		92.12	0.55	VPT 1			
Staphylococcus caprae		4.18	0.06				
Staphylococcus epidermidis		2.43	0.12				
707750	**Low discrimination**						**707750**
Staphylococcus piscifermentans		86.30	0.44	VPT 1			
Staphylococcus caprae		13.70	0.06				
707752	**Low discrimination**						**707752**
					COA		
Staphylococcus caprae		54.15	0.32	ESL 1	1		
Staphylococcus piscifermentans		26.94	0.48	MNS 88	1		
Staphylococcus aureus subsp. aureus		18.91	0.18		99		
707753	**Low discrimination**						**707753**
					COA	YEP	
Staphylococcus aureus subsp. aureus		82.44	0.59	ESL 1	99	84	
Staphylococcus caprae		17.48	0.50		1	1	
Staphylococcus chromogenes		0.06	0.07		1	95	
710253	**Good identification**						**710253**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. ureolyticus		100.0	0.33	ORN 1		
710342	**Good identification**					**710342**
					NOV	
Staphylococcus epidermidis		96.47	0.30	ORN 12	1	
Staphylococcus hominis subsp. novobiosepticus		3.53	-.07	NIT 97	99	
710343	**Good identification**					**710343**
Staphylococcus epidermidis		100.0	0.36	ORN 12		
710640	**Low discrimination**					**710640**
					PYR	
Staphylococcus warneri		57.66	0.47	ORN 1	8	
Staphylococcus petrasii subsp. petrasii		14.78	0.20		99	
Staphylococcus petrasii subsp. jettensis		13.03	0.10		99	
710641	**Acceptable identification**					**710641**
					OXI	
Staphylococcus petrasii subsp. petrasii		93.84	0.39	ORN 1	1	
Staphylococcus lugdunensis		4.37	0.13	NIT 90	1	
Kocuria kristinae		0.97	-.02		99	
710642	**Low discrimination**					**710642**
					PYR	
Staphylococcus warneri		62.02	0.37	ORN 1	8	
Staphylococcus petrasii subsp. petrasii		25.69	0.14		99	
Staphylococcus petrasii subsp. jettensis		7.42	-.05		99	
710643	**Low discrimination**					**710643**
Staphylococcus petrasii subsp. petrasii		56.25	0.34	ORN 1		
Staphylococcus lugdunensis		43.75	0.33	NIT 90		
710650	**Low discrimination**					**710650**
					NOV	
Staphylococcus warneri		76.28	0.44	ORN 1	1	
Staphylococcus saprophyticus subsp. saprophyticus		11.69	0.06		99	
Staphylococcus pasteuri		9.03	0.10		1	
710740	**Low discrimination**					**710740**
					PYR	
Staphylococcus warneri		64.88	0.36	ORN 1	8	
Staphylococcus petrasii subsp. petrasii		35.12	0.16		99	
710741	**Acceptable identification**					**710741**
Staphylococcus petrasii subsp. petrasii		93.48	0.36	ORN 1		
Staphylococcus lugdunensis		6.53	0.13	NIT 90		
710743	**Low discrimination**					**710743**
Staphylococcus lugdunensis		53.85	0.33	ARG 5		
Staphylococcus petrasii subsp. petrasii		46.15	0.30	NIT 99		
710750	**Acceptable identification**					**710750**
					PYR	
Staphylococcus warneri		92.33	0.33	ORN 1	8	
Staphylococcus petrasii subsp. petrasii		7.67	-.04		99	
712053	**Good identification**					**712053**
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	ORN 1		
712213	**Good identification**					**712213**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	ORN 1		
712240	**Low discrimination**					**712240**
					NOV	
Staphylococcus epidermidis		66.95	0.35	ORN 12	1	
Staphylococcus hominis subsp. hominis		21.59	0.29		1	
Staphylococcus hominis subsp. novobiosepticus		7.57	0.08		99	
712241	**Low discrimination**					**712241**
					PYR	
Staphylococcus epidermidis		70.09	0.41	ORN 12	1	
Staphylococcus capitis subsp. ureolyticus		13.69	0.13		1	
Staphylococcus petrasii subsp. petrasii		10.02	0.18		99	
712242	**Low discrimination**					**712242**
					NOV	
Staphylococcus epidermidis		87.16	0.46	ORN 12	1	
Staphylococcus hominis subsp. hominis		10.42	0.30		1	
Staphylococcus hominis subsp. novobiosepticus		1.68	0.03		99	
712243	**Low discrimination**					**712243**
					PYR	
Staphylococcus epidermidis		59.36	0.51	ORN 12	1	
Staphylococcus capitis subsp. ureolyticus		26.85	0.31		1	
Staphylococcus lugdunensis		12.35	0.33		96	
712251	**Good identification**					**712251**
					PYR	
Staphylococcus capitis subsp. ureolyticus		99.45	0.41	ORN 1	1	
Staphylococcus petrasii subsp. petrasii		0.34	-.02	LAC 88	99	
Staphylococcus epidermidis		0.21	0.00		1	
712253	**Very good identification**					**712253**
Staphylococcus capitis subsp. ureolyticus		99.91	0.59	ORN 1		
Staphylococcus epidermidis		0.09	0.10			
712340	**Low discrimination**					**712340**
					NOV	
Staphylococcus epidermidis		88.89	0.51	ORN 12	1	
Staphylococcus hominis subsp. hominis		5.48	0.29		1	
Staphylococcus hominis subsp. novobiosepticus		5.32	0.18		99	
712341	**Very good identification**					**712341**
					PYR	
Staphylococcus epidermidis		96.36	0.56	ORN 12	1	
Staphylococcus petrasii subsp. petrasii		1.62	0.14		99	
Staphylococcus lugdunensis		1.25	0.13		96	
712342	**Very good identification**					**712342**
					NOV	
Staphylococcus epidermidis		96.80	0.61	ORN 12	1	
Staphylococcus hominis subsp. hominis		2.21	0.31		1	
Staphylococcus hominis subsp. novobiosepticus		0.99	0.12		99	
712343	**Very good identification**					**712343**
					PYR	
Staphylococcus epidermidis		95.99	0.67	ORN 12	1	
Staphylococcus lugdunensis		3.52	0.33		96	
Staphylococcus petrasii subsp. petrasii		0.28	0.09		99	
712443	**Low discrimination**					**712443**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lugdunensis		89.54	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		10.47	0.12	SUC 99			
712543	**Acceptable identification**						**712543**
Staphylococcus lugdunensis		90.96	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		7.09	0.09	SUC 99			
Staphylococcus caprae		1.95	-.09				
712603	**Low discrimination**						**712603**
Staphylococcus lugdunensis		89.54	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		10.47	0.12	MLT 99			
712640	**Low discrimination**						**712640**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		46.21	0.39	ORN 1	99	1	
Staphylococcus hominis subsp. hominis		16.16	0.39	MNS 90	1	49	
Staphylococcus petrasii subsp. croceilyticus		14.05	0.18		90	99	
712641	**Low discrimination**						**712641**
					PYR		
Staphylococcus petrasii subsp. petrasii		66.08	0.59	ORN 1	99		
Staphylococcus lugdunensis		33.82	0.54		96		
Staphylococcus hominis subsp. hominis		0.05	0.04		1		
712642	**Low discrimination**						**712642**
					PYR		
Staphylococcus petrasii subsp. petrasii		35.89	0.34	ORN 1	99		
Staphylococcus lugdunensis		27.91	0.33	MNS 90	96		
Staphylococcus hominis subsp. hominis		27.35	0.40		1		
712643	**Low discrimination**						**712643**
					PYR		
Staphylococcus lugdunensis		89.46	0.74	ARG 5	96		
Staphylococcus petrasii subsp. petrasii		10.46	0.53		99		
Staphylococcus epidermidis		0.04	0.10		1		
712650	**Low discrimination**						**712650**
					PYR	YEP	
Staphylococcus warneri		32.86	0.36	ORN 1	8	43	
Staphylococcus petrasii subsp. petrasii		27.15	0.20		99	1	
Staphylococcus pasteuri		22.23	0.18		1	86	
712651	**Acceptable identification**						**712651**
					PYR		
Staphylococcus petrasii subsp. petrasii		92.88	0.39	ORN 1	99		
Staphylococcus lugdunensis		4.32	0.13	MAN 10	96		
Staphylococcus capitis subsp. ureolyticus		2.80	0.00		1		
712653	**Low discrimination**						**712653**
					PYR		
Staphylococcus petrasii subsp. petrasii		45.21	0.34	ORN 1	99		
Staphylococcus lugdunensis		35.16	0.33	MAN 10	96		
Staphylococcus capitis subsp. ureolyticus		18.53	0.18		1		
712663	**Low discrimination**						**712663**
Staphylococcus lugdunensis		89.54	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		10.47	0.12	XYL 1			
712703	**Acceptable identification**						**712703**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus lugdunensis		92.77	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		7.23	0.09	MLT 99			
712740	**Low discrimination**						**712740**
					PYR		
Staphylococcus petrasii subsp. petrasii		55.12	0.36	ORN 1	99		
Staphylococcus hominis subsp. hominis		31.32	0.40	MNS 90	1		
Staphylococcus warneri		4.85	0.29		8		
712741	**Low discrimination**						**712741**
					PYR		
Staphylococcus petrasii subsp. petrasii		56.33	0.55	ORN 1	99		
Staphylococcus lugdunensis		43.25	0.54		96		
Staphylococcus epidermidis		0.34	0.15		1		
712742	**Low discrimination**						**712742**
					PYR		
Staphylococcus hominis subsp. hominis		33.53	0.41	ARG 12	1		
Staphylococcus lugdunensis		31.58	0.33	ORN 1	96		
Staphylococcus petrasii subsp. petrasii		27.07	0.30		99		
712743	**Good identification**						**712743**
					PYR		
Staphylococcus lugdunensis		92.51	0.74	ARG 5	96		
Staphylococcus petrasii subsp. petrasii		7.21	0.50		99		
Staphylococcus epidermidis		0.26	0.26		1		
712751	**Acceptable identification**						**712751**
Staphylococcus petrasii subsp. petrasii		93.48	0.36	ORN 1			
Staphylococcus lugdunensis		6.53	0.13	MAN 10			
712753	**Low discrimination**						**712753**
					COA	PYR	
Staphylococcus lugdunensis		36.79	0.33	ARG 5	1	96	
Staphylococcus petrasii subsp. petrasii		31.53	0.30	MAN 1	1	99	
Staphylococcus aureus subsp. aureus		26.14	0.18		99	10	
712763	**Acceptable identification**						**712763**
Staphylococcus lugdunensis		92.77	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		7.23	0.09	XYL 1			
713643	**Low discrimination**						**713643**
Staphylococcus lugdunensis		89.54	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		10.47	0.12	ESL 1			
713743	**Acceptable identification**						**713743**
Staphylococcus lugdunensis		92.77	0.33	ARG 5			
Staphylococcus petrasii subsp. petrasii		7.23	0.09	ESL 1			
714242	**Good identification**						**714242**
Staphylococcus epidermidis		100.0	0.30	ORN 12			
714243	**Good identification**						**714243**
Staphylococcus epidermidis		100.0	0.36	ORN 12			
714340	**Good identification**						**714340**

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus epidermidis		100.0	0.35	ORN 12		
714341	**Good identification**					**714341**
Staphylococcus epidermidis		100.0	0.41	ORN 12		
714342	**Good identification**					**714342**
Staphylococcus epidermidis		100.0	0.46	ORN 12		
714343	**Good identification**					**714343**
Staphylococcus epidermidis		100.0	0.51	ORN 12		
714553	**Good identification**					**714553**
Staphylococcus caprae		100.0	0.39	ORN 1		
714753	**Low discrimination**					**714753**
					COA	
Staphylococcus caprae		70.00	0.30	ORN 1	1	
Staphylococcus aureus subsp. aureus		30.00	0.18	NIT 90	99	
716101	**Good identification**					**716101**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.19	0.37	ORN 1	99	
Staphylococcus schleiferi subsp. schleiferi		3.81	0.04	LAC 86	1	
716103	**Very good identification**					**716103**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.72	0.53	ORN 1	99	
Staphylococcus epidermidis		0.28	0.00		1	
716111	**Good identification**					**716111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	ORN 1		
716113	**Good identification**					**716113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.14	0.53	ORN 1	99	
Staphylococcus caprae		0.86	-.02		1	
716141	**Good identification**					**716141**
Staphylococcus epidermidis		100.0	0.31	ORN 12		
716142	**Good identification**					**716142**
Staphylococcus epidermidis		96.31	0.36	ORN 12		
Staphylococcus caprae		3.69	-.05	SUC 99		
716143	**Low discrimination**					**716143**
					COA	
Staphylococcus epidermidis		84.17	0.41	ORN 12	1	
Staphylococcus caprae		12.74	0.13	SUC 99	1	
Staphylococcus schleiferi subsp. coagulans		3.10	0.12		99	
716153	**Good identification**					**716153**
					COA	
Staphylococcus caprae		98.49	0.39	ORN 1	1	
Staphylococcus schleiferi subsp. coagulans		1.16	0.12	TRE 90	99	
Staphylococcus epidermidis		0.35	0.00		1	
716240	**Good identification**					**716240**
Staphylococcus epidermidis		100.0	0.51	ORN 12		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
716241	**Very good identification**						**716241**
Staphylococcus epidermidis		100.0	0.56	ORN 12			
716242	**Very good identification**						**716242**
Staphylococcus epidermidis		100.0	0.61	ORN 12			
716243	**Very good identification**						**716243**
Staphylococcus epidermidis		99.92	0.67	ORN 12			
Staphylococcus capitis subsp. ureolyticus		0.08	-.11				
716301	**Low discrimination**						**716301**
					COA		
Staphylococcus epidermidis		62.79	0.31	ORN 12	1		
Staphylococcus schleiferi subsp. coagulans		37.21	0.27	MLT 99	99		
716302	**Good identification**						**716302**
					COA		
Staphylococcus epidermidis		97.89	0.36	ORN 12	1		
Staphylococcus schleiferi subsp. coagulans		2.11	0.02	MLT 99	99		
716303	**Low discrimination**						**716303**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		53.23	0.43	ORN 1	99	1	
Staphylococcus epidermidis		46.31	0.41		1	1	
Staphylococcus chromogenes		0.46	-.04		1	95	
716313	**Good identification**						**716313**
					COA		
Staphylococcus schleiferi subsp. coagulans		98.07	0.42	ORN 1	99		
Staphylococcus caprae		0.99	-.11		1		
Staphylococcus epidermidis		0.93	0.00		1		
716340	**Very good identification**						**716340**
Staphylococcus epidermidis		100.0	0.66	ORN 12			
716341	**Very good identification**						**716341**
Staphylococcus epidermidis		100.0	0.72	ORN 12			
716342	**Very good identification**						**716342**
Staphylococcus epidermidis		100.0	0.77	ORN 12			
716343	**Very good identification**						**716343**
					COA		
Staphylococcus epidermidis		99.92	0.82	ORN 12	1		
Staphylococcus caprae		0.06	0.04		1		
Staphylococcus schleiferi subsp. coagulans		0.01	0.02		99		
716351	**Low discrimination**						**716351**
Staphylococcus epidermidis		84.95	0.31	ORN 12			
Staphylococcus caprae		15.05	0.04	MAN 1			
716352	**Low discrimination**						**716352**
Staphylococcus epidermidis		78.79	0.36	ORN 12			
Staphylococcus caprae		21.21	0.13	MAN 1			
716353	**Low discrimination**						**716353**
					COA		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus caprae		37.69	0.30	ORN 1	1	
Staphylococcus epidermidis		35.45	0.41	TRE 90	1	
Staphylococcus aureus subsp. aureus		16.16	0.18		99	
716361	**Good identification**					**716361**
Staphylococcus epidermidis		100.0	0.31	ORN 12		
716362	**Good identification**					**716362**
Staphylococcus epidermidis		100.0	0.36	ORN 12		
716363	**Good identification**					**716363**
Staphylococcus epidermidis		100.0	0.41	ORN 12		
716543	**Good identification**					**716543**
Staphylococcus caprae		99.26	0.33	ORN 1		
Staphylococcus epidermidis		0.74	0.00	MAN 95		
716551	**Good identification**					**716551**
Staphylococcus caprae		100.0	0.33	ORN 1		
716552	**Good identification**					**716552**
Staphylococcus caprae		100.0	0.41	ORN 1		
716553	**Very good identification**					**716553**
					COA	
Staphylococcus caprae		98.27	0.59	ORN 1	1	
Staphylococcus aureus subsp. aureus		1.73	0.18		99	
716643	**Low discrimination**					**716643**
					PYR	
Staphylococcus lugdunensis		70.12	0.33	ARG 5	96	
Staphylococcus epidermidis		19.29	0.26	PHS 1	1	
Staphylococcus petrasii subsp. petrasii		8.20	0.12		99	
716741	**Low discrimination**					**716741**
					PYR	
Staphylococcus epidermidis		60.85	0.31	ORN 12	1	
Staphylococcus petrasii subsp. petrasii		17.89	0.14	TRE 1	99	
Staphylococcus lugdunensis		13.73	0.13		96	
716742	**Low discrimination**					**716742**
Staphylococcus epidermidis		85.72	0.36	ORN 12		
Staphylococcus caprae		10.93	0.06	TRE 1		
Staphylococcus hominis subsp. hominis		3.35	0.10			
716743	**Low discrimination**					**716743**
					PYR	
Staphylococcus epidermidis		36.18	0.41	ORN 12	1	
Staphylococcus lugdunensis		23.21	0.33	TRE 1	96	
Staphylococcus caprae		18.22	0.24		75	
716752	**Low discrimination**					**716752**
					COA	
Staphylococcus caprae		74.11	0.32	ORN 1	1	
Staphylococcus aureus subsp. aureus		25.89	0.18	MNS 88	99	
716753	**Low discrimination**					**716753**
					COA YEP	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. aureus		82.44	0.59	ORN 1	99	84	
Staphylococcus caprae		17.48	0.50		1	1	
Staphylococcus chromogenes		0.06	0.07		1	95	
717101	**Good identification**						**717101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	ORN 1			
717103	**Very good identification**						**717103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	ORN 1			
717111	**Good identification**						**717111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	ORN 1			
717113	**Good identification**						**717113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	ORN 1			
717301	**Good identification**						**717301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.27	ORN 1			
717303	**Good identification**						**717303**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.09	0.43	ORN 1	99		
Staphylococcus epidermidis		0.91	0.00		1		
717313	**Good identification**						**717313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ORN 1			
717341	**Good identification**						**717341**
Staphylococcus epidermidis		100.0	0.31	ORN 12			
717342	**Good identification**						**717342**
Staphylococcus epidermidis		100.0	0.36	ORN 12			
717343	**Good identification**						**717343**
					COA		
Staphylococcus epidermidis		98.90	0.41	ORN 12	1		
Staphylococcus schleiferi subsp. coagulans		1.10	0.02	ESL 1	99		
720053	**Good identification**						**720053**
Staphylococcus capitis subsp. ureolyticus		100.0	0.28	BGA 12			
720213	**Good identification**						**720213**
Staphylococcus capitis subsp. ureolyticus		95.47	0.28	BGA 12			
Staphylococcus simulans		4.53	0.13	NIT 95			
720240	**Acceptable identification**						**720240**
					PYR		
Staphylococcus warneri		90.60	0.49	TRE 95	8		
Staphylococcus petrasii subsp. petrasii		4.46	0.07		99		
Staphylococcus epidermidis		3.12	0.10		1		
720241	**Low discrimination**						**720241**
					PYR		
Staphylococcus petrasii subsp. petrasii		61.93	0.27	NIT 90	99		
Staphylococcus capitis subsp. ureolyticus		21.86	0.10	TRE 99	1		
Staphylococcus epidermidis		8.93	0.15		1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
720242	**Low discrimination**						**720242**
					NOV		
Staphylococcus warneri		56.09	0.39	TRE 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		19.18	0.09		99		
Staphylococcus epidermidis		18.32	0.20		1		
720243	**Low discrimination**						**720243**
					PYR		
Staphylococcus capitis subsp. ureolyticus		72.23	0.28	BGA 12	1		
Staphylococcus petrasii subsp. petrasii		15.03	0.21	NIT 95	99		
Staphylococcus epidermidis		12.74	0.26	MAN 96	1		
720250	**Low discrimination**						**720250**
					NOV		
Staphylococcus warneri		58.18	0.46	TRE 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		38.78	0.22		99		
Staphylococcus capitis subsp. ureolyticus		3.04	-.03		1		
720251	**Good identification**						**720251**
					PYR		
Staphylococcus capitis subsp. ureolyticus		98.32	0.38	BGA 12	1		
Staphylococcus petrasii subsp. petrasii		1.29	0.07	NIT 95	99		
Staphylococcus warneri		0.39	0.11	LAC 88	8		
720252	**Low discrimination**						**720252**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		84.05	0.37	ARG 8	99		
Staphylococcus capitis subsp. ureolyticus		8.54	0.15	TRE 99	1		
Staphylococcus warneri		7.42	0.36		1		
720253	**Very good identification**						**720253**
					NOV	PYR	
Staphylococcus capitis subsp. ureolyticus		99.72	0.56	BGA 12	1	1	
Staphylococcus saprophyticus subsp. saprophyticus		0.10	-.04	NIT 95	99	7	
Staphylococcus petrasii subsp. petrasii		0.10	0.02		1	99	
720340	**Low discrimination**						**720340**
					PYR		
Staphylococcus warneri		55.76	0.39	TRE 95	8		
Staphylococcus epidermidis		34.41	0.25		1		
Staphylococcus petrasii subsp. petrasii		5.79	0.04		99		
720341	**Low discrimination**						**720341**
					PYR		
Staphylococcus epidermidis		55.07	0.31	NIT 97	1		
Staphylococcus petrasii subsp. petrasii		44.93	0.23		99		
720342	**Low discrimination**						**720342**
					PYR		
Staphylococcus epidermidis		82.02	0.36	NIT 97	1		
Staphylococcus warneri		13.99	0.29		8		
Staphylococcus petrasii subsp. petrasii		2.35	-.02		99		
720343	**Low discrimination**						**720343**
					PYR		
Staphylococcus epidermidis		87.04	0.41	NIT 97	1		
Staphylococcus petrasii subsp. petrasii		12.08	0.18		99		
Staphylococcus capitis subsp. ureolyticus		0.88	-.14		1		
720350	**Low discrimination**						**720350**
					NOV		
Staphylococcus warneri		84.49	0.36	TRE 95	1		

Code	Taxon	ID score	T index	tests against		Differentiation tests		Code
Staphylococcus saprophyticus subsp. saprophyticus		15.51	0.00			99		
720440	**Low discrimination**							**720440**
						NOV	PYR	
Staphylococcus warneri		79.04	0.49	SUC	95	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		15.56	0.14			99	7	
Staphylococcus petrasii subsp. petrasii		3.89	0.07			1	99	
720441	**Good identification**							**720441**
						PYR		
Staphylococcus petrasii subsp. petrasii		95.60	0.27	NIT	90	99		
Staphylococcus warneri		4.41	0.14	SUC	99	8		
720442	**Low discrimination**							**720442**
						NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		75.62	0.29	ARG	8	99	7	
Staphylococcus warneri		22.59	0.39	SUC	91	1	8	
Staphylococcus petrasii subsp. petrasii		1.80	0.02	MAN	96	1	99	
720450	**Low discrimination**							**720450**
						NOV		
Staphylococcus saprophyticus subsp. saprophyticus		86.71	0.42	ARG	8	99		
Staphylococcus warneri		13.29	0.46	SUC	91	1		
720452	**Very good identification**							**720452**
						NOV		
Staphylococcus saprophyticus subsp. saprophyticus		99.11	0.58	ARG	8	99		
Staphylococcus warneri		0.89	0.36	SUC	91	1		
720540	**Low discrimination**							**720540**
						NOV	PYR	
Staphylococcus warneri		86.35	0.39	SUC	95	1	8	
Staphylococcus petrasii subsp. petrasii		8.97	0.04			1	99	
Staphylococcus saprophyticus subsp. saprophyticus		4.68	-.08			99	7	
720542	**Low discrimination**							**720542**
						NOV	PYR	
Staphylococcus warneri		47.86	0.29	SUC	95	1	8	
Staphylococcus saprophyticus subsp. saprophyticus		44.11	0.07			99	7	
Staphylococcus petrasii subsp. petrasii		8.03	-.02			1	99	
720552	**Good identification**							**720552**
						NOV		
Staphylococcus saprophyticus subsp. saprophyticus		96.83	0.36	ARG	8	99		
Staphylococcus warneri		3.17	0.26	GAL	8	1		
720600	**Acceptable identification**							**720600**
						NOV	PYR	
Staphylococcus warneri		93.52	0.49	MLT	95	1	8	
Staphylococcus petrasii subsp. petrasii		4.60	0.07			1	99	
Staphylococcus saprophyticus subsp. saprophyticus		1.88	-.07			99	7	
720601	**Low discrimination**							**720601**
						OXI	PYR	
Staphylococcus petrasii subsp. petrasii		83.54	0.27	NIT	90	1	99	
Kocuria kristinae		12.61	0.10	MLT	99	99	93	
Staphylococcus warneri		3.85	0.14			1	8	
720602	**Low discrimination**							**720602**
						NOV		
Staphylococcus warneri		60.49	0.39	MLT	95	1		
Staphylococcus saprophyticus subsp. saprophyticus		20.68	0.09			99		
Staphylococcus simulans		14.02	0.16			1		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
720603	**Low discrimination**						**720603**
					OXI		
Staphylococcus simulans		50.33	0.27	VPT 4	1		
Staphylococcus petrasii subsp. petrasii		46.42	0.21	NIT 96	1		
Kocuria kristinae		3.25	-.03		99		
720610	**Low discrimination**						**720610**
					NOV		
Staphylococcus warneri		60.01	0.46	MLT 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		40.00	0.22		99		
720612	**Low discrimination**						**720612**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		83.25	0.37	ARG 8	99		
Staphylococcus simulans		9.40	0.29	MLT 99	1		
Staphylococcus warneri		7.35	0.36		1		
720613	**Acceptable identification**						**720613**
					NOV	PYR	
Staphylococcus simulans		94.03	0.40	VPT 4	1	70	
Staphylococcus saprophyticus subsp. saprophyticus		2.51	-.04	NIT 96	99	7	
Staphylococcus petrasii subsp. petrasii		2.41	0.02		1	99	
720640	**Low discrimination**						**720640**
					NOV	PYR	
Staphylococcus warneri		72.35	0.75		1	8	
Staphylococcus petrasii subsp. petrasii		18.55	0.48		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		7.58	0.34		99	7	
720641	**Very good identification**						**720641**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		97.75	0.68	NIT 90	1	99	
Kocuria kristinae		1.34	0.30		99	93	
Staphylococcus warneri		0.86	0.40		1	8	
720642	**Low discrimination**						**720642**
					NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		55.57	0.50	ARG 8	99	7	
Staphylococcus warneri		31.19	0.65	MAN 96	1	8	
Staphylococcus petrasii subsp. petrasii		12.92	0.43		1	99	
720643	**Very good identification**						**720643**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		97.84	0.62	NIT 90	1	99	
Kocuria kristinae		0.62	0.17		99	93	
Staphylococcus warneri		0.54	0.31		1	8	
720650	**Low discrimination**						**720650**
					NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		76.89	0.63	ARG 8	99	7	
Staphylococcus warneri		22.14	0.72		1	8	
Staphylococcus petrasii subsp. petrasii		0.87	0.29		1	99	
720651	**Low discrimination**						**720651**
					NOV	PYR	
Staphylococcus petrasii subsp. petrasii		85.50	0.48	NIT 90	1	99	
Staphylococcus saprophyticus subsp. saprophyticus		8.47	0.22	MAN 10	99	7	
Staphylococcus warneri		4.93	0.38		1	8	
720652	**Very good identification**						**720652**
					NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		98.17	0.78	ARG 8	99	7	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		1.66	0.63		1	8	
Staphylococcus petrasii subsp. petrasii		0.11	0.23		1	99	
720653	**Low discrimination**						**720653**
					NOV	PYR	
Staphylococcus saprophyticus subsp. saprophyticus		44.15	0.37	ARG 8	99	7	
Staphylococcus petrasii subsp. petrasii		42.35	0.43	MNS 1	1	99	
Staphylococcus simulans		7.50	0.32		1	70	
720660	**Low discrimination**						**720660**
					NOV	PYR	
Staphylococcus warneri		73.47	0.34	XYL 1	1	8	
Staphylococcus petrasii subsp. petrasii		18.84	0.07		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		7.70	-.07		99	7	
720661	**Good identification**						**720661**
Staphylococcus petrasii subsp. petrasii		100.0	0.27	NIT 90			
720672	**Good identification**						**720672**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		98.33	0.37	ARG 8	99		
Staphylococcus warneri		1.67	0.21	XYL 1	1		
720700	**Acceptable identification**						**720700**
					PYR		
Staphylococcus warneri		90.59	0.39	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		9.41	0.04		99		
720702	**Low discrimination**						**720702**
					PYR		
Staphylococcus warneri		85.63	0.29	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		14.37	-.02		99		
720710	**Low discrimination**						**720710**
					NOV		
Staphylococcus warneri		84.49	0.36	MLT 95	1		
Staphylococcus saprophyticus subsp. saprophyticus		15.51	0.00		99		
720740	**Low discrimination**						**720740**
					NOV	PYR	
Staphylococcus warneri		63.57	0.65		1	8	
Staphylococcus petrasii subsp. petrasii		34.41	0.45		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		1.83	0.12		99	7	
720741	**Very good identification**						**720741**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		98.91	0.64	NIT 90	1	99	
Kocuria kristinae		0.68	0.20		99	93	
Staphylococcus warneri		0.41	0.30		1	8	
720742	**Low discrimination**						**720742**
					NOV	PYR	
Staphylococcus warneri		42.29	0.55		1	8	
Staphylococcus petrasii subsp. petrasii		36.97	0.39		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		20.74	0.28		99	7	
720743	**Very good identification**						**720743**
					OXI	PYR	
Staphylococcus petrasii subsp. petrasii		99.35	0.59	NIT 90	1	99	
Kocuria kristinae		0.32	0.07		99	93	
Staphylococcus warneri		0.26	0.20		1	8	
720750	**Low discrimination**						**720750**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	PYR		
Staphylococcus warneri		48.91	0.62		1	8		
Staphylococcus saprophyticus subsp. saprophyticus		46.77	0.41		99	7		
Staphylococcus petrasii subsp. petrasii		4.06	0.25		1	99		
720751	**Good identification**							**720751**
					NOV	PYR		
Staphylococcus petrasii subsp. petrasii		96.13	0.45	NIT 90	1	99		
Staphylococcus warneri		2.63	0.27	MAN 10	1	8		
Staphylococcus saprophyticus subsp. saprophyticus		1.24	0.00		99	7		
720752	**Good identification**							**720752**
					NOV	PYR		
Staphylococcus saprophyticus subsp. saprophyticus		93.43	0.56	ARG 8	99	7		
Staphylococcus warneri		5.75	0.52	GAL 8	1	8		
Staphylococcus petrasii subsp. petrasii		0.77	0.20		1	99		
720753	**Low discrimination**							**720753**
					COA	NOV	PYR	
Staphylococcus petrasii subsp. petrasii		81.40	0.39	NIT 90	1	1	99	
Staphylococcus saprophyticus subsp. saprophyticus		11.07	0.15	MAN 10	1	99	7	
Staphylococcus aureus subsp. aureus		2.73	-.02		99	4	10	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
721640	**Low discrimination**						**721640**
					NOV	PYR	
Staphylococcus warneri		73.47	0.34	ESL 1	1	8	
Staphylococcus petrasii subsp. petrasii		18.84	0.07		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		7.70	-.07		99	7	
721641	**Low discrimination**						**721641**
					OXI		
Staphylococcus petrasii subsp. petrasii		74.65	0.27	ESL 1	1		
Kocuria kristinae		25.35	0.17	NIT 90	99		
721652	**Good identification**						**721652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		98.33	0.37	ARG 8	99		
Staphylococcus warneri		1.67	0.21	ESL 1	1		
722000	**Good identification**						**722000**
Kocuria varians		100.0	0.29	VPT 14			
722051	**Good identification**						**722051**
Staphylococcus capitis subsp. ureolyticus		100.0	0.36	BGA 12			
722053	**Very good identification**						**722053**
Staphylococcus capitis subsp. ureolyticus		99.89	0.54	BGA 12			
Staphylococcus simulans		0.11	0.06	SUC 96			
722100	**Good identification**						**722100**
					COA		
Kocuria varians		96.61	0.33	VPT 14	1		
Staphylococcus schleiferi subsp. coagulans		3.39	0.02	ARG 1	99		
722101	**Acceptable identification**						**722101**
					COA		
Staphylococcus schleiferi subsp. coagulans		94.61	0.43	PHS 99	99		
Staphylococcus schleiferi subsp. schleiferi		3.03	0.08	LAC 86	1		
Kocuria varians		2.37	0.11		1		
722103	**Very good identification**						**722103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	PHS 99			
722111	**Good identification**						**722111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	PHS 99			
722113	**Very good identification**						**722113**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	PHS 99	99		
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1		
722143	**Low discrimination**						**722143**
					COA	PYR	
Staphylococcus epidermidis		80.02	0.31	SUC 99	1	1	
Staphylococcus schleiferi subsp. coagulans		16.89	0.18		99	20	
Staphylococcus petrasii subsp. petrasii		3.09	-.04		1	99	
722211	**Good identification**						**722211**
Staphylococcus capitis subsp. ureolyticus		95.05	0.36	BGA 12			
Staphylococcus capitis subsp. capitis		4.53	0.18	MLT 96			
Staphylococcus simulans		0.42	0.01	LAC 88			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
722212	**Low discrimination**						**722212**
Staphylococcus simulans		63.95	0.31	VPT 4			
Staphylococcus capitis subsp. ureolyticus		36.05	0.13	TRE 95			
722213	**Good identification**						**722213**
					COA		
Staphylococcus capitis subsp. ureolyticus		94.25	0.54	BGA 12	1		
Staphylococcus simulans		5.65	0.42	MLT 96	1		
Staphylococcus schleiferi subsp. coagulans		0.10	0.07		99		
722240	**Low discrimination**						**722240**
					PYR		
Staphylococcus epidermidis		46.21	0.41		1		
Staphylococcus petrasii subsp. petrasii		18.41	0.27		99		
Staphylococcus warneri		17.81	0.41		8		
722241	**Low discrimination**						**722241**
					PYR		
Staphylococcus petrasii subsp. petrasii		44.11	0.47	TRE 99	99		
Staphylococcus capitis subsp. ureolyticus		32.86	0.36		1		
Staphylococcus epidermidis		22.85	0.46		1		
722242	**Low discrimination**						**722242**
					PYR		
Staphylococcus epidermidis		78.29	0.51		1		
Staphylococcus petrasii subsp. petrasii		5.30	0.21		99		
Staphylococcus hominis subsp. hominis		5.11	0.30		1		
722243	**Low discrimination**						**722243**
					PYR		
Staphylococcus capitis subsp. ureolyticus		71.15	0.54	BGA 12	1		
Staphylococcus epidermidis		21.36	0.57	MAN 96	1		
Staphylococcus petrasii subsp. petrasii		7.01	0.41		99		
722251	**Very good identification**						**722251**
					PYR		
Staphylococcus capitis subsp. ureolyticus		99.35	0.64	BGA 12	1		
Staphylococcus petrasii subsp. petrasii		0.62	0.27	LAC 88	99		
Staphylococcus epidermidis		0.03	0.05		1		
722252	**Acceptable identification**						**722252**
Staphylococcus capitis subsp. ureolyticus		92.30	0.41	BGA 12			
Staphylococcus simulans		3.07	0.24	MNS 99			
Staphylococcus warneri		1.81	0.29				
722253	**Very good identification**						**722253**
					PYR		
Staphylococcus capitis subsp. ureolyticus		99.83	0.82	BGA 12	1		
Staphylococcus simulans		0.11	0.35		70		
Staphylococcus petrasii subsp. petrasii		0.05	0.21		99		
722273	**Good identification**						**722273**
Staphylococcus capitis subsp. ureolyticus		100.0	0.41	BGA 12			
722301	**Low discrimination**						**722301**
					COA	PYR	
Staphylococcus schleiferi subsp. coagulans		73.49	0.32	PHS 99	99	20	
Staphylococcus epidermidis		21.61	0.21	LAC 86	1	1	
Staphylococcus petrasii subsp. petrasii		4.91	0.02		1	99	

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
722303	**Low discrimination**						**722303**
					COA		
Staphylococcus schleiferi subsp. coagulans		85.55	0.49	PHS 99	99		
Staphylococcus epidermidis		12.97	0.31		1		
Staphylococcus simulans		0.98	0.07		1		
722311	**Acceptable identification**						**722311**
					COA		
Staphylococcus schleiferi subsp. coagulans		92.52	0.32	PHS 99	99		
Staphylococcus capitis subsp. ureolyticus		7.48	-.05	LAC 86	1		
722313	**Low discrimination**						**722313**
					COA		
Staphylococcus schleiferi subsp. coagulans		86.77	0.48	PHS 99	99		
Staphylococcus capitis subsp. ureolyticus		8.37	0.13		1		
Staphylococcus simulans		4.33	0.20		1		
722340	**Low discrimination**						**722340**
					PYR		
Staphylococcus epidermidis		86.44	0.56		1		
Staphylococcus petrasii subsp. petrasii		4.05	0.23		99		
Staphylococcus hominis subsp. hominis		2.91	0.29		1		
722341	**Low discrimination**						**722341**
					PYR	NOV	
Staphylococcus epidermidis		81.03	0.62		1	1	
Staphylococcus petrasii subsp. petrasii		18.40	0.43		99	1	
Staphylococcus hominis subsp. novobiosepticus		0.36	0.05		1	99	
722342	**Very good identification**						**722342**
					PYR		
Staphylococcus epidermidis		96.91	0.67		1		
Staphylococcus hominis subsp. hominis		1.21	0.31		1		
Staphylococcus petrasii subsp. petrasii		0.77	0.18		99		
722343	**Very good identification**						**722343**
					PYR		
Staphylococcus epidermidis		95.57	0.72		1		
Staphylococcus petrasii subsp. petrasii		3.69	0.37		99		
Staphylococcus capitis subsp. ureolyticus		0.57	0.13		1		
722350	**Low discrimination**						**722350**
					PYR		
Staphylococcus warneri		30.17	0.28	TRE 95	8		
Staphylococcus haemolyticus		23.39	0.19		98		
Staphylococcus epidermidis		19.59	0.15		1		
722353	**Low discrimination**						**722353**
					PYR		
Staphylococcus capitis subsp. ureolyticus		88.31	0.41	BGA 12	1		
Staphylococcus epidermidis		6.26	0.31	GAL 1	1		
Staphylococcus petrasii subsp. petrasii		2.66	0.18		99		
722363	**Good identification**						**722363**
					PYR		
Staphylococcus epidermidis		96.28	0.31	XYL 1	1		
Staphylococcus petrasii subsp. petrasii		3.72	-.04		99		
722403	**Low discrimination**						**722403**
Staphylococcus simulans		88.07	0.27	VPT 4			
Staphylococcus petrasii subsp. petrasii		7.39	0.00	SUC 96			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus carnosus subsp. carnosus		4.55	-.02				
722412	**Good identification**						**722412**
Staphylococcus simulans		100.0	0.29	VPT 4			
722413	**Acceptable identification**						**722413**
Staphylococcus simulans		92.03	0.40	VPT 4			
Staphylococcus carnosus subsp. carnosus		4.75	0.10	SUC 96			
Staphylococcus felis		3.22	0.01				
722440	**Low discrimination**						**722440**
					PYR		
Staphylococcus petrasii subsp. petrasii		41.84	0.27	SUC 99	99		
Staphylococcus warneri		40.49	0.41	MNS 90	8		
Staphylococcus auricularis		16.21	0.25		40		
722441	**Good identification**						**722441**
Staphylococcus petrasii subsp. petrasii		99.52	0.47	SUC 99			
Staphylococcus auricularis		0.48	0.05				
722443	**Acceptable identification**						**722443**
Staphylococcus petrasii subsp. petrasii		94.87	0.41	SUC 99			
Staphylococcus simulans		5.13	0.20				
722450	**Low discrimination**						**722450**
					NOV	PYR	
Staphylococcus warneri		56.64	0.38	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		25.66	0.22		99	99	
Staphylococcus petrasii subsp. petrasii		8.98	0.07		1	99	
722451	**Low discrimination**						**722451**
					PYR	NOV	
Staphylococcus petrasii subsp. petrasii		87.18	0.27	SUC 99	99	1	
Staphylococcus saprophyticus subsp. bovis		6.92	0.09	MAN 10	99	99	
Staphylococcus capitis subsp. ureolyticus		5.90	-.05		1	1	
722453	**Low discrimination**						**722453**
					PYR		
Staphylococcus simulans		50.34	0.32	VPT 4	70		
Staphylococcus petrasii subsp. petrasii		25.84	0.21	SUC 96	99		
Staphylococcus capitis subsp. ureolyticus		23.83	0.13		1		
722541	**Good identification**						**722541**
Staphylococcus petrasii subsp. petrasii		100.0	0.43	SUC 99			
722543	**Good identification**						**722543**
Staphylococcus petrasii subsp. petrasii		98.90	0.37	SUC 99			
Staphylococcus caprae		1.10	-.09				
722550	**Low discrimination**						**722550**
					NOV	PYR	
Staphylococcus saprophyticus subsp. bovis		81.13	0.34	ARG 1	99	99	
Staphylococcus warneri		14.14	0.28		1	8	
Staphylococcus petrasii subsp. petrasii		4.73	0.04		1	99	
722600	**Low discrimination**						**722600**
					PYR		
Staphylococcus petrasii subsp. petrasii		45.50	0.27	MLT 99	99		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		44.03	0.41	MNS 90	8	
Staphylococcus auricularis		6.97	0.17		40	
722601	**Good identification**					**722601**
Staphylococcus petrasii subsp. petrasii		98.45	0.47	MLT 99		
Staphylococcus simulans		1.55	0.15			
722602	**Low discrimination**					**722602**
					PYR	
Staphylococcus simulans		82.94	0.45	VPT 4	70	
Staphylococcus petrasii subsp. petrasii		10.67	0.21		99	
Staphylococcus warneri		6.39	0.32		8	
722603	**Low discrimination**					**722603**
Staphylococcus simulans		74.20	0.56	VPT 4		
Staphylococcus petrasii subsp. petrasii		25.67	0.41			
Staphylococcus felis		0.13	-.10			
722610	**Low discrimination**					**722610**
					PYR	
Staphylococcus warneri		67.37	0.38	MLT 95	8	
Staphylococcus simulans		16.26	0.16		70	
Staphylococcus petrasii subsp. petrasii		10.68	0.07		99	
722611	**Low discrimination**					**722611**
					PYR	
Staphylococcus petrasii subsp. petrasii		61.21	0.27	MAN 10	99	
Staphylococcus simulans		34.65	0.27	MLT 99	70	
Staphylococcus capitis subsp. ureolyticus		4.15	-.05		1	
722612	**Very good identification**					**722612**
					NOV	
Staphylococcus simulans		97.92	0.57	VPT 4	1	
Staphylococcus warneri		1.37	0.29		1	
Staphylococcus saprophyticus subsp. saprophyticus		0.37	-.04		99	
722613	**Very good identification**					**722613**
					PYR	
Staphylococcus simulans		97.90	0.68	VPT 4	70	
Staphylococcus petrasii subsp. petrasii		0.94	0.21		99	
Staphylococcus capitis subsp. ureolyticus		0.87	0.13		1	
722633	**Good identification**					**722633**
Staphylococcus simulans		100.0	0.27	VPT 4		
722640	**Low discrimination**					**722640**
					PYR	
Staphylococcus petrasii subsp. petrasii		79.30	0.68	MNS 90	99	
Staphylococcus warneri		14.73	0.68		8	
Staphylococcus haemolyticus		2.76	0.46		98	
722641	**Very good identification**					**722641**
					PYR	
Staphylococcus petrasii subsp. petrasii		99.90	0.88		99	
Staphylococcus warneri		0.04	0.33		8	
Staphylococcus haemolyticus		0.01	0.14		98	
722642	**Low discrimination**					**722642**
					PYR	
Staphylococcus petrasii subsp. petrasii		81.47	0.62	MNS 90	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		9.37	0.58		8		
Staphylococcus haemolyticus		2.96	0.40		98		
722643	**Very good identification**						**722643**
					PYR		
Staphylococcus petrasii subsp. petrasii		98.44	0.82		99		
Staphylococcus simulans		1.29	0.48		70		
Staphylococcus capitis subsp. ureolyticus		0.10	0.13		1		
722650	**Low discrimination**						**722650**
					PYR		
Staphylococcus warneri		45.37	0.65		8		
Staphylococcus petrasii subsp. petrasii		37.49	0.48		99		
Staphylococcus haemolyticus		6.89	0.41		98		
722651	**Very good identification**						**722651**
					PYR	NOV	
Staphylococcus petrasii subsp. petrasii		97.51	0.68	MAN 10	99	1	
Staphylococcus capitis subsp. ureolyticus		1.60	0.23		1	1	
Staphylococcus saprophyticus subsp. bovis		0.31	0.22		99	99	
722652	**Low discrimination**						**722652**
					NOV	PYR	
Staphylococcus simulans		28.90	0.50	VPT 4	1	70	
Staphylococcus saprophyticus subsp. saprophyticus		23.73	0.37		99	7	
Staphylococcus petrasii subsp. petrasii		22.76	0.43		1	99	
722653	**Low discrimination**						**722653**
					PYR		
Staphylococcus petrasii subsp. petrasii		58.62	0.62	MAN 10	99		
Staphylococcus simulans		27.69	0.61		70		
Staphylococcus capitis subsp. ureolyticus		13.11	0.41		1		
722660	**Low discrimination**						**722660**
					PYR		
Staphylococcus petrasii subsp. petrasii		81.93	0.27	XYL 1	99		
Staphylococcus warneri		15.22	0.27	MNS 90	8		
Staphylococcus haemolyticus		2.85	0.04		98		
722661	**Good identification**						**722661**
Staphylococcus petrasii subsp. petrasii		100.0	0.47	XYL 1			
722663	**Good identification**						**722663**
Staphylococcus petrasii subsp. petrasii		98.71	0.41	XYL 1			
Staphylococcus simulans		1.30	0.07				
722671	**Good identification**						**722671**
Staphylococcus petrasii subsp. petrasii		100.0	0.27	MAN 10			
722701	**Good identification**						**722701**
					COA		
Staphylococcus petrasii subsp. petrasii		99.58	0.43	MLT 99	1		
Staphylococcus intermedius		0.42	-.03		99		
722703	**Low discrimination**						**722703**
					COA		
Staphylococcus petrasii subsp. petrasii		65.07	0.37	MLT 99	1		
Staphylococcus simulans		24.54	0.34		1		
Staphylococcus intermedius		3.11	0.14		99		
722710	**Low discrimination**						**722710**

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
					NOV	PYR		
Staphylococcus warneri		63.83	0.28	MLT 95	1	8		
Staphylococcus petrasii subsp. petrasii		21.37	0.04		1	99		
Staphylococcus saprophyticus subsp. bovis		14.80	0.05		99	99		
722712	**Acceptable identification**							**722712**
					PYR			
Staphylococcus simulans		92.76	0.35	VPT 4	70			
Staphylococcus warneri		4.70	0.18	GAL 8	8			
Staphylococcus petrasii subsp. petrasii		2.54	-.02		99			
722713	**Low discrimination**							**722713**
Staphylococcus simulans		66.42	0.46	VPT 4				
Staphylococcus condimenti		12.43	0.18	GAL 8				
Staphylococcus felis		7.57	0.18					
722740	**Low discrimination**							**722740**
					PYR			
Staphylococcus petrasii subsp. petrasii		85.23	0.64	MNS 90	99			
Staphylococcus warneri		7.50	0.57		8			
Staphylococcus haemolyticus		5.01	0.47		98			
722741	**Very good identification**							**722741**
					PYR			
Staphylococcus petrasii subsp. petrasii		99.89	0.84		99			
Staphylococcus epidermidis		0.05	0.21		1			
Staphylococcus warneri		0.02	0.23		8			
722742	**Low discrimination**							**722742**
					PYR			
Staphylococcus petrasii subsp. petrasii		84.51	0.59	MNS 90	99			
Staphylococcus haemolyticus		5.19	0.41		98			
Staphylococcus warneri		4.60	0.48		8			
722743	**Very good identification**							**722743**
					PYR			
Staphylococcus petrasii subsp. petrasii		99.24	0.78		99			
Staphylococcus epidermidis		0.26	0.31		1			
Staphylococcus simulans		0.17	0.27		70			
722750	**Low discrimination**							**722750**
					NOV	PYR		
Staphylococcus petrasii subsp. petrasii		38.16	0.45	MAN 10	1	99		
Staphylococcus saprophyticus subsp. bovis		26.43	0.47	MNS 90	99	99		
Staphylococcus warneri		21.88	0.54		1	8		
722751	**Very good identification**							**722751**
					COA	NOV	PYR	
Staphylococcus petrasii subsp. petrasii		97.55	0.64	MAN 10	1	1	99	
Staphylococcus saprophyticus subsp. bovis		1.88	0.34		1	99	99	
Staphylococcus aureus subsp. aureus		0.20	-.02		99	4	10	
722752	**Low discrimination**							**722752**
					PYR			
Staphylococcus petrasii subsp. petrasii		41.45	0.39	MAN 10	99			
Staphylococcus warneri		14.71	0.45	MNS 90	8			
Staphylococcus haemolyticus		13.46	0.37		98			
722753	**Low discrimination**							**722753**
					COA	PYR		
Staphylococcus petrasii subsp. petrasii		68.03	0.59	MAN 10	1	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus aureus subsp. aureus		25.06	0.39		99	10	
Staphylococcus simulans		4.19	0.39		1	70	
722761	**Good identification**						**722761**
Staphylococcus petrasii subsp. petrasii		100.0	0.43	XYL 1			
722763	**Good identification**						**722763**
Staphylococcus petrasii subsp. petrasii		100.0	0.37	XYL 1			
723101	**Good identification**						**723101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	PHS 99			
723103	**Very good identification**						**723103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	PHS 99			
723111	**Good identification**						**723111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	PHS 99			
723113	**Very good identification**						**723113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	PHS 99			
723253	**Good identification**						**723253**
Staphylococcus capitis subsp. ureolyticus		100.0	0.41	BGA 12			
723301	**Good identification**						**723301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	PHS 99			
723303	**Good identification**						**723303**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	PHS 99			
723311	**Good identification**						**723311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	PHS 99			
723313	**Good identification**						**723313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	PHS 99			
723343	**Acceptable identification**						**723343**
					COA	PYR	
Staphylococcus epidermidis		90.69	0.31	ESL 1	1	1	
Staphylococcus schleiferi subsp. coagulans		5.81	0.07		99	20	
Staphylococcus petrasii subsp. petrasii		3.50	-.04		1	99	
723613	**Good identification**						**723613**
Staphylococcus simulans		100.0	0.27	VPT 4			
723640	**Low discrimination**						**723640**
					PYR		
Staphylococcus petrasii subsp. petrasii		80.18	0.27	ESL 1	99		
Staphylococcus warneri		14.89	0.27	MNS 90	8		
Staphylococcus haemolyticus		2.79	0.04		98		
723641	**Good identification**						**723641**
					OXI		
Staphylococcus petrasii subsp. petrasii		99.72	0.47	ESL 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Kocuria kristinae		0.28	-.06		99		
723643	**Good identification**						**723643**
Staphylococcus petrasii subsp. petrasii		98.71	0.41	ESL 1			
Staphylococcus simulans		1.30	0.07				
723651	**Good identification**						**723651**
Staphylococcus petrasii subsp. petrasii		100.0	0.27	ESL 1			
723741	**Good identification**						**723741**
Staphylococcus petrasii subsp. petrasii		97.69	0.43	ESL 1			
Rothia mucilaginosa		2.31	0.05				
723743	**Good identification**						**723743**
Staphylococcus petrasii subsp. petrasii		100.0	0.37	ESL 1			
724001	**Good identification**						**724001**
					COA		
Staphylococcus schleiferi subsp. schleiferi		97.21	0.30	URE 1	1		
Staphylococcus schleiferi subsp. coagulans		2.79	0.02	NIT 90	99		
724101	**Low discrimination**						**724101**
					COA		
Staphylococcus schleiferi subsp. coagulans		73.98	0.43	NIT 99	99		
Staphylococcus schleiferi subsp. schleiferi		26.02	0.30	LAC 86	1		
724103	**Very good identification**						**724103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	NIT 99			
724111	**Good identification**						**724111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	NIT 99			
724113	**Very good identification**						**724113**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	NIT 99	99		
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1		
724241	**Good identification**						**724241**
Staphylococcus epidermidis		100.0	0.31	NIT 97			
724242	**Good identification**						**724242**
Staphylococcus epidermidis		100.0	0.36	NIT 97			
724243	**Good identification**						**724243**
Staphylococcus epidermidis		99.00	0.41	NIT 97			
Staphylococcus capitis subsp. ureolyticus		1.00	-.14				
724301	**Good identification**						**724301**
					COA		
Staphylococcus schleiferi subsp. coagulans		95.10	0.32	NIT 99	99		
Staphylococcus epidermidis		4.90	0.05	LAC 86	1		
724303	**Good identification**						**724303**
					COA	PYR	
Staphylococcus schleiferi subsp. coagulans		96.93	0.49	NIT 99	99	20	
Staphylococcus epidermidis		2.58	0.16		1	1	
Staphylococcus intermedius		0.49	-.02		99	90	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
724311	**Good identification**						**724311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	NIT 99			
724313	**Good identification**						**724313**
Staphylococcus schleiferi subsp. coagulans		98.54	0.48	NIT 99			
Staphylococcus intermedius		1.47	0.07				
724340	**Good identification**						**724340**
Staphylococcus epidermidis		100.0	0.41	NIT 97			
724341	**Good identification**						**724341**
Staphylococcus epidermidis		100.0	0.46	NIT 97			
724342	**Good identification**						**724342**
Staphylococcus epidermidis		100.0	0.51	NIT 97			
724343	**Very good identification**						**724343**
					COA	PYR	
Staphylococcus epidermidis		99.33	0.57	NIT 97	1	1	
Staphylococcus schleiferi subsp. coagulans		0.38	0.08		99	20	
Staphylococcus intermedius		0.29	0.02		99	90	
724401	**Good identification**						**724401**
Staphylococcus schleiferi subsp. schleiferi		100.0	0.39	URE 1			
724501	**Good identification**						**724501**
					COA		
Staphylococcus schleiferi subsp. schleiferi		99.05	0.39	URE 1	1		
Staphylococcus schleiferi subsp. coagulans		0.95	0.02	NIT 90	99		
724550	**Low discrimination**						**724550**
					NOV	YEP	
Staphylococcus kloosii		74.18	0.33	VPT 10	99	99	
Staphylococcus caprae		25.82	-.05	ARG 1	1	1	
724553	**Good identification**						**724553**
					COA	NOV	
Staphylococcus caprae		97.91	0.39	BGA 1	1	1	
Staphylococcus aureus subsp. aureus		1.73	-.02	NIT 90	99	4	
Staphylococcus cohnii subsp. urealyticum		0.36	0.06		1	99	
724613	**Low discrimination**						**724613**
Staphylococcus simulans		82.43	0.30	VPT 4			
Staphylococcus felis		17.57	0.07	NIT 96			
724640	**Low discrimination**						**724640**
					NOV	PYR	
Staphylococcus warneri		73.47	0.34	PHS 1	1	8	
Staphylococcus petrasii subsp. petrasii		18.84	0.07		1	99	
Staphylococcus saprophyticus subsp. saprophyticus		7.70	-.07		99	7	
724641	**Good identification**						**724641**
Staphylococcus petrasii subsp. petrasii		100.0	0.27	NIT 90			
724652	**Good identification**						**724652**
					NOV		
Staphylococcus saprophyticus subsp. saprophyticus		96.73	0.37	ARG 8	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		1.64	0.21	PHS 1	1		
Staphylococcus simulans		1.64	0.12		1		
724713	**Low discrimination**						**724713**
					COA		
Staphylococcus intermedius		36.30	0.30	VPT 1	99		
Staphylococcus condimenti		32.58	0.18	NIT 93	1		
Staphylococcus felis		19.84	0.18		1		
724753	**Low discrimination**						**724753**
					COA	PYR	
Staphylococcus aureus subsp. aureus		74.82	0.39	BGA 10	99	10	
Staphylococcus caprae		15.87	0.30	NIT 99	1	75	
Staphylococcus intermedius		8.71	0.34		99	90	
725101	**Good identification**						**725101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	NIT 99			
725103	**Very good identification**						**725103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	NIT 99			
725111	**Good identification**						**725111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	NIT 99			
725113	**Very good identification**						**725113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	NIT 99			
725301	**Good identification**						**725301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	NIT 99			
725303	**Good identification**						**725303**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	NIT 99			
725311	**Good identification**						**725311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	NIT 99			
725313	**Good identification**						**725313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	NIT 99			
725450	**Good identification**						**725450**
Staphylococcus kloosii		100.0	0.27	VPT 10			
725452	**Good identification**						**725452**
Staphylococcus kloosii		100.0	0.28	VPT 10			
725550	**Good identification**						**725550**
Staphylococcus kloosii		100.0	0.33	VPT 10			
725552	**Good identification**						**725552**
Staphylococcus kloosii		100.0	0.33	VPT 10			
726001	**Low discrimination**						**726001**
					COA		
Staphylococcus schleiferi subsp. schleiferi		66.34	0.49	URE 1	1		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus schleiferi subsp. coagulans		20.96	0.43		99	
Staphylococcus carnosus subsp. carnosus		12.69	0.34		1	
726003	**Low discrimination**					**726003**
					COA	
Staphylococcus schleiferi subsp. coagulans		71.45	0.59	GAL 99	99	
Staphylococcus carnosus subsp. carnosus		28.18	0.47		1	
Staphylococcus schleiferi subsp. schleiferi		0.37	0.08		1	
726011	**Low discrimination**					**726011**
					COA	
Staphylococcus carnosus subsp. carnosus		71.72	0.47	URE 1	1	
Staphylococcus schleiferi subsp. coagulans		27.33	0.42		99	
Staphylococcus schleiferi subsp. schleiferi		0.95	0.08		1	
726013	**Low discrimination**					**726013**
					COA	
Staphylococcus carnosus subsp. carnosus		62.91	0.59	URE 1	1	
Staphylococcus schleiferi subsp. coagulans		36.81	0.58		99	
Staphylococcus felis		0.19	0.01		1	
726043	**Low discrimination**					**726043**
					COA	
Staphylococcus epidermidis		77.26	0.31	SUC 99	1	
Staphylococcus schleiferi subsp. coagulans		16.31	0.18		99	
Staphylococcus carnosus subsp. carnosus		6.43	0.05		1	
726100	**Good identification**					**726100**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.90	0.43	MNS 99	99	
Staphylococcus schleiferi subsp. schleiferi		3.10	0.08	LAC 86	1	
726101	**Very good identification**					**726101**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.76	0.84	LAC 86	99	
Staphylococcus schleiferi subsp. schleiferi		3.09	0.49		1	
Staphylococcus carnosus subsp. carnosus		0.15	0.22		1	
726102	**Very good identification**					**726102**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.85	0.59	MNS 99	99	
Staphylococcus epidermidis		0.15	0.00		1	
726103	**Excellent identification**					**726103**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.89	1.00		99	
Staphylococcus carnosus subsp. carnosus		0.10	0.34		1	
Staphylococcus schleiferi subsp. schleiferi		0.01	0.08		1	
726110	**Good identification**					**726110**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	MNS 99		
726111	**Very good identification**					**726111**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.31	0.83	LAC 86	99	
Staphylococcus carnosus subsp. carnosus		0.66	0.34		1	
Staphylococcus schleiferi subsp. schleiferi		0.04	0.08		1	
726112	**Very good identification**					**726112**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	MNS 99	99	
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1	

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
726113	**Excellent identification**					**726113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.51	0.99		99	
Staphylococcus carnosus subsp. carnosus		0.43	0.47		1	
Staphylococcus condimenti		0.04	0.14		1	
726121	**Good identification**					**726121**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.90	0.43	XYL 1	99	
Staphylococcus schleiferi subsp. schleiferi		3.10	0.08	LAC 86	1	
726123	**Very good identification**					**726123**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	XYL 1		
726131	**Good identification**					**726131**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	XYL 1		
726133	**Very good identification**					**726133**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	XYL 1	99	
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1	
726140	**Good identification**					**726140**
					COA	
Staphylococcus epidermidis		96.56	0.31	SUC 99	1	
Staphylococcus schleiferi subsp. coagulans		3.44	0.02		99	
726141	**Low discrimination**					**726141**
					COA	
Staphylococcus schleiferi subsp. coagulans		64.18	0.43	MLT 1	99	
Staphylococcus epidermidis		33.77	0.36	LAC 86	1	
Staphylococcus schleiferi subsp. schleiferi		2.05	0.08		1	
726142	**Acceptable identification**					**726142**
					COA	
Staphylococcus epidermidis		91.74	0.41	SUC 99	1	
Staphylococcus schleiferi subsp. coagulans		6.35	0.18		99	
Staphylococcus caprae		1.92	-.05		1	
726143	**Low discrimination**					**726143**
					COA	
Staphylococcus schleiferi subsp. coagulans		77.30	0.59	MLT 1	99	
Staphylococcus epidermidis		20.97	0.47		1	
Staphylococcus caprae		1.73	0.13		1	
726151	**Low discrimination**					**726151**
					COA	
Staphylococcus schleiferi subsp. coagulans		87.04	0.42	MLT 1	99	
Staphylococcus caprae		12.96	0.13	LAC 86	1	
726153	**Low discrimination**					**726153**
					COA	
Staphylococcus schleiferi subsp. coagulans		67.75	0.58	MLT 1	99	
Staphylococcus caprae		31.21	0.39		1	
Staphylococcus aureus subsp. aureus		0.55	-.02		99	
726201	**Low discrimination**					**726201**
					COA	
Staphylococcus schleiferi subsp. coagulans		71.67	0.32	GAL 99	99	
Staphylococcus epidermidis		21.07	0.21	LAC 86	1	
Staphylococcus schleiferi subsp. schleiferi		7.26	0.08		1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
726203	**Low discrimination**						**726203**
					COA		
Staphylococcus schleiferi subsp. coagulans		81.01	0.49	GAL 99	99		
Staphylococcus epidermidis		12.28	0.31		1		
Staphylococcus simulans		3.57	0.19		1		
726211	**Low discrimination**						**726211**
					COA		
Staphylococcus schleiferi subsp. coagulans		82.51	0.32	GAL 99	99		
Staphylococcus carnosus subsp. carnosus		6.93	0.05	LAC 86	1		
Staphylococcus capitis subsp. ureolyticus		6.67	-.05		1		
726213	**Low discrimination**						**726213**
					COA		
Staphylococcus schleiferi subsp. coagulans		72.41	0.48	GAL 99	99		
Staphylococcus simulans		13.83	0.32		1		
Staphylococcus capitis subsp. ureolyticus		6.99	0.13		1		
726240	**Very good identification**						**726240**
Staphylococcus epidermidis		100.0	0.56				
726241	**Very good identification**						**726241**
					PYR		
Staphylococcus epidermidis		99.40	0.62		1		
Staphylococcus petrasii subsp. petrasii		0.34	0.05		99		
Staphylococcus capitis subsp. ureolyticus		0.26	-.05		1		
726242	**Very good identification**						**726242**
Staphylococcus epidermidis		99.93	0.67				
Staphylococcus simulans		0.07	0.02				
726243	**Very good identification**						**726243**
					YEP		
Staphylococcus epidermidis		99.01	0.72		1		
Staphylococcus capitis subsp. ureolyticus		0.59	0.13		77		
Staphylococcus chromogenes		0.14	0.10		95		
726253	**Low discrimination**						**726253**
					COA		
Staphylococcus capitis subsp. ureolyticus		83.08	0.41	BGA 12	1		
Staphylococcus epidermidis		5.89	0.31	PHS 1	1		
Staphylococcus delphini		5.30	0.18		99		
726263	**Good identification**						**726263**
Staphylococcus epidermidis		100.0	0.31	XYL 1			
726300	**Low discrimination**						**726300**
					COA		
Staphylococcus schleiferi subsp. coagulans		52.71	0.32	MNS 99	99		
Staphylococcus epidermidis		47.29	0.31	LAC 86	1		
726301	**Very good identification**						**726301**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		98.05	0.74	LAC 86	99	1	
Staphylococcus epidermidis		1.65	0.36		1	1	
Staphylococcus chromogenes		0.13	0.10		1	95	
726302	**Low discrimination**						**726302**
					COA		
Staphylococcus schleiferi subsp. coagulans		68.38	0.49	MNS 99	99		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		31.62	0.41		1		
726303	**Very good identification**						**726303**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		98.85	0.90		99	1	
Staphylococcus epidermidis		0.86	0.47		1	1	
Staphylococcus chromogenes		0.20	0.30		1	95	
726310	**Good identification**						**726310**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	MNS 99			
726311	**Very good identification**						**726311**
					COA	YEP	
Staphylococcus schleiferi subsp. coagulans		99.76	0.73	LAC 86	99	1	
Staphylococcus intermedius		0.20	0.14		99	1	
Staphylococcus chromogenes		0.04	-.01		1	95	
726312	**Good identification**						**726312**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.01	0.48	MNS 99	99		
Staphylococcus epidermidis		0.50	0.00		1		
Staphylococcus simulans		0.49	-.01		1		
726313	**Very good identification**						**726313**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.38	0.89		99		
Staphylococcus intermedius		0.20	0.30		99		
Staphylococcus condimenti		0.18	0.18		1		
726321	**Good identification**						**726321**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	XYL 1			
726323	**Good identification**						**726323**
					COA		
Staphylococcus schleiferi subsp. coagulans		99.14	0.49	XYL 1	99		
Staphylococcus epidermidis		0.86	0.05		1		
726331	**Good identification**						**726331**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	XYL 1			
726333	**Good identification**						**726333**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	XYL 1			
726340	**Very good identification**						**726340**
Staphylococcus epidermidis		100.0	0.72				
726341	**Very good identification**						**726341**
					COA	YEP	
Staphylococcus epidermidis		99.22	0.77		1	1	
Staphylococcus schleiferi subsp. coagulans		0.60	0.32		99	1	
Staphylococcus chromogenes		0.08	0.10		1	95	
726342	**Very good identification**						**726342**
					COA		
Staphylococcus epidermidis		99.98	0.82		1		
Staphylococcus schleiferi subsp. coagulans		0.02	0.08		99		
726343	**Very good identification**						**726343**
					COA	YEP	
Staphylococcus epidermidis		98.43	0.88		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus schleiferi subsp. coagulans		1.16	0.49		99	1	
Staphylococcus chromogenes		0.23	0.30		1	95	
726350	**Good identification**						**726350**
Staphylococcus epidermidis		100.0	0.31	MAN 1			
726351	**Low discrimination**						**726351**
					COA	PYR	
Staphylococcus epidermidis		51.95	0.36	MAN 1	1	1	
Staphylococcus schleiferi subsp. coagulans		28.78	0.32		99	20	
Staphylococcus intermedius		8.53	0.18		99	90	
726352	**Low discrimination**						**726352**
					COA		
Staphylococcus epidermidis		82.08	0.41	MAN 1	1		
Staphylococcus caprae		12.06	0.13		1		
Staphylococcus aureus subsp. aureus		4.21	-.02		99		
726353	**Low discrimination**						**726353**
					COA		
Staphylococcus aureus subsp. aureus		45.85	0.39	BGA 10	99		
Staphylococcus schleiferi subsp. coagulans		18.02	0.48	TRE 99	99		
Staphylococcus epidermidis		16.77	0.47		1		
726360	**Good identification**						**726360**
Staphylococcus epidermidis		100.0	0.31	XYL 1			
726361	**Good identification**						**726361**
Staphylococcus epidermidis		100.0	0.36	XYL 1			
726362	**Good identification**						**726362**
Staphylococcus epidermidis		100.0	0.41	XYL 1			
726363	**Good identification**						**726363**
					COA		
Staphylococcus epidermidis		98.84	0.47	XYL 1	1		
Staphylococcus schleiferi subsp. coagulans		1.16	0.08		99		
726401	**Very good identification**						**726401**
					COA		
Staphylococcus schleiferi subsp. schleiferi		97.24	0.59	URE 1	1		
Staphylococcus carnosus subsp. carnosus		2.66	0.27		1		
Staphylococcus schleiferi subsp. coagulans		0.10	0.02		99		
726403	**Low discrimination**						**726403**
Staphylococcus carnosus subsp. carnosus		61.63	0.39	URE 1			
Staphylococcus felis		24.97	0.25				
Staphylococcus schleiferi subsp. schleiferi		5.69	0.18				
726411	**Low discrimination**						**726411**
Staphylococcus carnosus subsp. carnosus		88.59	0.39	URE 1			
Staphylococcus schleiferi subsp. schleiferi		8.18	0.18				
Staphylococcus felis		2.43	0.01				
726413	**Low discrimination**						**726413**
Staphylococcus carnosus subsp. carnosus		55.31	0.51	URE 1			
Staphylococcus felis		37.50	0.42				

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		3.61	0.30				
726501	**Low discrimination**						**726501**
					COA		
Staphylococcus schleiferi subsp. schleiferi		89.80	0.59	URE 1	1		
Staphylococcus schleiferi subsp. coagulans		9.46	0.43		99		
Staphylococcus carnosus subsp. carnosus		0.61	0.14		1		
726503	**Low discrimination**						**726503**
					COA		
Staphylococcus schleiferi subsp. coagulans		74.98	0.59	TRE 1	99		
Staphylococcus felis		17.19	0.36		1		
Staphylococcus carnosus subsp. carnosus		3.17	0.27		1		
726511	**Low discrimination**						**726511**
					COA		
Staphylococcus schleiferi subsp. coagulans		57.77	0.42	TRE 1	99		
Staphylococcus carnosus subsp. carnosus		16.24	0.27	LAC 86	1		
Staphylococcus condimenti		13.24	0.14		1		
726513	**Low discrimination**						**726513**
					COA		
Staphylococcus condimenti		55.94	0.55	VPT 1	1		
Staphylococcus felis		25.17	0.53		1		
Staphylococcus schleiferi subsp. coagulans		15.14	0.58		99		
726543	**Low discrimination**						**726543**
					COA		
Staphylococcus caprae		84.85	0.33	BGA 1	1		
Staphylococcus schleiferi subsp. coagulans		4.26	0.18	MAN 95	99		
Staphylococcus aureus subsp. aureus		3.16	-.02		99		
726551	**Acceptable identification**						**726551**
					COA	NOV	
Staphylococcus caprae		92.90	0.33	BGA 1	1	1	
Staphylococcus aureus subsp. aureus		3.46	-.02	LAC 95	99	4	
Staphylococcus saprophyticus subsp. bovis		2.07	0.09		1	99	
726552	**Good identification**						**726552**
					COA	NOV	
Staphylococcus caprae		97.76	0.41	BGA 1	1	1	
Staphylococcus aureus subsp. aureus		1.40	-.02	MNS 88	99	4	
Staphylococcus saprophyticus subsp. bovis		0.84	0.09		1	99	
726553	**Low discrimination**						**726553**
					COA		
Staphylococcus caprae		82.39	0.59	BGA 1	1		
Staphylococcus aureus subsp. aureus		15.97	0.39		99		
Staphylococcus condimenti		0.74	0.14		1		
726602	**Good identification**						**726602**
Staphylococcus simulans		98.25	0.35	VPT 4			
Staphylococcus felis		1.75	-.10				
726603	**Low discrimination**						**726603**
					YEP		
Staphylococcus simulans		53.07	0.46	VPT 4	1		
Staphylococcus felis		27.89	0.31		1		
Staphylococcus chromogenes		15.84	0.32		95		
726612	**Good identification**						**726612**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus simulans		97.11	0.47	VPT 4			
Staphylococcus felis		2.89	0.07				
726613	**Low discrimination**						**726613**
Staphylococcus simulans		50.50	0.58	VPT 4			
Staphylococcus felis		44.41	0.48				
Staphylococcus condimenti		2.47	0.18				
726640	**Low discrimination**						**726640**
					PYR		
Staphylococcus petrasii subsp. petrasii		71.05	0.27	PHS 1	99		
Staphylococcus warneri		13.20	0.27	MNS 90	8		
Staphylococcus epidermidis		10.21	0.15		1		
726641	**Good identification**						**726641**
					YEP	PYR	
Staphylococcus petrasii subsp. petrasii		95.52	0.47	PHS 1	1	99	
Staphylococcus epidermidis		2.83	0.21		1	1	
Staphylococcus chromogenes		1.64	0.12		95	45	
726642	**Low discrimination**						**726642**
					PYR		
Staphylococcus simulans		36.17	0.28	VPT 4	70		
Staphylococcus petrasii subsp. petrasii		31.07	0.21		99		
Staphylococcus epidermidis		26.26	0.26		1		
726643	**Low discrimination**						**726643**
					YEP		
Staphylococcus petrasii subsp. petrasii		50.74	0.41	PHS 1	1		
Staphylococcus simulans		21.97	0.39		1		
Staphylococcus chromogenes		14.60	0.32		95		
726651	**Low discrimination**						**726651**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		66.30	0.27	PHS 1	1	99	
Staphylococcus aureus subsp. aureus		17.72	0.05	MAN 10	99	10	
Staphylococcus simulans		5.62	0.10		1	70	
726652	**Low discrimination**						**726652**
					COA	NOV	
Staphylococcus simulans		82.06	0.40	VPT 4	1	1	
Staphylococcus aureus subsp. aureus		8.75	0.05		99	4	
Staphylococcus saprophyticus subsp. bovis		2.60	0.09		1	99	
726653	**Low discrimination**						**726653**
					COA	PYR	
Staphylococcus aureus subsp. aureus		72.43	0.46	BGA 10	99	10	
Staphylococcus simulans		22.98	0.51	GAL 98	1	70	
Staphylococcus petrasii subsp. petrasii		1.47	0.21		1	99	
726701	**Low discrimination**						**726701**
					COA	YEP	
Staphylococcus chromogenes		38.41	0.32	VPT 1	1	95	
Staphylococcus schleiferi subsp. coagulans		25.01	0.32	LAC 90	99	1	
Staphylococcus intermedius		22.40	0.28		99	1	
726703	**Low discrimination**						**726703**
					COA	YEP	
Staphylococcus chromogenes		36.94	0.51	VPT 1	1	95	
Staphylococcus felis		24.20	0.42		1	1	
Staphylococcus schleiferi subsp. coagulans		16.42	0.49		99	1	

Code Taxon		ID score	T index	tests against	Differentiation tests			Code
726711	**Low discrimination**							**726711**
					COA			
Staphylococcus intermedius		43.71	0.37	VPT 1	99			
Staphylococcus condimenti		18.14	0.18	LAC 86	1			
Staphylococcus schleiferi subsp. coagulans		16.66	0.32		99			
726713	**Low discrimination**							**726713**
					COA			
Staphylococcus condimenti		49.49	0.59	VPT 1	1			
Staphylococcus felis		30.14	0.59		1			
Staphylococcus intermedius		7.40	0.54		99			
726740	**Low discrimination**							**726740**
					PYR			
Staphylococcus epidermidis		50.09	0.31	TRE 1	1			
Staphylococcus petrasii subsp. petrasii		41.01	0.23		99			
Staphylococcus warneri		3.61	0.16		8			
726741	**Low discrimination**							**726741**
					YEP	PYR		
Staphylococcus petrasii subsp. petrasii		58.10	0.43	PHS 1	1	99		
Staphylococcus epidermidis		14.64	0.36		1	1		
Staphylococcus chromogenes		13.50	0.32		95	45		
726742	**Low discrimination**							**726742**
					PYR			
Staphylococcus epidermidis		72.41	0.41	TRE 1	1			
Staphylococcus petrasii subsp. petrasii		10.08	0.18		99			
Staphylococcus caprae		5.04	0.06		75			
726743	**Low discrimination**							**726743**
					COA	PYR	YEP	
Staphylococcus aureus subsp. aureus		30.60	0.39	BGA 10	99	10	84	
Staphylococcus chromogenes		29.32	0.51	MAN 99	1	45	95	
Staphylococcus intermedius		17.51	0.48		99	90	1	
726750	**Low discrimination**							**726750**
					COA	NOV		
Staphylococcus saprophyticus subsp. bovis		73.35	0.34	ARG 1	1	99		
Staphylococcus caprae		10.36	0.06		1	1		
Staphylococcus aureus subsp. aureus		7.64	-.02		99	4		
726751	**Low discrimination**							**726751**
					COA	PYR		
Staphylococcus aureus subsp. aureus		68.61	0.39	BGA 10	99	10		
Staphylococcus intermedius		17.28	0.41	LAC 99	99	90		
Staphylococcus caprae		6.89	0.24		1	75		
726752	**Low discrimination**							**726752**
					COA	NOV		
Staphylococcus aureus subsp. aureus		75.50	0.39	BGA 10	99	4		
Staphylococcus caprae		19.65	0.32	MNS 99	1	1		
Staphylococcus saprophyticus subsp. bovis		1.83	0.22		1	99		
726753	**Very good identification**							**726753**
					COA	PYR		
Staphylococcus aureus subsp. aureus		96.18	0.80	BGA 10	99	10		
Staphylococcus caprae		1.85	0.50		1	75		
Staphylococcus intermedius		1.50	0.57		99	90		
726773	**Acceptable identification**							**726773**
					COA			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus aureus subsp. aureus		92.43	0.39	BGA 10	99	
Staphylococcus lutrae		3.58	0.18	XYL 1	99	
Staphylococcus caprae		1.78	0.09		1	
727001	**Acceptable identification**					**727001**
					COA	
Staphylococcus schleiferi subsp. coagulans		90.64	0.42	GAL 99	99	
Staphylococcus carnosus subsp. carnosus		6.35	0.15	LAC 86	1	
Staphylococcus schleiferi subsp. schleiferi		3.02	0.08		1	
727003	**Very good identification**					**727003**
					COA	
Staphylococcus schleiferi subsp. coagulans		95.64	0.59	GAL 99	99	
Staphylococcus carnosus subsp. carnosus		4.36	0.27		1	
727011	**Low discrimination**					**727011**
					COA	
Staphylococcus schleiferi subsp. coagulans		76.72	0.42	GAL 99	99	
Staphylococcus carnosus subsp. carnosus		23.28	0.27	LAC 86	1	
727013	**Low discrimination**					**727013**
					COA	
Staphylococcus schleiferi subsp. coagulans		83.50	0.58	GAL 99	99	
Staphylococcus carnosus subsp. carnosus		16.50	0.39		1	
727100	**Good identification**					**727100**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	MNS 99		
727101	**Very good identification**					**727101**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.95	0.83	LAC 86	99	
Staphylococcus schleiferi subsp. schleiferi		0.03	0.08		1	
Staphylococcus carnosus subsp. carnosus		0.02	0.02		1	
727102	**Very good identification**					**727102**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	MNS 99		
727103	**Excellent identification**					**727103**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.99	1.00		99	
Staphylococcus carnosus subsp. carnosus		0.01	0.15		1	
727110	**Good identification**					**727110**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	MNS 99		
727111	**Very good identification**					**727111**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.92	0.83	LAC 86	99	
Staphylococcus carnosus subsp. carnosus		0.08	0.15		1	
727112	**Very good identification**					**727112**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	MNS 99		
727113	**Excellent identification**					**727113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.95	0.99		99	
Staphylococcus carnosus subsp. carnosus		0.05	0.27		1	
727121	**Good identification**					**727121**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	XYL 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
727123	**Very good identification**					**727123**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	XYL 1		
727131	**Good identification**					**727131**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	XYL 1		
727133	**Very good identification**					**727133**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	XYL 1		
727141	**Good identification**					**727141**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	MLT 1		
727143	**Very good identification**					**727143**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.72	0.59	MLT 1	99	
Staphylococcus epidermidis		0.28	0.05		1	
727151	**Good identification**					**727151**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	MLT 1		
727153	**Very good identification**					**727153**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.52	0.58	MLT 1	99	
Staphylococcus caprae		0.48	-.02		1	
727201	**Good identification**					**727201**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	GAL 99		
727203	**Good identification**					**727203**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	GAL 99		
727211	**Good identification**					**727211**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	GAL 99		
727213	**Good identification**					**727213**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.37	0.48	GAL 99	99	
Staphylococcus carnosus subsp. carnosus		0.63	-.02		1	
727243	**Acceptable identification**					**727243**
					COA	
Staphylococcus epidermidis		93.98	0.31	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		6.02	0.07		99	
727300	**Good identification**					**727300**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	MNS 99		
727301	**Very good identification**					**727301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.73	LAC 86		
727302	**Good identification**					**727302**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.52	0.48	MNS 99	99	
Staphylococcus epidermidis		0.48	0.00		1	
727303	**Very good identification**					**727303**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					COA	
Staphylococcus schleiferi subsp. coagulans		99.99	0.89		99	
Staphylococcus epidermidis		0.01	0.05		1	
727310	**Good identification**					**727310**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	MNS 99		
727311	**Very good identification**					**727311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.72	LAC 86		
727312	**Good identification**					**727312**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	MNS 99		
727313	**Very good identification**					**727313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.89			
727321	**Good identification**					**727321**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	XYL 1		
727323	**Good identification**					**727323**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	XYL 1		
727331	**Good identification**					**727331**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	XYL 1		
727333	**Good identification**					**727333**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	XYL 1		
727340	**Good identification**					**727340**
Staphylococcus epidermidis		100.0	0.31	ESL 1		
727341	**Low discrimination**					**727341**
					COA	
Staphylococcus epidermidis		63.43	0.36	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		36.58	0.32		99	
727342	**Good identification**					**727342**
					COA	
Staphylococcus epidermidis		97.95	0.41	ESL 1	1	
Staphylococcus schleiferi subsp. coagulans		2.06	0.07		99	
727343	**Low discrimination**					**727343**
					COA	
Staphylococcus schleiferi subsp. coagulans		52.80	0.48	MLT 1	99	
Staphylococcus epidermidis		47.20	0.47		1	
727351	**Good identification**					**727351**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	MLT 1		
727353	**Good identification**					**727353**
					COA	
Staphylococcus schleiferi subsp. coagulans		95.95	0.48	MLT 1	99	
Staphylococcus aureus subsp. aureus		2.57	-.02		99	
Staphylococcus epidermidis		0.94	0.05		1	
727413	**Low discrimination**					**727413**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
					COA		
	Staphylococcus carnosus subsp. carnosus	84.68	0.32	URE 1	1		
	Staphylococcus schleiferi subsp. coagulans	10.10	0.17	ESL 10	99		
	Staphylococcus felis	5.22	0.01		1		
727501	**Acceptable identification**						**727501**
					COA		
	Staphylococcus schleiferi subsp. coagulans	90.92	0.42	TRE 1	99		
	Staphylococcus schleiferi subsp. schleiferi	9.08	0.18	LAC 86	1		
727503	**Very good identification**						**727503**
					COA		
	Staphylococcus schleiferi subsp. coagulans	99.28	0.59	TRE 1	99		
	Staphylococcus carnosus subsp. carnosus	0.49	0.07		1		
	Staphylococcus felis	0.24	-.06		1		
727511	**Good identification**						**727511**
					COA		
	Staphylococcus schleiferi subsp. coagulans	96.85	0.42	TRE 1	99		
	Staphylococcus carnosus subsp. carnosus	3.15	0.07	LAC 86	1		
727513	**Good identification**						**727513**
					COA		
	Staphylococcus schleiferi subsp. coagulans	92.81	0.58	TRE 1	99		
	Staphylococcus condimenti	3.61	0.14		1		
	Staphylococcus carnosus subsp. carnosus	1.97	0.19		1		
727701	**Good identification**						**727701**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.32	TRE 1			
727703	**Good identification**						**727703**
					COA	YEP	
	Staphylococcus schleiferi subsp. coagulans	95.37	0.48	TRE 1	99	1	
	Staphylococcus chromogenes	2.26	0.10		1	95	
	Staphylococcus felis	1.48	0.01		1	1	
727711	**Good identification**						**727711**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.31	TRE 1			
727713	**Low discrimination**						**727713**
					COA		
	Staphylococcus schleiferi subsp. coagulans	73.62	0.48	TRE 1	99		
	Staphylococcus condimenti	13.58	0.18		1		
	Staphylococcus felis	8.27	0.18		1		
727753	**Good identification**						**727753**
					COA	PYR	
	Staphylococcus aureus subsp. aureus	96.02	0.39	BGA 10	99	10	
	Staphylococcus caprae	1.85	0.09	ESL 1	1	75	
	Staphylococcus intermedius	1.50	0.16		99	90	
730640	**Low discrimination**						**730640**
					NOV	PYR	
	Staphylococcus warneri	73.47	0.34	ORN 1	1	8	
	Staphylococcus petrasii subsp. petrasii	18.84	0.07		1	99	
	Staphylococcus saprophyticus subsp. saprophyticus	7.70	-.07		99	7	
730641	**Good identification**						**730641**
	Staphylococcus petrasii subsp. petrasii	95.55	0.27	ORN 1			
	Staphylococcus lugdunensis	4.45	0.01	NIT 90			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
730652	**Good identification**					**730652**
					NOV	
Staphylococcus saprophyticus subsp. saprophyticus		98.33	0.37	ARG 8	99	
Staphylococcus warneri		1.67	0.21	ORN 1	1	
732241	**Low discrimination**					**732241**
					PYR	
Staphylococcus epidermidis		75.60	0.29	ORN 12	1	
Staphylococcus petrasii subsp. petrasii		10.81	0.05		99	
Staphylococcus capitis subsp. ureolyticus		8.05	-.05		1	
732242	**Good identification**					**732242**
Staphylococcus epidermidis		100.0	0.33	ORN 12		
732243	**Low discrimination**					**732243**
					PYR	
Staphylococcus epidermidis		67.61	0.39	ORN 12	1	
Staphylococcus capitis subsp. ureolyticus		16.68	0.13		1	
Staphylococcus lugdunensis		14.07	0.20		96	
732253	**Good identification**					**732253**
Staphylococcus capitis subsp. ureolyticus		100.0	0.41	ORN 1		
732340	**Good identification**					**732340**
Staphylococcus epidermidis		100.0	0.38	ORN 12		
732341	**Good identification**					**732341**
					PYR	
Staphylococcus epidermidis		97.11	0.44	ORN 12	1	
Staphylococcus petrasii subsp. petrasii		1.63	0.02		99	
Staphylococcus lugdunensis		1.25	0.01		96	
732342	**Good identification**					**732342**
Staphylococcus epidermidis		100.0	0.49	ORN 12		
732343	**Very good identification**					**732343**
					PYR	
Staphylococcus epidermidis		96.19	0.54	ORN 12	1	
Staphylococcus lugdunensis		3.53	0.20		96	
Staphylococcus petrasii subsp. petrasii		0.28	-.04		99	
732613	**Good identification**					**732613**
Staphylococcus simulans		100.0	0.27	VPT 4		
732640	**Low discrimination**					**732640**
					PYR	
Staphylococcus petrasii subsp. petrasii		78.93	0.27	ORN 1	99	
Staphylococcus warneri		14.66	0.27	MNS 90	8	
Staphylococcus lugdunensis		3.67	0.01		96	
732641	**Low discrimination**					**732641**
Staphylococcus petrasii subsp. petrasii		66.14	0.47	ORN 1		
Staphylococcus lugdunensis		33.86	0.42			
732643	**Low discrimination**					**732643**
Staphylococcus lugdunensis		89.41	0.61	ARG 5		
Staphylococcus petrasii subsp. petrasii		10.45	0.41			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		0.14	0.07			
732651	**Good identification**					**732651**
Staphylococcus petrasii subsp. petrasii		95.55	0.27	ORN 1		
Staphylococcus lugdunensis		4.45	0.01	MAN 10		
732741	**Low discrimination**					**732741**
					PYR	
Staphylococcus petrasii subsp. petrasii		56.37	0.43	ORN 1	99	
Staphylococcus lugdunensis		43.28	0.42		96	
Staphylococcus epidermidis		0.34	0.03		1	
732743	**Good identification**					**732743**
					PYR	
Staphylococcus lugdunensis		92.53	0.61	ARG 5	96	
Staphylococcus petrasii subsp. petrasii		7.21	0.37		99	
Staphylococcus epidermidis		0.26	0.13		1	
734341	**Good identification**					**734341**
Staphylococcus epidermidis		100.0	0.28	ORN 12		
734342	**Good identification**					**734342**
Staphylococcus epidermidis		100.0	0.33	ORN 12		
734343	**Good identification**					**734343**
Staphylococcus epidermidis		100.0	0.39	ORN 12		
736101	**Good identification**					**736101**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.90	0.43	ORN 1	99	
Staphylococcus schleiferi subsp. schleiferi		3.10	0.08	LAC 86	1	
736103	**Very good identification**					**736103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	ORN 1		
736111	**Good identification**					**736111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ORN 1		
736113	**Very good identification**					**736113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	ORN 1	99	
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1	
736143	**Low discrimination**					**736143**
					COA	
Staphylococcus epidermidis		78.55	0.29	ORN 12	1	
Staphylococcus schleiferi subsp. coagulans		21.45	0.18	SUC 99	99	
736240	**Good identification**					**736240**
Staphylococcus epidermidis		100.0	0.38	ORN 12		
736241	**Good identification**					**736241**
Staphylococcus epidermidis		100.0	0.44	ORN 12		
736242	**Good identification**					**736242**
Staphylococcus epidermidis		100.0	0.49	ORN 12		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
736243	**Very good identification**						**736243**
	Staphylococcus epidermidis	100.0	0.54	ORN 12			
736301	**Low discrimination**						**736301**
					COA		
	Staphylococcus schleiferi subsp. coagulans	81.49	0.32	ORN 1	99		
	Staphylococcus epidermidis	18.51	0.18	LAC 86	1		
736303	**Low discrimination**						**736303**
					COA		
	Staphylococcus schleiferi subsp. coagulans	89.52	0.49	ORN 1	99		
	Staphylococcus epidermidis	10.48	0.29		1		
736311	**Good identification**						**736311**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.32	ORN 1			
736313	**Good identification**						**736313**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.48	ORN 1			
736340	**Very good identification**						**736340**
	Staphylococcus epidermidis	100.0	0.54	ORN 12			
736341	**Very good identification**						**736341**
	Staphylococcus epidermidis	100.0	0.60	ORN 12			
736342	**Very good identification**						**736342**
	Staphylococcus epidermidis	100.0	0.64	ORN 12			
736343	**Very good identification**						**736343**
					COA		
	Staphylococcus epidermidis	99.91	0.70	ORN 12	1		
	Staphylococcus schleiferi subsp. coagulans	0.09	0.08		99		
736353	**Low discrimination**						**736353**
					COA		
	Staphylococcus epidermidis	75.46	0.29	ORN 12	1		
	Staphylococcus aureus subsp. aureus	15.29	-.02	MAN 1	99		
	Staphylococcus schleiferi subsp. coagulans	6.01	0.07		99		
736363	**Good identification**						**736363**
	Staphylococcus epidermidis	100.0	0.29	ORN 12			
736743	**Low discrimination**						**736743**
					COA	PYR	
	Staphylococcus epidermidis	45.37	0.29	ORN 12	1	1	
	Staphylococcus lugdunensis	29.10	0.20	TRE 1	1	96	
	Staphylococcus aureus subsp. aureus	9.19	-.02		99	10	
736753	**Good identification**						**736753**
					COA	PYR	
	Staphylococcus aureus subsp. aureus	96.37	0.39	ORN 1	99	10	
	Staphylococcus caprae	1.86	0.09	BGA 10	1	75	
	Staphylococcus intermedius	1.51	0.16		99	90	
737101	**Good identification**						**737101**
	Staphylococcus schleiferi subsp. coagulans	100.0	0.42	ORN 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
737103	**Very good identification**					**737103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	ORN 1		
737111	**Good identification**					**737111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	ORN 1		
737113	**Very good identification**					**737113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	ORN 1		
737301	**Good identification**					**737301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	ORN 1		
737303	**Good identification**					**737303**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	ORN 1		
737311	**Good identification**					**737311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	ORN 1		
737313	**Good identification**					**737313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	ORN 1		
737343	**Acceptable identification**					**737343**
					COA	
Staphylococcus epidermidis		92.35	0.29	ORN 12	1	
Staphylococcus schleiferi subsp. coagulans		7.65	0.07	ESL 1	99	
740040	**Low discrimination**					**740040**
					PYR	
Staphylococcus warneri		86.21	0.47	SUC 95	8	
Staphylococcus petrasii subsp. croceilyticus		13.79	0.05	TRE 95	90	
740042	**Good identification**					**740042**
Staphylococcus warneri		100.0	0.38	SUC 95		
740050	**Acceptable identification**					**740050**
Staphylococcus warneri		90.26	0.45	SUC 95		
Staphylococcus pasteuri		9.74	0.10	TRE 95		
740052	**Low discrimination**					**740052**
Staphylococcus warneri		88.32	0.35	SUC 95		
Staphylococcus pasteuri		11.68	0.02	TRE 95		
740140	**Good identification**					**740140**
Staphylococcus warneri		100.0	0.37	SUC 95		
740142	**Good identification**					**740142**
Staphylococcus warneri		100.0	0.27	SUC 95		
740150	**Good identification**					**740150**
Staphylococcus warneri		100.0	0.34	SUC 95		
740200	**Low discrimination**					**740200**
					PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		73.26	0.47	TRE 95	8		
Staphylococcus pasteuri		15.03	0.19	MLT 95	1		
Staphylococcus petrasii subsp. croceilyticus		11.72	0.05		90		
740202	**Low discrimination**						**740202**
Staphylococcus warneri		79.91	0.38	TRE 95			
Staphylococcus pasteuri		20.09	0.11	MLT 95			
740210	**Low discrimination**						**740210**
Staphylococcus pasteuri		63.50	0.35	TRE 99			
Staphylococcus warneri		36.50	0.45	MLT 86			
740212	**Low discrimination**						**740212**
Staphylococcus pasteuri		68.07	0.27	TRE 99			
Staphylococcus warneri		31.93	0.35	MLT 86			
740240	**Low discrimination**						**740240**
					PYR		
Staphylococcus warneri		52.64	0.74	TRE 95	8		
Staphylococcus petrasii subsp. croceilyticus		43.87	0.47		90		
Staphylococcus pasteuri		3.49	0.35		1		
740241	**Low discrimination**						**740241**
					PYR	YEP	
Staphylococcus warneri		66.55	0.39	TRE 95	8	43	
Staphylococcus petrasii subsp. croceilyticus		27.45	0.05	MNS 2	90	99	
Staphylococcus petrasii subsp. petrasii		3.65	-.02		99	1	
740242	**Good identification**						**740242**
					PYR		
Staphylococcus warneri		90.37	0.64	TRE 95	8		
Staphylococcus pasteuri		7.34	0.27		1		
Staphylococcus petrasii subsp. croceilyticus		2.28	0.05		90		
740243	**Low discrimination**						**740243**
Staphylococcus warneri		56.18	0.29	TRE 95			
Staphylococcus capitis subsp. ureolyticus		43.82	0.04	MNS 2			
740250	**Low discrimination**						**740250**
					PYR		
Staphylococcus warneri		63.53	0.71	TRE 95	8		
Staphylococcus pasteuri		35.74	0.51		1		
Staphylococcus petrasii subsp. croceilyticus		0.74	0.05		90		
740252	**Low discrimination**						**740252**
Staphylococcus warneri		59.01	0.61	TRE 95			
Staphylococcus pasteuri		40.68	0.43				
Staphylococcus capitis subsp. ureolyticus		0.31	-.09				
740253	**Good identification**						**740253**
Staphylococcus capitis subsp. ureolyticus		95.07	0.33	GLR 1			
Staphylococcus warneri		3.68	0.26	NIT 95			
Staphylococcus pasteuri		1.26	0.02				
740260	**Low discrimination**						**740260**
					PYR		
Staphylococcus warneri		54.55	0.33	TRE 95	8		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	XYL 1	90		
740270	**Low discrimination**						**740270**
Staphylococcus warneri		64.00	0.30	TRE 95			
Staphylococcus pasteuri		36.00	0.10	XYL 1			
740300	**Good identification**						**740300**
Staphylococcus warneri		100.0	0.37	TRE 95			
740302	**Good identification**						**740302**
Staphylococcus warneri		100.0	0.27	TRE 95			
740310	**Good identification**						**740310**
Staphylococcus warneri		100.0	0.34	TRE 95			
740340	**Very good identification**						**740340**
					NOV	PYR	
Staphylococcus warneri		96.98	0.63	TRE 95	1	8	
Staphylococcus petrasii subsp. croceilyticus		2.59	0.05		1	90	
Staphylococcus hominis subsp. novobiosepticus		0.44	-.02		99	1	
740341	**Low discrimination**						**740341**
					PYR		
Staphylococcus warneri		79.53	0.29	TRE 95	8		
Staphylococcus epidermidis		11.28	0.02	MNS 2	1		
Staphylococcus petrasii subsp. petrasii		9.20	-.05		99		
740342	**Very good identification**						**740342**
					NOV		
Staphylococcus warneri		97.84	0.54	TRE 95	1		
Staphylococcus epidermidis		1.45	0.07		1		
Staphylococcus hominis subsp. novobiosepticus		0.72	-.07		99		
740350	**Very good identification**						**740350**
Staphylococcus warneri		98.23	0.61	TRE 95			
Staphylococcus pasteuri		1.77	0.10				
740352	**Good identification**						**740352**
Staphylococcus warneri		97.84	0.51	TRE 95			
Staphylococcus pasteuri		2.16	0.02				
740400	**Low discrimination**						**740400**
					PYR		
Staphylococcus warneri		73.26	0.47	SUC 95	8		
Staphylococcus pasteuri		15.03	0.19	MLT 95	1		
Staphylococcus petrasii subsp. croceilyticus		11.72	0.05		90		
740402	**Low discrimination**						**740402**
Staphylococcus warneri		79.91	0.38	SUC 95			
Staphylococcus pasteuri		20.09	0.11	MLT 95			
740410	**Low discrimination**						**740410**
Staphylococcus pasteuri		63.50	0.35	SUC 99			
Staphylococcus warneri		36.50	0.45	MLT 86			
740412	**Low discrimination**						**740412**

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		68.07	0.27	SUC 99			
Staphylococcus warneri		31.93	0.35	MLT 86			
740413	**Good identification**						**740413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1			
740440	**Low discrimination**						**740440**
					PYR		
Staphylococcus warneri		52.64	0.74	SUC 95	8		
Staphylococcus petrasii subsp. croceilyticus		43.87	0.47		90		
Staphylococcus pasteuri		3.49	0.35		1		
740441	**Low discrimination**						**740441**
					PYR	YEP	
Staphylococcus warneri		66.44	0.39	SUC 95	8	43	
Staphylococcus petrasii subsp. croceilyticus		27.40	0.05	MNS 2	90	99	
Staphylococcus petrasii subsp. petrasii		3.64	-.02		99	1	
740442	**Good identification**						**740442**
					PYR		
Staphylococcus warneri		90.37	0.64	SUC 95	8		
Staphylococcus pasteuri		7.34	0.27		1		
Staphylococcus petrasii subsp. croceilyticus		2.28	0.05		90		
740450	**Low discrimination**						**740450**
					PYR		
Staphylococcus warneri		63.47	0.71	SUC 95	8		
Staphylococcus pasteuri		35.70	0.51		1		
Staphylococcus petrasii subsp. croceilyticus		0.74	0.05		90		
740451	**Low discrimination**						**740451**
					NOV		
Staphylococcus warneri		55.34	0.36	SUC 95	1		
Staphylococcus cohnii subsp. urealyticum		29.25	0.26	MNS 2	99		
Staphylococcus pasteuri		15.41	0.10		1		
740452	**Low discrimination**						**740452**
					NOV		
Staphylococcus warneri		58.22	0.61	SUC 95	1		
Staphylococcus pasteuri		40.13	0.43		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.45	0.07		99		
740453	**Acceptable identification**						**740453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		92.28	0.47	ARG 1	99		
Staphylococcus warneri		5.76	0.26		1		
Staphylococcus pasteuri		1.96	0.02		1		
740460	**Low discrimination**						**740460**
					PYR		
Staphylococcus warneri		54.55	0.33	SUC 95	8		
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	XYL 1	90		
740470	**Low discrimination**						**740470**
Staphylococcus warneri		64.00	0.30	SUC 95			
Staphylococcus pasteuri		36.00	0.10	XYL 1			
740500	**Good identification**						**740500**
Staphylococcus warneri		100.0	0.37	SUC 95			
740502	**Good identification**						**740502**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		100.0	0.27	SUC 95		
740510	**Good identification**					**740510**
Staphylococcus warneri		100.0	0.34	SUC 95		
740540	**Very good identification**					**740540**
					PYR	
Staphylococcus warneri		97.40	0.63	SUC 95	8	
Staphylococcus petrasii subsp. croceilyticus		2.60	0.05		90	
740541	**Low discrimination**					**740541**
					PYR	
Staphylococcus warneri		89.63	0.29	SUC 95	8	
Staphylococcus petrasii subsp. petrasii		10.37	-.05	MNS 2	99	
740542	**Very good identification**					**740542**
Staphylococcus warneri		100.0	0.54	SUC 95		
740550	**Very good identification**					**740550**
Staphylococcus warneri		98.23	0.61	SUC 95		
Staphylococcus pasteuri		1.77	0.10			
740552	**Good identification**					**740552**
Staphylococcus warneri		97.84	0.51	SUC 95		
Staphylococcus pasteuri		2.16	0.02			
740553	**Acceptable identification**					**740553**
					NOV	
Staphylococcus cohnii subsp. urealyticum		90.25	0.39	ARG 1	99	
Staphylococcus caprae		5.60	-.02		1	
Staphylococcus warneri		4.15	0.16		1	
740600	**Low discrimination**					**740600**
					PYR	
Staphylococcus pasteuri		36.83	0.60	MAN 86	1	
Staphylococcus warneri		34.46	0.74	MLT 86	8	
Staphylococcus petrasii subsp. croceilyticus		28.71	0.47		90	
740601	**Low discrimination**					**740601**
					PYR	
Staphylococcus warneri		50.10	0.39	MLT 95	8	
Staphylococcus pasteuri		26.50	0.19	MNS 2	1	
Staphylococcus petrasii subsp. croceilyticus		20.66	0.05		90	
740602	**Low discrimination**					**740602**
					PYR	
Staphylococcus pasteuri		56.03	0.52	MAN 86	1	
Staphylococcus warneri		42.79	0.64	MLT 86	8	
Staphylococcus petrasii subsp. croceilyticus		1.08	0.05		90	
740603	**Low discrimination**					**740603**
					YEP	
Staphylococcus warneri		48.92	0.29	MLT 95	43	
Staphylococcus pasteuri		31.71	0.11	MNS 2	86	
Staphylococcus simulans		19.37	0.11		1	
740610	**Low discrimination**					**740610**
					PYR	
Staphylococcus pasteuri		89.96	0.76	MLT 86	1	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		9.92	0.71		8		
Staphylococcus petrasii subsp. croceilyticus		0.12	0.05		90		
740611	**Low discrimination**						**740611**
Staphylococcus pasteuri		81.78	0.35	MLT 86			
Staphylococcus warneri		18.22	0.36	MNS 1			
740612	**Good identification**						**740612**
					YEP		
Staphylococcus pasteuri		91.64	0.68	MLT 86	86		
Staphylococcus warneri		8.25	0.61		43		
Staphylococcus simulans		0.11	0.13		1		
740613	**Low discrimination**						**740613**
					YEP		
Staphylococcus pasteuri		63.31	0.27	MLT 86	86		
Staphylococcus simulans		25.18	0.24	MNS 1	1		
Staphylococcus warneri		11.51	0.26		43		
740630	**Acceptable identification**						**740630**
Staphylococcus pasteuri		90.07	0.35	XYL 1			
Staphylococcus warneri		9.93	0.30	MLT 86			
740632	**Acceptable identification**						**740632**
Staphylococcus pasteuri		91.74	0.27	XYL 1			
Staphylococcus warneri		8.26	0.20	MLT 86			
740640	**Low discrimination**						**740640**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		76.32	0.88		90		
Staphylococcus warneri		17.58	1.00		8		
Staphylococcus pasteuri		6.07	0.76		1		
740641	**Low discrimination**						**740641**
					PYR	YEP	
Staphylococcus petrasii subsp. croceilyticus		59.55	0.47	MNS 1	90	99	
Staphylococcus warneri		27.71	0.65		8	43	
Staphylococcus petrasii subsp. petrasii		7.91	0.39		99	1	
740642	**Low discrimination**						**740642**
					PYR		
Staphylococcus warneri		64.24	0.90		8		
Staphylococcus pasteuri		27.20	0.68		1		
Staphylococcus petrasii subsp. croceilyticus		8.45	0.47		90		
740643	**Low discrimination**						**740643**
					PYR	YEP	
Staphylococcus warneri		57.30	0.55	MNS 2	8	43	
Staphylococcus petrasii subsp. petrasii		26.43	0.34		99	1	
Staphylococcus pasteuri		12.01	0.27		1	86	
740650	**Low discrimination**						**740650**
					PYR		
Staphylococcus pasteuri		73.42	0.92		1		
Staphylococcus warneri		25.05	0.97		8		
Staphylococcus petrasii subsp. croceilyticus		1.52	0.47		90		
740651	**Low discrimination**						**740651**
					PYR	YEP	
Staphylococcus pasteuri		57.47	0.51	MNS 1	1	86	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		39.61	0.62		8	43	
Staphylococcus petrasii subsp. petrasii		1.74	0.20		99	1	
740652	**Low discrimination**						**740652**
					NOV		
Staphylococcus pasteuri		77.96	0.84		1		
Staphylococcus warneri		21.70	0.87		1		
Staphylococcus saprophyticus subsp. saprophyticus		0.29	0.28		99		
740653	**Low discrimination**						**740653**
					PYR	YEP	
Staphylococcus pasteuri		60.47	0.43	MNS 1	1	86	
Staphylococcus warneri		34.01	0.53		8	43	
Staphylococcus petrasii subsp. petrasii		2.41	0.14		99	1	
740660	**Low discrimination**						**740660**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		76.34	0.47	XYL 1	90		
Staphylococcus warneri		17.58	0.59		8		
Staphylococcus pasteuri		6.08	0.35		1		
740662	**Low discrimination**						**740662**
					PYR		
Staphylococcus warneri		64.31	0.49	XYL 1	8		
Staphylococcus pasteuri		27.23	0.27		1		
Staphylococcus petrasii subsp. croceilyticus		8.46	0.05		90		
740670	**Low discrimination**						**740670**
					PYR		
Staphylococcus pasteuri		73.43	0.51	XYL 1	1		
Staphylococcus warneri		25.05	0.56		8		
Staphylococcus petrasii subsp. croceilyticus		1.52	0.05		90		
740672	**Low discrimination**						**740672**
Staphylococcus pasteuri		78.22	0.43	XYL 1			
Staphylococcus warneri		21.78	0.46				
740700	**Good identification**						**740700**
					PYR		
Staphylococcus warneri		94.27	0.63	MLT 95	8		
Staphylococcus pasteuri		3.22	0.19		1		
Staphylococcus petrasii subsp. croceilyticus		2.51	0.05		90		
740701	**Low discrimination**						**740701**
					PYR		
Staphylococcus warneri		89.63	0.29	MLT 95	8		
Staphylococcus petrasii subsp. petrasii		10.37	-.05	MNS 2	99		
740702	**Very good identification**						**740702**
Staphylococcus warneri		95.98	0.54	MLT 95			
Staphylococcus pasteuri		4.02	0.11				
740710	**Low discrimination**						**740710**
Staphylococcus warneri		77.52	0.61	MLT 95			
Staphylococcus pasteuri		22.48	0.35				
740712	**Low discrimination**						**740712**
Staphylococcus warneri		73.78	0.51	MLT 95			
Staphylococcus pasteuri		26.22	0.27				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
740740	**Low discrimination**						**740740**
					PYR		
Staphylococcus warneri		86.84	0.90		8		
Staphylococcus petrasii subsp. croceilyticus		12.06	0.47		90		
Staphylococcus pasteuri		0.96	0.35		1		
740741	**Low discrimination**						**740741**
					PYR	YEP	
Staphylococcus warneri		59.83	0.55	MNS 2	8	43	
Staphylococcus petrasii subsp. petrasii		36.06	0.36		99	1	
Staphylococcus petrasii subsp. croceilyticus		4.11	0.05		90	99	
740742	**Very good identification**						**740742**
					PYR		
Staphylococcus warneri		98.01	0.80		8		
Staphylococcus pasteuri		1.33	0.27		1		
Staphylococcus petrasii subsp. croceilyticus		0.41	0.05		90		
740743	**Low discrimination**						**740743**
					PYR		
Staphylococcus warneri		50.67	0.45	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		49.34	0.30		99		
740750	**Good identification**						**740750**
					PYR		
Staphylococcus warneri		91.22	0.87		8		
Staphylococcus pasteuri		8.55	0.51		1		
Staphylococcus petrasii subsp. croceilyticus		0.18	0.05		90		
740751	**Low discrimination**						**740751**
					PYR	YEP	
Staphylococcus warneri		87.80	0.52	MNS 2	8	43	
Staphylococcus petrasii subsp. petrasii		8.12	0.16		99	1	
Staphylococcus pasteuri		4.08	0.10		1	86	
740752	**Low discrimination**						**740752**
					NOV		
Staphylococcus warneri		89.40	0.77		1		
Staphylococcus pasteuri		10.27	0.43		1		
Staphylococcus saprophyticus subsp. saprophyticus		0.33	0.06		99		
740753	**Low discrimination**						**740753**
					PYR	YEP	
Staphylococcus warneri		78.87	0.42	MNS 2	8	43	
Staphylococcus petrasii subsp. petrasii		11.78	0.11		99	1	
Staphylococcus pasteuri		4.49	0.02		1	86	
740760	**Low discrimination**						**740760**
					PYR		
Staphylococcus warneri		87.81	0.49	XYL 1	8		
Staphylococcus petrasii subsp. croceilyticus		12.19	0.05		90		
740762	**Good identification**						**740762**
Staphylococcus warneri		100.0	0.39	XYL 1			
740770	**Acceptable identification**						**740770**
Staphylococcus warneri		91.43	0.46	XYL 1			
Staphylococcus pasteuri		8.57	0.10				
740772	**Low discrimination**						**740772**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		89.70	0.36	XYL 1		
Staphylococcus pasteuri		10.31	0.02			
741240	**Low discrimination**					**741240**
					PYR	
Staphylococcus warneri		54.55	0.33	ESL 1	8	
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	TRE 95	90	
741250	**Low discrimination**					**741250**
Staphylococcus warneri		64.00	0.30	ESL 1		
Staphylococcus pasteuri		36.00	0.10	TRE 95		
741440	**Low discrimination**					**741440**
					PYR	
Staphylococcus warneri		54.55	0.33	ESL 1	8	
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	SUC 95	90	
741450	**Low discrimination**					**741450**
Staphylococcus warneri		64.00	0.30	ESL 1		
Staphylococcus pasteuri		36.00	0.10	SUC 95		
741453	**Good identification**					**741453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.39	ARG 1		
741553	**Good identification**					**741553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1		
741610	**Acceptable identification**					**741610**
Staphylococcus pasteuri		90.07	0.35	ESL 1		
Staphylococcus warneri		9.93	0.30	MLT 86		
741612	**Acceptable identification**					**741612**
Staphylococcus pasteuri		91.74	0.27	ESL 1		
Staphylococcus warneri		8.26	0.20	MLT 86		
741640	**Low discrimination**					**741640**
					PYR	
Staphylococcus petrasii subsp. croceilyticus		76.34	0.47	ESL 1	90	
Staphylococcus warneri		17.58	0.59		8	
Staphylococcus pasteuri		6.08	0.35		1	
741642	**Low discrimination**					**741642**
					PYR	
Staphylococcus warneri		64.31	0.49	ESL 1	8	
Staphylococcus pasteuri		27.23	0.27		1	
Staphylococcus petrasii subsp. croceilyticus		8.46	0.05		90	
741650	**Low discrimination**					**741650**
					PYR	
Staphylococcus pasteuri		73.43	0.51	ESL 1	1	
Staphylococcus warneri		25.05	0.56		8	
Staphylococcus petrasii subsp. croceilyticus		1.52	0.05		90	
741652	**Low discrimination**					**741652**
Staphylococcus pasteuri		78.22	0.43	ESL 1		
Staphylococcus warneri		21.78	0.46			
741740	**Low discrimination**					**741740**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus warneri		87.81	0.49	ESL 1	8	
Staphylococcus petrasii subsp. croceilyticus		12.19	0.05		90	
741742	**Good identification**					**741742**
Staphylococcus warneri		100.0	0.39	ESL 1		
741750	**Acceptable identification**					**741750**
Staphylococcus warneri		91.43	0.46	ESL 1		
Staphylococcus pasteuri		8.57	0.10			
741752	**Low discrimination**					**741752**
Staphylococcus warneri		89.70	0.36	ESL 1		
Staphylococcus pasteuri		10.31	0.02			
742042	**Good identification**					**742042**
Staphylococcus warneri		100.0	0.30	SUC 95		
742050	**Low discrimination**					**742050**
Staphylococcus warneri		61.85	0.37	SUC 95		
Staphylococcus pasteuri		38.15	0.18	TRE 95		
742052	**Low discrimination**					**742052**
Staphylococcus warneri		51.47	0.27	SUC 95		
Staphylococcus pasteuri		38.90	0.10	TRE 95		
Staphylococcus capitis subsp. ureolyticus		9.63	-.11			
742053	**Good identification**					**742053**
Staphylococcus capitis subsp. ureolyticus		100.0	0.31	GLR 1		
742140	**Good identification**					**742140**
Staphylococcus warneri		100.0	0.30	SUC 95		
742150	**Good identification**					**742150**
Staphylococcus warneri		100.0	0.27	SUC 95		
742210	**Acceptable identification**					**742210**
Staphylococcus pasteuri		90.86	0.43	TRE 99		
Staphylococcus warneri		9.14	0.37	MLT 86		
742212	**Low discrimination**					**742212**
					YEP	
Staphylococcus pasteuri		86.30	0.35	TRE 99	86	
Staphylococcus warneri		7.09	0.27	MLT 86	43	
Staphylococcus simulans		5.29	0.15		1	
742213	**Low discrimination**					**742213**
Staphylococcus capitis subsp. ureolyticus		88.11	0.31	GLR 1		
Staphylococcus simulans		11.89	0.26	MLT 96		
742240	**Low discrimination**					**742240**
					PYR	
Staphylococcus petrasii subsp. croceilyticus		84.33	0.59	TRE 99	90	
Staphylococcus warneri		10.84	0.66		8	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		4.11	0.43		1		
742242	**Low discrimination**						**742242**
					PYR		
Staphylococcus warneri		52.15	0.56	TRE 95	8		
Staphylococcus pasteuri		24.21	0.35		1		
Staphylococcus petrasii subsp. croceilyticus		12.29	0.18		90		
742243	**Low discrimination**						**742243**
					PYR		
Staphylococcus capitis subsp. ureolyticus		79.26	0.31	GLR 1	1		
Staphylococcus epidermidis		12.98	0.28	MAN 96	1		
Staphylococcus petrasii subsp. petrasii		4.26	0.12		99		
742250	**Low discrimination**						**742250**
					PYR		
Staphylococcus pasteuri		73.81	0.59	TRE 99	1		
Staphylococcus warneri		22.97	0.63		8		
Staphylococcus petrasii subsp. croceilyticus		2.49	0.18		90		
742251	**Acceptable identification**						**742251**
Staphylococcus capitis subsp. ureolyticus		94.96	0.41	GLR 1			
Staphylococcus pasteuri		2.90	0.18	LAC 88			
Staphylococcus warneri		1.82	0.28				
742252	**Low discrimination**						**742252**
Staphylococcus pasteuri		75.17	0.51	TRE 99			
Staphylococcus warneri		19.09	0.53				
Staphylococcus capitis subsp. ureolyticus		4.51	0.18				
742253	**Very good identification**						**742253**
					YEP		
Staphylococcus capitis subsp. ureolyticus		99.49	0.59	GLR 1	77		
Staphylococcus simulans		0.25	0.19		1		
Staphylococcus pasteuri		0.17	0.10		86		
742300	**Good identification**						**742300**
Staphylococcus warneri		100.0	0.30	TRE 95			
742310	**Low discrimination**						**742310**
Staphylococcus warneri		75.88	0.27	TRE 95			
Staphylococcus pasteuri		24.13	0.02	MLT 95			
742340	**Low discrimination**						**742340**
					PYR	YEP	
Staphylococcus warneri		60.82	0.56	TRE 95	8	43	
Staphylococcus petrasii subsp. croceilyticus		15.13	0.18		90	99	
Staphylococcus epidermidis		7.15	0.28		1	1	
742341	**Low discrimination**						**742341**
					PYR	NOV	
Staphylococcus epidermidis		69.96	0.33	GLR 1	1	1	
Staphylococcus petrasii subsp. petrasii		15.89	0.14		99	1	
Staphylococcus hominis subsp. novobiosepticus		7.61	0.05		1	99	
742342	**Low discrimination**						**742342**
Staphylococcus epidermidis		40.93	0.38	GLR 1			
Staphylococcus warneri		36.65	0.46				

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus hominis subsp. hominis		12.63	0.31				
742343	**Acceptable identification**						**742343**
					PYR	NOV	
Staphylococcus epidermidis		92.78	0.43	GLR 1	1	1	
Staphylococcus petrasii subsp. petrasii		3.58	0.09		99	1	
Staphylococcus hominis subsp. novobiosepticus		1.72	0.00		1	99	
742350	**Low discrimination**						**742350**
					PYR		
Staphylococcus warneri		84.91	0.53	TRE 95	8		
Staphylococcus pasteuri		8.73	0.18		1		
Staphylococcus haemolyticus		4.64	0.20		98		
742352	**Low discrimination**						**742352**
					PYR		
Staphylococcus warneri		77.56	0.43	TRE 95	8		
Staphylococcus pasteuri		9.77	0.10		1		
Staphylococcus haemolyticus		7.16	0.15		98		
742410	**Acceptable identification**						**742410**
Staphylococcus pasteuri		90.86	0.43	SUC 99			
Staphylococcus warneri		9.14	0.37	MLT 86			
742412	**Low discrimination**						**742412**
					YEP		
Staphylococcus pasteuri		88.45	0.35	SUC 99	86		
Staphylococcus warneri		7.26	0.27	MLT 86	43		
Staphylococcus simulans		4.29	0.13		1		
742440	**Low discrimination**						**742440**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		84.85	0.59	SUC 99	90		
Staphylococcus warneri		10.91	0.66		8		
Staphylococcus pasteuri		4.13	0.43		1		
742442	**Low discrimination**						**742442**
					PYR		
Staphylococcus warneri		58.83	0.56	SUC 95	8		
Staphylococcus pasteuri		27.31	0.35		1		
Staphylococcus petrasii subsp. croceilyticus		13.86	0.18		90		
742450	**Low discrimination**						**742450**
					PYR		
Staphylococcus pasteuri		74.27	0.59	SUC 99	1		
Staphylococcus warneri		23.11	0.63		8		
Staphylococcus petrasii subsp. croceilyticus		2.51	0.18		90		
742452	**Low discrimination**						**742452**
					YEP		
Staphylococcus pasteuri		79.52	0.51	SUC 99	86		
Staphylococcus warneri		20.19	0.53		43		
Staphylococcus simulans		0.28	0.06		1		
742453	**Low discrimination**						**742453**
					NOV	YEP	
Staphylococcus cohnii subsp. urealyticum		53.56	0.31	ARG 1	99	64	
Staphylococcus simulans		18.60	0.16		1	1	
Staphylococcus pasteuri		15.81	0.10		1	86	
742500	**Good identification**						**742500**

Code / Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		100.0	0.30	SUC 95			
742510	**Low discrimination**						**742510**
Staphylococcus warneri		75.88	0.27	SUC 95			
Staphylococcus pasteuri		24.13	0.02	MLT 95			
742540	**Low discrimination**						**742540**
					PYR		
Staphylococcus warneri		78.41	0.56	SUC 95	8		
Staphylococcus petrasii subsp. croceilyticus		19.50	0.18		90		
Staphylococcus pasteuri		0.95	0.02		1		
742542	**Good identification**						**742542**
					PYR		
Staphylococcus warneri		98.50	0.46	SUC 95	8		
Staphylococcus haemolyticus		1.50	0.01		98		
742550	**Low discrimination**						**742550**
					NOV	PYR	
Staphylococcus warneri		85.48	0.53	SUC 95	1	8	
Staphylococcus pasteuri		8.79	0.18		1	1	
Staphylococcus saprophyticus subsp. bovis		5.11	0.18		99	99	
742552	**Low discrimination**						**742552**
					YEP		
Staphylococcus warneri		80.02	0.43	SUC 95	43		
Staphylococcus pasteuri		10.08	0.10		86		
Staphylococcus caprae		6.31	0.00		1		
742600	**Low discrimination**						**742600**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		52.23	0.59	MLT 99	90		
Staphylococcus pasteuri		41.01	0.68		1		
Staphylococcus warneri		6.72	0.66		8		
742602	**Low discrimination**						**742602**
					PYR		
Staphylococcus pasteuri		84.51	0.60	MAN 86	1		
Staphylococcus warneri		11.30	0.56	MLT 86	8		
Staphylococcus petrasii subsp. croceilyticus		2.66	0.18		90		
742603	**Low discrimination**						**742603**
					YEP		
Staphylococcus simulans		69.35	0.40	VPT 4	1		
Staphylococcus pasteuri		11.58	0.19	BGA 90	86		
Staphylococcus hyicus		10.12	0.15		1		
742610	**Very good identification**						**742610**
					PYR		
Staphylococcus pasteuri		97.90	0.84	MLT 86	1		
Staphylococcus warneri		1.89	0.63		8		
Staphylococcus petrasii subsp. croceilyticus		0.21	0.18		90		
742611	**Acceptable identification**						**742611**
					YEP		
Staphylococcus pasteuri		94.16	0.43	MLT 86	86		
Staphylococcus warneri		3.67	0.28	MNS 1	43		
Staphylococcus simulans		1.52	0.11		1		
742612	**Very good identification**						**742612**
					YEP		
Staphylococcus pasteuri		97.32	0.76	MLT 86	86		

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		1.53	0.53		43		
Staphylococcus simulans		1.15	0.41		1		
742613	**Low discrimination**						**742613**
					YEP		
Staphylococcus simulans		78.83	0.52	VPT 4	1		
Staphylococcus pasteuri		20.22	0.35	BGA 90	86		
Staphylococcus warneri		0.64	0.19		43		
742630	**Good identification**						**742630**
Staphylococcus pasteuri		98.11	0.43	XYL 1			
Staphylococcus warneri		1.89	0.22	MLT 86			
742632	**Good identification**						**742632**
					YEP		
Staphylococcus pasteuri		97.32	0.35	XYL 1	86		
Staphylococcus warneri		1.53	0.12	MLT 86	43		
Staphylococcus simulans		1.15	0.00		1		
742640	**Good identification**						**742640**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		93.09	1.00		90		
Staphylococcus pasteuri		4.53	0.84		1		
Staphylococcus warneri		2.30	0.92		8		
742641	**Low discrimination**						**742641**
					PYR	YEP	
Staphylococcus petrasii subsp. croceilyticus		71.49	0.59	MNS 1	90	99	
Staphylococcus petrasii subsp. petrasii		21.37	0.59		99	1	
Staphylococcus warneri		3.56	0.58		8	43	
742642	**Low discrimination**						**742642**
					PYR		
Staphylococcus pasteuri		51.33	0.76	MAN 86	1		
Staphylococcus petrasii subsp. croceilyticus		26.06	0.59		90		
Staphylococcus warneri		21.22	0.83		8		
742643	**Low discrimination**						**742643**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		73.93	0.53	GLR 1	99	1	
Staphylococcus pasteuri		9.14	0.35		1	86	
Staphylococcus warneri		7.63	0.48		8	43	
742650	**Good identification**						**742650**
					PYR		
Staphylococcus pasteuri		91.38	1.00		1		
Staphylococcus warneri		5.46	0.90		8		
Staphylococcus petrasii subsp. croceilyticus		3.09	0.59		90		
742651	**Low discrimination**						**742651**
					PYR	YEP	
Staphylococcus pasteuri		78.53	0.59	MNS 1	1	86	
Staphylococcus warneri		9.48	0.55		8	43	
Staphylococcus petrasii subsp. petrasii		8.72	0.39		99	1	
742652	**Very good identification**						**742652**
					PYR		
Staphylococcus pasteuri		95.09	0.92		1		
Staphylococcus warneri		4.63	0.80		8		
Staphylococcus haemolyticus		0.08	0.37		98		
742653	**Low discrimination**						**742653**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					YEP	PYR	
Staphylococcus pasteuri		62.72	0.51	MNS 1	86	1	
Staphylococcus simulans		17.89	0.45		1	70	
Staphylococcus petrasii subsp. petrasii		9.18	0.34		1	99	
742660	**Good identification**						**742660**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		93.11	0.59	XYL 1	90		
Staphylococcus pasteuri		4.54	0.43		1		
Staphylococcus warneri		2.30	0.51		8		
742662	**Low discrimination**						**742662**
					PYR		
Staphylococcus pasteuri		52.05	0.35	MAN 86	1		
Staphylococcus petrasii subsp. croceilyticus		26.43	0.18	XYL 1	90		
Staphylococcus warneri		21.52	0.42		8		
742670	**Good identification**						**742670**
					PYR		
Staphylococcus pasteuri		91.45	0.59	XYL 1	1		
Staphylococcus warneri		5.46	0.49		8		
Staphylococcus petrasii subsp. croceilyticus		3.09	0.18		90		
742672	**Good identification**						**742672**
Staphylococcus pasteuri		95.35	0.51	XYL 1			
Staphylococcus warneri		4.65	0.39				
742700	**Low discrimination**						**742700**
					PYR		
Staphylococcus warneri		68.56	0.56	MLT 95	8		
Staphylococcus petrasii subsp. croceilyticus		17.06	0.18		90		
Staphylococcus pasteuri		13.39	0.27		1		
742702	**Low discrimination**						**742702**
					YEP		
Staphylococcus warneri		75.98	0.46	MLT 95	43		
Staphylococcus pasteuri		18.18	0.19		86		
Staphylococcus simulans		2.83	0.07		1		
742703	**Acceptable identification**						**742703**
					PYR		
Staphylococcus hyicus		91.44	0.37	VPT 1	1		
Staphylococcus simulans		4.74	0.18	PHS 92	70		
Staphylococcus petrasii subsp. petrasii		3.05	0.09		99		
742710	**Low discrimination**						**742710**
					PYR		
Staphylococcus pasteuri		62.19	0.43	GAL 1	1		
Staphylococcus warneri		37.54	0.53	MLT 86	8		
Staphylococcus haemolyticus		0.28	0.02		98		
742712	**Low discrimination**						**742712**
					YEP		
Staphylococcus pasteuri		62.74	0.35	GAL 1	86		
Staphylococcus warneri		30.91	0.43	MLT 86	43		
Staphylococcus simulans		6.35	0.19		1		
742713	**Good identification**						**742713**
Staphylococcus simulans		95.35	0.30	VPT 4			
Staphylococcus hyicus		4.65	-.04	BGA 90			

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
742740	**Low discrimination**						**742740**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		52.87	0.59	GAL 1	90		
Staphylococcus warneri		40.79	0.82		8		
Staphylococcus pasteuri		2.58	0.43		1		
742741	**Low discrimination**						**742741**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		87.79	0.55	GLR 1	99	1	
Staphylococcus warneri		6.94	0.47		8	43	
Staphylococcus petrasii subsp. croceilyticus		4.45	0.18		90	99	
742742	**Low discrimination**						**742742**
					PYR		
Staphylococcus warneri		76.87	0.72		8		
Staphylococcus haemolyticus		6.11	0.43		98		
Staphylococcus pasteuri		5.95	0.35		1		
742743	**Acceptable identification**						**742743**
					PYR		
Staphylococcus petrasii subsp. petrasii		93.20	0.50	GLR 1	99		
Staphylococcus warneri		4.56	0.38		8		
Staphylococcus simulans		0.66	0.11		70		
742750	**Low discrimination**						**742750**
					PYR		
Staphylococcus warneri		62.02	0.79		8		
Staphylococcus pasteuri		33.22	0.59		1		
Staphylococcus haemolyticus		2.37	0.43		98		
742751	**Low discrimination**						**742751**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		56.35	0.36	GLR 1	99	1	
Staphylococcus warneri		29.01	0.44	MAN 10	8	43	
Staphylococcus pasteuri		7.69	0.18		1	86	
742752	**Low discrimination**						**742752**
					PYR		
Staphylococcus warneri		56.96	0.69		8		
Staphylococcus pasteuri		37.38	0.51		1		
Staphylococcus haemolyticus		3.67	0.38		98		
742753	**Low discrimination**						**742753**
					COA	PYR	
Staphylococcus petrasii subsp. petrasii		35.41	0.30	GLR 1	1	99	
Staphylococcus aureus subsp. aureus		29.35	0.18	MAN 10	99	10	
Staphylococcus warneri		11.29	0.35		1	8	
742762	**Acceptable identification**						**742762**
					PYR		
Staphylococcus warneri		92.63	0.31	XYL 1	8		
Staphylococcus haemolyticus		7.37	0.01		98		
742770	**Low discrimination**						**742770**
					PYR		
Staphylococcus warneri		63.54	0.38	XYL 1	8		
Staphylococcus pasteuri		34.03	0.18		1		
Staphylococcus haemolyticus		2.43	0.02		98		
742772	**Low discrimination**						**742772**
Staphylococcus warneri		60.38	0.28	XYL 1			

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus pasteuri		39.63	0.10			
742773	**Good identification**					**742773**
Staphylococcus xylosus		100.0	0.30	VPT 14		
743610	**Good identification**					**743610**
Staphylococcus pasteuri		98.11	0.43	ESL 1		
Staphylococcus warneri		1.89	0.22	MLT 86		
743612	**Good identification**					**743612**
					YEP	
Staphylococcus pasteuri		97.32	0.35	ESL 1	86	
Staphylococcus warneri		1.53	0.12	MLT 86	43	
Staphylococcus simulans		1.15	0.00		1	
743640	**Good identification**					**743640**
					PYR	
Staphylococcus petrasii subsp. croceilyticus		93.09	0.59	ESL 1	90	
Staphylococcus pasteuri		4.53	0.43		1	
Staphylococcus warneri		2.30	0.51		8	
743642	**Low discrimination**					**743642**
					PYR	
Staphylococcus pasteuri		51.61	0.35	ESL 1	1	
Staphylococcus petrasii subsp. croceilyticus		26.20	0.18	MAN 86	90	
Staphylococcus warneri		21.34	0.42		8	
743650	**Good identification**					**743650**
					PYR	
Staphylococcus pasteuri		91.45	0.59	ESL 1	1	
Staphylococcus warneri		5.46	0.49		8	
Staphylococcus petrasii subsp. croceilyticus		3.09	0.18		90	
743652	**Good identification**					**743652**
Staphylococcus pasteuri		95.35	0.51	ESL 1		
Staphylococcus warneri		4.65	0.39			
743742	**Low discrimination**					**743742**
					PYR	
Staphylococcus warneri		78.67	0.31	ESL 1	8	
Staphylococcus piscifermentans		15.07	0.18		15	
Staphylococcus haemolyticus		6.26	0.01		98	
743750	**Low discrimination**					**743750**
					PYR	
Staphylococcus warneri		63.54	0.38	ESL 1	8	
Staphylococcus pasteuri		34.03	0.18		1	
Staphylococcus haemolyticus		2.43	0.02		98	
743752	**Low discrimination**					**743752**
					YEP	
Staphylococcus warneri		56.51	0.28	ESL 1	43	
Staphylococcus pasteuri		37.09	0.10		86	
Staphylococcus piscifermentans		6.41	0.10		1	
744240	**Low discrimination**					**744240**
					PYR	
Staphylococcus warneri		54.55	0.33	PHS 1	8	
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	TRE 95	90	
744250	**Low discrimination**					**744250**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		64.00	0.30	PHS 1			
Staphylococcus pasteuri		36.00	0.10	TRE 95			
744343	**Good identification**						**744343**
Staphylococcus epidermidis		100.0	0.28	GLR 1			
744413	**Good identification**						**744413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ARG 1			
744440	**Low discrimination**						**744440**
					PYR		
Staphylococcus warneri		54.55	0.33	PHS 1	8		
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	SUC 95	90		
744443	**Good identification**						**744443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ARG 1			
744450	**Low discrimination**						**744450**
					NOV		
Staphylococcus warneri		53.29	0.30	PHS 1	1		
Staphylococcus pasteuri		29.98	0.10	SUC 95	1		
Staphylococcus kloosii		16.74	0.27		99		
744451	**Good identification**						**744451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ARG 1			
744452	**Low discrimination**						**744452**
					NOV		
Staphylococcus kloosii		31.89	0.28	VPT 10	99		
Staphylococcus warneri		31.24	0.20	ARG 1	1		
Staphylococcus pasteuri		21.54	0.02		1		
744453	**Good identification**						**744453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		98.26	0.51	ARG 1	99		
Staphylococcus caprae		1.74	-.02		1		
744513	**Low discrimination**						**744513**
					NOV		
Staphylococcus cohnii subsp. urealyticum		81.02	0.28	ARG 1	99		
Staphylococcus caprae		18.98	-.02		1		
744550	**Low discrimination**						**744550**
					NOV	YEP	
Staphylococcus kloosii		53.98	0.33	VPT 10	99	99	
Staphylococcus warneri		27.96	0.19	ARG 1	1	43	
Staphylococcus caprae		18.06	-.05		1	1	
744553	**Low discrimination**						**744553**
					NOV		
Staphylococcus caprae		80.37	0.39	GLR 1	1		
Staphylococcus cohnii subsp. urealyticum		19.63	0.43	NIT 90	99		
744610	**Acceptable identification**						**744610**
Staphylococcus pasteuri		90.07	0.35	PHS 1			
Staphylococcus warneri		9.93	0.30	MLT 86			
744612	**Low discrimination**						**744612**
					YEP		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
Staphylococcus pasteuri		88.53	0.27	PHS 1	86		
Staphylococcus warneri		7.97	0.20	MLT 86	43		
Staphylococcus simulans		3.51	0.03		1		
744640	**Low discrimination**						**744640**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		76.34	0.47	PHS 1	90		
Staphylococcus warneri		17.58	0.59		8		
Staphylococcus pasteuri		6.08	0.35		1		
744642	**Low discrimination**						**744642**
					PYR		
Staphylococcus warneri		64.31	0.49	PHS 1	8		
Staphylococcus pasteuri		27.23	0.27		1		
Staphylococcus petrasii subsp. croceilyticus		8.46	0.05		90		
744650	**Low discrimination**						**744650**
					PYR		
Staphylococcus pasteuri		73.43	0.51	PHS 1	1		
Staphylococcus warneri		25.05	0.56		8		
Staphylococcus petrasii subsp. croceilyticus		1.52	0.05		90		
744652	**Low discrimination**						**744652**
Staphylococcus pasteuri		78.22	0.43	PHS 1			
Staphylococcus warneri		21.78	0.46				
744740	**Low discrimination**						**744740**
					PYR		
Staphylococcus warneri		87.81	0.49	PHS 1	8		
Staphylococcus petrasii subsp. croceilyticus		12.19	0.05		90		
744742	**Good identification**						**744742**
Staphylococcus warneri		100.0	0.39	PHS 1			
744750	**Acceptable identification**						**744750**
Staphylococcus warneri		91.43	0.46	PHS 1			
Staphylococcus pasteuri		8.57	0.10				
744752	**Low discrimination**						**744752**
					YEP		
Staphylococcus warneri		54.60	0.36	PHS 1	43		
Staphylococcus caprae		39.12	0.13		1		
Staphylococcus pasteuri		6.27	0.02		86		
744753	**Low discrimination**						**744753**
					COA	NOV	
Staphylococcus caprae		69.01	0.30	GLR 1	1	1	
Staphylococcus aureus subsp. aureus		29.58	0.18	NIT 90	99	4	
Staphylococcus cohnii subsp. urealyticum		1.41	0.12		1	99	
745413	**Good identification**						**745413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1			
745450	**Good identification**						**745450**
Staphylococcus kloosii		100.0	0.27	VPT 10			
745452	**Good identification**						**745452**
Staphylococcus kloosii		100.0	0.28	VPT 10			

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
745453	**Good identification**					**745453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	ARG 1		
745550	**Good identification**					**745550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
745552	**Good identification**					**745552**
Staphylococcus kloosii		100.0	0.34	VPT 10		
745553	**Acceptable identification**					**745553**
					NOV	
Staphylococcus cohnii subsp. urealyticum		90.39	0.35	ARG 1	99	
Staphylococcus caprae		9.61	-.02		1	
746101	**Good identification**					**746101**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.19	0.37	GLR 1	99	
Staphylococcus schleiferi subsp. schleiferi		3.81	0.04	LAC 86	1	
746103	**Very good identification**					**746103**
Staphylococcus schleiferi subsp. coagulans		99.59	0.53	GLR 1		
Staphylococcus hyicus		0.42	-.04			
746111	**Good identification**					**746111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	GLR 1		
746113	**Good identification**					**746113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.14	0.53	GLR 1	99	
Staphylococcus caprae		0.86	-.02		1	
746153	**Good identification**					**746153**
					COA	
Staphylococcus caprae		98.83	0.39	GLR 1	1	
Staphylococcus schleiferi subsp. coagulans		1.17	0.12	TRE 90	99	
746241	**Good identification**					**746241**
Staphylococcus epidermidis		100.0	0.33	GLR 1		
746242	**Good identification**					**746242**
Staphylococcus epidermidis		97.22	0.38	GLR 1		
Staphylococcus warneri		2.79	0.15			
746243	**Good identification**					**746243**
Staphylococcus epidermidis		98.92	0.43	GLR 1		
Staphylococcus capitis subsp. ureolyticus		1.08	-.11			
746301	**Low discrimination**					**746301**
					COA	
Staphylococcus schleiferi subsp. coagulans		54.87	0.27	GLR 1	99	
Staphylococcus hyicus		38.27	0.15	LAC 86	24	
Staphylococcus epidermidis		6.86	0.08		1	
746303	**Low discrimination**					**746303**
					COA	
Staphylococcus hyicus		54.89	0.37	VPT 1	24	
Staphylococcus schleiferi subsp. coagulans		42.04	0.43	TRE 92	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus epidermidis		2.71	0.18		1		
746313	**Good identification**						**746313**
					COA		
Staphylococcus schleiferi subsp. coagulans		97.62	0.42	GLR 1	99		
Staphylococcus hyicus		1.40	-.04		24		
Staphylococcus caprae		0.99	-.11		1		
746340	**Good identification**						**746340**
Staphylococcus epidermidis		98.51	0.43	GLR 1			
Staphylococcus warneri		1.49	0.15				
746341	**Good identification**						**746341**
Staphylococcus epidermidis		100.0	0.49	GLR 1			
746342	**Very good identification**						**746342**
Staphylococcus epidermidis		100.0	0.53	GLR 1			
746343	**Very good identification**						**746343**
Staphylococcus epidermidis		98.75	0.59	GLR 1			
Staphylococcus caprae		0.75	0.04				
Staphylococcus hyicus		0.20	-.04				
746353	**Low discrimination**						**746353**
					COA		
Staphylococcus caprae		56.11	0.30	GLR 1	1		
Staphylococcus aureus subsp. aureus		24.05	0.18	TRE 90	99		
Staphylococcus delphini		15.36	0.18		99		
746453	**Low discrimination**						**746453**
					NOV		
Staphylococcus cohnii subsp. urealyticum		49.68	0.35	ARG 1	99		
Staphylococcus caprae		44.89	0.18		1		
Staphylococcus simulans		3.83	0.07		1		
746543	**Good identification**						**746543**
					NOV		
Staphylococcus caprae		99.11	0.33	GLR 1	1		
Staphylococcus cohnii subsp. urealyticum		0.89	0.07	MAN 95	99		
746551	**Good identification**						**746551**
					NOV		
Staphylococcus caprae		99.11	0.33	GLR 1	1		
Staphylococcus cohnii subsp. urealyticum		0.89	0.07	LAC 95	99		
746552	**Good identification**						**746552**
Staphylococcus caprae		100.0	0.41	GLR 1			
746553	**Very good identification**						**746553**
					COA	NOV	
Staphylococcus caprae		97.81	0.59	GLR 1	1	1	
Staphylococcus aureus subsp. aureus		1.72	0.18		99	4	
Staphylococcus cohnii subsp. urealyticum		0.47	0.28		1	99	
746603	**Low discrimination**						**746603**
					YEP		
Staphylococcus hyicus		82.85	0.37	VPT 1	1		
Staphylococcus simulans		16.45	0.30	GAL 92	1		
Staphylococcus chromogenes		0.69	-.02		95		

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
746610	**Good identification**						**746610**
	Staphylococcus pasteuri	98.11	0.43	PHS 1			
	Staphylococcus warneri	1.89	0.22	MLT 86			
746612	**Low discrimination**						**746612**
					YEP		
	Staphylococcus pasteuri	71.23	0.35	PHS 1	86		
	Staphylococcus simulans	27.65	0.31	MLT 86	1		
	Staphylococcus warneri	1.12	0.12		43		
746613	**Good identification**						**746613**
	Staphylococcus simulans	97.09	0.42	VPT 4			
	Staphylococcus hyicus	1.24	-.04	BGA 90			
	Staphylococcus felis	0.91	-.09				
746640	**Good identification**						**746640**
					PYR		
	Staphylococcus petrasii subsp. croceilyticus	93.07	0.59	PHS 1	90		
	Staphylococcus pasteuri	4.53	0.43		1		
	Staphylococcus warneri	2.30	0.51		8		
746642	**Low discrimination**						**746642**
					PYR		
	Staphylococcus pasteuri	50.36	0.35	PHS 1	1		
	Staphylococcus petrasii subsp. croceilyticus	25.56	0.18	MAN 86	90		
	Staphylococcus warneri	20.82	0.42		8		
746650	**Good identification**						**746650**
					PYR		
	Staphylococcus pasteuri	91.33	0.59	PHS 1	1		
	Staphylococcus warneri	5.45	0.49		8		
	Staphylococcus petrasii subsp. croceilyticus	3.08	0.18		90		
746652	**Acceptable identification**						**746652**
					YEP		
	Staphylococcus pasteuri	92.50	0.51	PHS 1	86		
	Staphylococcus warneri	4.51	0.39		43		
	Staphylococcus simulans	2.63	0.24		1		
746701	**Good identification**						**746701**
					YEP		
	Staphylococcus hyicus	99.17	0.37	VPT 1	1		
	Staphylococcus chromogenes	0.83	-.02	LAC 92	95		
746703	**Very good identification**						**746703**
					YEP		
	Staphylococcus hyicus	99.13	0.59	VPT 1	1		
	Staphylococcus chromogenes	0.65	0.18		95		
	Staphylococcus simulans	0.15	0.08		1		
746742	**Low discrimination**						**746742**
	Staphylococcus warneri	46.26	0.31	PHS 1			
	Staphylococcus caprae	26.53	0.06				
	Staphylococcus epidermidis	15.41	0.12				
746750	**Low discrimination**						**746750**
					NOV	PYR	
	Staphylococcus warneri	46.53	0.38	PHS 1	1	8	
	Staphylococcus pasteuri	24.92	0.18		1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus saprophyticus subsp. bovis		14.49	0.18		99 99	
746752	**Low discrimination**					**746752**
					COA	
Staphylococcus caprae		67.32	0.32	GLR 1	1	
Staphylococcus aureus subsp. aureus		23.51	0.18	MNS 88	99	
Staphylococcus warneri		4.47	0.28		1	
746753	**Low discrimination**					**746753**
					COA	
Staphylococcus aureus subsp. aureus		82.37	0.59	GLR 1	99	
Staphylococcus caprae		17.47	0.50		1	
Staphylococcus simulans		0.09	0.13		1	
747101	**Good identification**					**747101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.37	GLR 1		
747103	**Very good identification**					**747103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.53	GLR 1		
747111	**Good identification**					**747111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.36	GLR 1		
747113	**Good identification**					**747113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.52	GLR 1		
747301	**Good identification**					**747301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.27	GLR 1		
747303	**Good identification**					**747303**
Staphylococcus schleiferi subsp. coagulans		98.65	0.43	GLR 1		
Staphylococcus hyicus		1.35	-.04			
747313	**Good identification**					**747313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	GLR 1		
747453	**Good identification**					**747453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1		
750240	**Low discrimination**					**750240**
					PYR	
Staphylococcus warneri		54.55	0.33	ORN 1	8	
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	TRE 95	90	
750250	**Low discrimination**					**750250**
Staphylococcus warneri		64.00	0.30	ORN 1		
Staphylococcus pasteuri		36.00	0.10	TRE 95		
750440	**Low discrimination**					**750440**
					PYR	
Staphylococcus warneri		54.55	0.33	ORN 1	8	
Staphylococcus petrasii subsp. croceilyticus		45.45	0.05	SUC 95	90	
750450	**Low discrimination**					**750450**
Staphylococcus warneri		64.00	0.30	ORN 1		
Staphylococcus pasteuri		36.00	0.10	SUC 95		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
750610	**Acceptable identification**					**750610**
	Staphylococcus pasteuri	90.07	0.35	ORN 1		
	Staphylococcus warneri	9.93	0.30	MLT 86		
750612	**Acceptable identification**					**750612**
	Staphylococcus pasteuri	91.74	0.27	ORN 1		
	Staphylococcus warneri	8.26	0.20	MLT 86		
750640	**Low discrimination**					**750640**
					PYR	
	Staphylococcus petrasii subsp. croceilyticus	76.34	0.47	ORN 1	90	
	Staphylococcus warneri	17.58	0.59		8	
	Staphylococcus pasteuri	6.08	0.35		1	
750642	**Low discrimination**					**750642**
					PYR	
	Staphylococcus warneri	64.31	0.49	ORN 1	8	
	Staphylococcus pasteuri	27.23	0.27		1	
	Staphylococcus petrasii subsp. croceilyticus	8.46	0.05		90	
750650	**Low discrimination**					**750650**
					PYR	
	Staphylococcus pasteuri	73.43	0.51	ORN 1	1	
	Staphylococcus warneri	25.05	0.56		8	
	Staphylococcus petrasii subsp. croceilyticus	1.52	0.05		90	
750652	**Low discrimination**					**750652**
	Staphylococcus pasteuri	78.22	0.43	ORN 1		
	Staphylococcus warneri	21.78	0.46			
750740	**Low discrimination**					**750740**
					PYR	
	Staphylococcus warneri	87.81	0.49	ORN 1	8	
	Staphylococcus petrasii subsp. croceilyticus	12.19	0.05		90	
750742	**Good identification**					**750742**
	Staphylococcus warneri	100.0	0.39	ORN 1		
750750	**Acceptable identification**					**750750**
	Staphylococcus warneri	91.43	0.46	ORN 1		
	Staphylococcus pasteuri	8.57	0.10			
750752	**Low discrimination**					**750752**
	Staphylococcus warneri	89.70	0.36	ORN 1		
	Staphylococcus pasteuri	10.31	0.02			
752610	**Good identification**					**752610**
	Staphylococcus pasteuri	98.11	0.43	ORN 1		
	Staphylococcus warneri	1.89	0.22	MLT 86		
752612	**Good identification**					**752612**
					YEP	
	Staphylococcus pasteuri	97.32	0.35	ORN 1	86	
	Staphylococcus warneri	1.53	0.12	MLT 86	43	
	Staphylococcus simulans	1.15	0.00		1	
752640	**Good identification**					**752640**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus petrasii subsp. croceilyticus		93.11	0.59	ORN 1	90	
Staphylococcus pasteuri		4.54	0.43		1	
Staphylococcus warneri		2.30	0.51		8	
752642	**Low discrimination**					**752642**
					PYR	
Staphylococcus pasteuri		51.67	0.35	ORN 1	1	
Staphylococcus petrasii subsp. croceilyticus		26.23	0.18	MAN 86	90	
Staphylococcus warneri		21.36	0.42		8	
752643	**Acceptable identification**					**752643**
Staphylococcus lugdunensis		94.53	0.39	ARG 5		
Staphylococcus petrasii subsp. petrasii		5.47	0.12	GLR 2		
752650	**Good identification**					**752650**
					PYR	
Staphylococcus pasteuri		91.45	0.59	ORN 1	1	
Staphylococcus warneri		5.46	0.49		8	
Staphylococcus petrasii subsp. croceilyticus		3.09	0.18		90	
752652	**Good identification**					**752652**
Staphylococcus pasteuri		95.35	0.51	ORN 1		
Staphylococcus warneri		4.65	0.39			
752742	**Low discrimination**					**752742**
					PYR	
Staphylococcus warneri		84.12	0.31	ORN 1	8	
Staphylococcus lugdunensis		9.19	-.02		96	
Staphylococcus haemolyticus		6.69	0.01		98	
752743	**Good identification**					**752743**
Staphylococcus lugdunensis		96.29	0.39	ARG 5		
Staphylococcus petrasii subsp. petrasii		3.71	0.09	GLR 2		
752750	**Low discrimination**					**752750**
					PYR	
Staphylococcus warneri		63.54	0.38	ORN 1	8	
Staphylococcus pasteuri		34.03	0.18		1	
Staphylococcus haemolyticus		2.43	0.02		98	
752752	**Low discrimination**					**752752**
Staphylococcus warneri		60.38	0.28	ORN 1		
Staphylococcus pasteuri		39.63	0.10			
756341	**Good identification**					**756341**
Staphylococcus epidermidis		100.0	0.31	ORN 12		
756342	**Good identification**					**756342**
Staphylococcus epidermidis		100.0	0.36	ORN 12		
756343	**Good identification**					**756343**
Staphylococcus epidermidis		100.0	0.41	ORN 12		
760040	**Good identification**					**760040**
Staphylococcus warneri		100.0	0.35	SUC 95		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
760050	**Good identification**					**760050**
Staphylococcus warneri		100.0	0.32	SUC 95		
760200	**Good identification**					**760200**
Staphylococcus warneri		100.0	0.35	TRE 95		
760210	**Good identification**					**760210**
Staphylococcus warneri		100.0	0.32	TRE 95		
760240	**Very good identification**					**760240**
					PYR	
Staphylococcus warneri		96.74	0.61	TRE 95	8	
Staphylococcus petrasii subsp. croceilyticus		3.26	0.05		90	
760241	**Good identification**					**760241**
Staphylococcus warneri		100.0	0.27	TRE 95		
760242	**Good identification**					**760242**
Staphylococcus warneri		100.0	0.52	TRE 95		
760250	**Very good identification**					**760250**
Staphylococcus warneri		97.78	0.58	TRE 95		
Staphylococcus pasteuri		2.22	0.10			
760252	**Acceptable identification**					**760252**
					NOV	
Staphylococcus warneri		91.16	0.49	TRE 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		5.27	0.02		99	
Staphylococcus pasteuri		2.54	0.02		1	
760340	**Good identification**					**760340**
Staphylococcus warneri		100.0	0.51	TRE 95		
760342	**Good identification**					**760342**
Staphylococcus warneri		100.0	0.41	TRE 95		
760350	**Good identification**					**760350**
Staphylococcus warneri		100.0	0.48	TRE 95		
760352	**Good identification**					**760352**
Staphylococcus warneri		100.0	0.38	TRE 95		
760400	**Good identification**					**760400**
Staphylococcus warneri		100.0	0.35	SUC 95		
760410	**Good identification**					**760410**
Staphylococcus warneri		100.0	0.32	SUC 95		
760413	**Low discrimination**					**760413**
					NOV	
Staphylococcus cohnii subsp. urealyticum		83.81	0.35	ARG 1	99	
Staphylococcus simulans		16.20	0.15		1	
760440	**Very good identification**					**760440**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus warneri		96.74	0.61	SUC 95	8	
Staphylococcus petrasii subsp. croceilyticus		3.26	0.05		90	
760441	**Low discrimination**					**760441**
					NOV	
Staphylococcus warneri		81.49	0.27	SUC 95	1	
Staphylococcus cohnii subsp. urealyticum		18.51	0.09	MNS 2	99	
760442	**Good identification**					**760442**
					NOV	
Staphylococcus warneri		98.32	0.52	SUC 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		1.68	-.06		99	
760443	**Low discrimination**					**760443**
					NOV	
Staphylococcus cohnii subsp. urealyticum		87.32	0.30	ARG 1	99	
Staphylococcus warneri		12.68	0.17	MAN 91	1	
760450	**Good identification**					**760450**
					NOV	
Staphylococcus warneri		94.70	0.58	SUC 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		3.15	0.07		99	
Staphylococcus pasteuri		2.15	0.10		1	
760451	**Low discrimination**					**760451**
					NOV	
Staphylococcus cohnii subsp. urealyticum		76.03	0.30	ARG 1	99	
Staphylococcus warneri		23.98	0.24	LAC 91	1	
760452	**Low discrimination**					**760452**
					NOV	
Staphylococcus warneri		61.97	0.49	SUC 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		35.08	0.23		99	
Staphylococcus pasteuri		1.73	0.02		1	
760453	**Good identification**					**760453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		97.49	0.51	ARG 1	99	
Staphylococcus simulans		1.49	0.08		1	
Staphylococcus warneri		1.01	0.14		1	
760513	**Good identification**					**760513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.28	ARG 1		
760540	**Good identification**					**760540**
Staphylococcus warneri		100.0	0.51	SUC 95		
760542	**Good identification**					**760542**
Staphylococcus warneri		100.0	0.41	SUC 95		
760550	**Good identification**					**760550**
Staphylococcus warneri		100.0	0.48	SUC 95		
760552	**Low discrimination**					**760552**
					NOV	
Staphylococcus warneri		86.52	0.38	SUC 95	1	
Staphylococcus saprophyticus subsp. saprophyticus		13.49	0.01		99	
760553	**Good identification**					**760553**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus cohnii subsp. urealyticum		100.0	0.43	ARG 1			
760600	**Good identification**						**760600**
					PYR		
Staphylococcus warneri		92.86	0.61	MLT 95	8		
Staphylococcus pasteuri		4.01	0.19		1		
Staphylococcus petrasii subsp. croceilyticus		3.13	0.05		90		
760601	**Good identification**						**760601**
Staphylococcus warneri		100.0	0.27	MLT 95			
760602	**Low discrimination**						**760602**
					YEP		
Staphylococcus warneri		87.74	0.52	MLT 95	43		
Staphylococcus simulans		7.62	0.20		1		
Staphylococcus pasteuri		4.64	0.11		86		
760603	**Acceptable identification**						**760603**
Staphylococcus simulans		93.44	0.31	VPT 4			
Staphylococcus warneri		6.56	0.17	NIT 96			
760610	**Low discrimination**						**760610**
Staphylococcus warneri		73.19	0.58	MLT 95			
Staphylococcus pasteuri		26.81	0.35				
760612	**Low discrimination**						**760612**
					YEP		
Staphylococcus warneri		50.34	0.49	MLT 95	43		
Staphylococcus simulans		24.16	0.32		1		
Staphylococcus pasteuri		22.60	0.27		86		
760613	**Good identification**						**760613**
Staphylococcus simulans		98.75	0.43	VPT 4			
Staphylococcus warneri		1.25	0.14	NIT 96			
760640	**Low discrimination**						**760640**
					PYR		
Staphylococcus warneri		83.99	0.88		8		
Staphylococcus petrasii subsp. croceilyticus		14.73	0.47		90		
Staphylococcus pasteuri		1.17	0.35		1		
760641	**Low discrimination**						**760641**
					PYR	YEP	
Staphylococcus warneri		72.87	0.53	MNS 2	8	43	
Staphylococcus petrasii subsp. petrasii		20.81	0.27		99	1	
Staphylococcus petrasii subsp. croceilyticus		6.33	0.05		90	99	
760642	**Very good identification**						**760642**
					NOV		
Staphylococcus warneri		96.66	0.78		1		
Staphylococcus pasteuri		1.65	0.27		1		
Staphylococcus saprophyticus subsp. saprophyticus		0.88	0.15		99		
760643	**Low discrimination**						**760643**
					PYR		
Staphylococcus warneri		55.61	0.43	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		25.65	0.21		99		
Staphylococcus simulans		18.74	0.24		70		
760650	**Low discrimination**						**760650**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					NOV		
Staphylococcus warneri		87.83	0.85		1		
Staphylococcus pasteuri		10.40	0.51		1		
Staphylococcus saprophyticus subsp. saprophyticus		1.56	0.28		99		
760651	**Acceptable identification**						**760651**
					PYR	YEP	
Staphylococcus warneri		90.71	0.50	MNS 2	8	43	
Staphylococcus pasteuri		5.32	0.10		1	86	
Staphylococcus petrasii subsp. petrasii		3.97	0.07		99	1	
760652	**Low discrimination**						**760652**
					NOV		
Staphylococcus warneri		68.60	0.75		1		
Staphylococcus saprophyticus subsp. saprophyticus		20.66	0.43		99		
Staphylococcus pasteuri		9.96	0.43		1		
760653	**Low discrimination**						**760653**
					NOV		
Staphylococcus simulans		56.24	0.36	VPT 4	1		
Staphylococcus warneri		30.21	0.40	NIT 96	1		
Staphylococcus cohnii subsp. urealyticum		4.73	0.20		99		
760660	**Low discrimination**						**760660**
					PYR		
Staphylococcus warneri		85.08	0.47	XYL 1	8		
Staphylococcus petrasii subsp. croceilyticus		14.92	0.05		90		
760662	**Good identification**						**760662**
Staphylococcus warneri		100.0	0.37	XYL 1			
760670	**Low discrimination**						**760670**
Staphylococcus warneri		89.41	0.44	XYL 1			
Staphylococcus pasteuri		10.59	0.10				
760672	**Low discrimination**						**760672**
					NOV		
Staphylococcus warneri		69.14	0.34	XYL 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		20.83	0.02		99		
Staphylococcus pasteuri		10.03	0.02		1		
760700	**Good identification**						**760700**
Staphylococcus warneri		100.0	0.51	MLT 95			
760702	**Good identification**						**760702**
Staphylococcus warneri		100.0	0.41	MLT 95			
760710	**Good identification**						**760710**
Staphylococcus warneri		100.0	0.48	MLT 95			
760712	**Low discrimination**						**760712**
Staphylococcus warneri		88.33	0.38	MLT 95			
Staphylococcus simulans		11.67	0.11				
760740	**Very good identification**						**760740**
					PYR	YEP	
Staphylococcus warneri		99.31	0.77		8	43	
Staphylococcus petrasii subsp. croceilyticus		0.56	0.05		90	99	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus petrasii subsp. petrasii		0.14	0.04		99	1	
760741	**Low discrimination**						**760741**
					PYR		
Staphylococcus warneri		62.39	0.43	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		37.61	0.23		99		
760742	**Very good identification**						**760742**
					NOV	PYR	
Staphylococcus warneri		99.53	0.68		1	8	
Staphylococcus saprophyticus subsp. saprophyticus		0.25	-.07		99	7	
Staphylococcus petrasii subsp. petrasii		0.22	-.02		1	99	
760743	**Low discrimination**						**760743**
					PYR		
Staphylococcus warneri		48.39	0.33	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		47.12	0.18		99		
Staphylococcus simulans		4.49	0.02		70		
760750	**Very good identification**						**760750**
					NOV		
Staphylococcus warneri		99.14	0.74		1		
Staphylococcus saprophyticus subsp. saprophyticus		0.48	0.06		99		
Staphylococcus pasteuri		0.38	0.10		1		
760751	**Acceptable identification**						**760751**
					PYR		
Staphylococcus warneri		91.53	0.40	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		8.47	0.04		99		
760752	**Good identification**						**760752**
					NOV		
Staphylococcus warneri		91.68	0.65		1		
Staphylococcus saprophyticus subsp. saprophyticus		7.61	0.22		99		
Staphylococcus pasteuri		0.43	0.02		1		
760753	**Low discrimination**						**760753**
					NOV		
Staphylococcus warneri		48.22	0.30	MNS 2	1		
Staphylococcus simulans		24.72	0.14		1		
Staphylococcus cohnii subsp. urealyticum		10.25	0.12		99		
760760	**Good identification**						**760760**
Staphylococcus warneri		100.0	0.36	XYL 1			
760770	**Good identification**						**760770**
Staphylococcus warneri		100.0	0.33	XYL 1			
761413	**Good identification**						**761413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1			
761453	**Good identification**						**761453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.42	ARG 1			
761553	**Good identification**						**761553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.35	ARG 1			
761640	**Low discrimination**						**761640**
					PYR		
Staphylococcus warneri		85.08	0.47	ESL 1	8		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus petrasii subsp. croceilyticus		14.92	0.05		90	
761642	**Good identification**					**761642**
Staphylococcus warneri		100.0	0.37	ESL 1		
761650	**Low discrimination**					**761650**
Staphylococcus warneri		89.41	0.44	ESL 1		
Staphylococcus pasteuri		10.59	0.10			
761652	**Low discrimination**					**761652**
					NOV	
Staphylococcus warneri		69.14	0.34	ESL 1	1	
Staphylococcus saprophyticus subsp. saprophyticus		20.83	0.02		99	
Staphylococcus pasteuri		10.03	0.02		1	
761740	**Good identification**					**761740**
Staphylococcus warneri		100.0	0.36	ESL 1		
761750	**Good identification**					**761750**
Staphylococcus warneri		100.0	0.33	ESL 1		
762040	**Good identification**					**762040**
Staphylococcus warneri		100.0	0.27	SUC 95		
762200	**Good identification**					**762200**
Staphylococcus warneri		100.0	0.27	TRE 95		
762203	**Good identification**					**762203**
Staphylococcus simulans		100.0	0.33	VPT 4		
762212	**Good identification**					**762212**
Staphylococcus simulans		96.41	0.35	VPT 4		
Staphylococcus warneri		3.59	0.15	TRE 95		
762213	**Low discrimination**					**762213**
Staphylococcus simulans		89.91	0.45	VPT 4		
Staphylococcus capitis subsp. ureolyticus		10.09	0.13	TRE 95		
762240	**Low discrimination**					**762240**
					PYR	
Staphylococcus warneri		73.80	0.54	TRE 95	8	
Staphylococcus petrasii subsp. croceilyticus		23.19	0.18		90	
Staphylococcus haemolyticus		1.39	0.11		98	
762242	**Low discrimination**					**762242**
Staphylococcus warneri		82.70	0.44	TRE 95		
Staphylococcus simulans		9.52	0.15			
Staphylococcus epidermidis		5.15	0.10			
762250	**Low discrimination**					**762250**
					PYR	
Staphylococcus warneri		87.32	0.51	TRE 95	8	
Staphylococcus pasteuri		11.34	0.18		1	
Staphylococcus haemolyticus		1.34	0.06		98	
762252	**Low discrimination**					**762252**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					YEP	
Staphylococcus warneri		51.99	0.41	TRE 95	43	
Staphylococcus simulans		33.04	0.27		1	
Staphylococcus pasteuri		8.27	0.10		86	
762253	**Low discrimination**					**762253**
Staphylococcus capitis subsp. ureolyticus		85.71	0.41	BGA 12		
Staphylococcus simulans		14.29	0.38	GLR 1		
762340	**Low discrimination**					**762340**
					PYR	
Staphylococcus warneri		84.39	0.43	TRE 95	8	
Staphylococcus epidermidis		9.92	0.15		1	
Staphylococcus haemolyticus		5.69	0.12		98	
762343	**Acceptable identification**					**762343**
					PYR	
Staphylococcus epidermidis		91.76	0.31	GLR 1	1	
Staphylococcus simulans		4.69	0.04		70	
Staphylococcus petrasii subsp. petrasii		3.54	-.04		99	
762350	**Acceptable identification**					**762350**
					PYR	
Staphylococcus warneri		94.82	0.41	TRE 95	8	
Staphylococcus haemolyticus		5.18	0.07		98	
762352	**Low discrimination**					**762352**
					PYR	
Staphylococcus warneri		78.91	0.31	TRE 95	8	
Staphylococcus simulans		13.81	0.06		70	
Staphylococcus haemolyticus		7.28	0.02		98	
762400	**Good identification**					**762400**
Staphylococcus warneri		100.0	0.27	SUC 95		
762403	**Good identification**					**762403**
Staphylococcus simulans		100.0	0.31	VPT 4		
762412	**Good identification**					**762412**
Staphylococcus simulans		95.51	0.32	VPT 4		
Staphylococcus warneri		4.49	0.15	SUC 96		
762413	**Good identification**					**762413**
					NOV	
Staphylococcus simulans		96.34	0.43	VPT 4	1	
Staphylococcus cohnii subsp. urealyticum		3.67	0.20	SUC 96	99	
762440	**Low discrimination**					**762440**
					PYR	
Staphylococcus warneri		75.21	0.54	SUC 95	8	
Staphylococcus petrasii subsp. croceilyticus		23.63	0.18		90	
Staphylococcus pasteuri		1.15	0.02		1	
762442	**Acceptable identification**					**762442**
Staphylococcus warneri		91.65	0.44	SUC 95		
Staphylococcus simulans		8.35	0.13			
762450	**Low discrimination**					**762450**
					NOV PYR	

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
Staphylococcus warneri		86.34	0.51	SUC 95	1	8	
Staphylococcus pasteuri		11.21	0.18		1	1	
Staphylococcus saprophyticus subsp. bovis		2.45	0.09		99	99	
762452	**Low discrimination**						**762452**
					YEP		
Staphylococcus warneri		60.16	0.41	SUC 95	43		
Staphylococcus simulans		30.27	0.25		1		
Staphylococcus pasteuri		9.57	0.10		86		
762453	**Low discrimination**						**762453**
					NOV		
Staphylococcus simulans		67.57	0.36	VPT 4	1		
Staphylococcus cohnii subsp. urealyticum		32.43	0.35	SUC 96	99		
762540	**Good identification**						**762540**
Staphylococcus warneri		100.0	0.43	SUC 95			
762542	**Good identification**						**762542**
Staphylococcus warneri		100.0	0.34	SUC 95			
762550	**Low discrimination**						**762550**
					NOV	PYR	
Staphylococcus warneri		73.60	0.41	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		26.40	0.22		99	99	
762552	**Low discrimination**						**762552**
					NOV	PYR	
Staphylococcus warneri		71.05	0.31	SUC 95	1	8	
Staphylococcus saprophyticus subsp. bovis		19.11	0.09		99	99	
Staphylococcus simulans		9.84	0.03		1	70	
762553	**Low discrimination**						**762553**
					NOV		
Staphylococcus cohnii subsp. urealyticum		70.29	0.28	ARG 1	99		
Staphylococcus simulans		29.72	0.14		1		
762600	**Low discrimination**						**762600**
					PYR		
Staphylococcus warneri		63.40	0.54	MLT 95	8		
Staphylococcus petrasii subsp. croceilyticus		19.92	0.18		90		
Staphylococcus pasteuri		15.64	0.27		1		
762602	**Low discrimination**						**762602**
					YEP		
Staphylococcus simulans		78.89	0.48	VPT 4	1		
Staphylococcus warneri		16.21	0.44		43		
Staphylococcus pasteuri		4.90	0.19		86		
762603	**Very good identification**						**762603**
					PYR		
Staphylococcus simulans		99.46	0.59	VPT 4	70		
Staphylococcus petrasii subsp. petrasii		0.23	0.00		99		
Staphylococcus hyicus		0.18	-.04		1		
762610	**Low discrimination**						**762610**
					YEP		
Staphylococcus pasteuri		65.74	0.43	BGA 1	86		
Staphylococcus warneri		31.42	0.51	MLT 86	43		
Staphylococcus simulans		2.84	0.20		1		
762611	**Low discrimination**						**762611**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					YEP		
Staphylococcus simulans		87.94	0.31	VPT 4	1		
Staphylococcus pasteuri		6.14	0.02	LAC 99	86		
Staphylococcus warneri		5.92	0.16		43		
762612	**Low discrimination**						**762612**
					YEP		
Staphylococcus simulans		88.29	0.61	VPT 4	1		
Staphylococcus pasteuri		8.42	0.35		86		
Staphylococcus warneri		3.29	0.41		43		
762613	**Very good identification**						**762613**
Staphylococcus simulans		100.0	0.72	VPT 4			
762633	**Good identification**						**762633**
					NOV		
Staphylococcus simulans		95.84	0.31	VPT 4	1		
Staphylococcus xylosus		4.16	-.01	XYL 1	99		
762640	**Low discrimination**						**762640**
					PYR		
Staphylococcus petrasii subsp. croceilyticus		59.66	0.59	BGA 1	90		
Staphylococcus warneri		36.44	0.80		8		
Staphylococcus pasteuri		2.91	0.43		1		
762641	**Low discrimination**						**762641**
					PYR	YEP	
Staphylococcus petrasii subsp. petrasii		75.62	0.47	GLR 1	99	1	
Staphylococcus warneri		12.61	0.45		8	43	
Staphylococcus petrasii subsp. croceilyticus		10.22	0.18		90	99	
762642	**Low discrimination**						**762642**
					YEP		
Staphylococcus warneri		76.53	0.70		43		
Staphylococcus simulans		8.81	0.41		1		
Staphylococcus pasteuri		7.48	0.35		86		
762643	**Low discrimination**						**762643**
					PYR		
Staphylococcus simulans		63.85	0.52	VPT 4	70		
Staphylococcus petrasii subsp. petrasii		32.77	0.41		99		
Staphylococcus warneri		3.38	0.35		8		
762650	**Low discrimination**						**762650**
					PYR		
Staphylococcus warneri		58.13	0.77		8		
Staphylococcus pasteuri		39.33	0.59		1		
Staphylococcus petrasii subsp. croceilyticus		1.33	0.18		90		
762651	**Low discrimination**						**762651**
					PYR		
Staphylococcus warneri		37.33	0.42	MNS 2	8		
Staphylococcus petrasii subsp. petrasii		34.34	0.27		99		
Staphylococcus simulans		13.10	0.23		70		
762652	**Low discrimination**						**762652**
					YEP		
Staphylococcus warneri		40.04	0.67		43		
Staphylococcus pasteuri		33.19	0.51		86		
Staphylococcus simulans		25.45	0.54		1		
762653	**Very good identification**						**762653**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
					PYR	
Staphylococcus simulans		96.89	0.64	VPT 4	70	
Staphylococcus petrasii subsp. petrasii		1.38	0.21		99	
Staphylococcus warneri		0.93	0.33		8	
762662	**Low discrimination**					**762662**
Staphylococcus warneri		89.68	0.29	XYL 1		
Staphylococcus simulans		10.32	0.00			
762670	**Low discrimination**					**762670**
Staphylococcus warneri		59.65	0.36	XYL 1		
Staphylococcus pasteuri		40.35	0.18			
762700	**Good identification**					**762700**
Staphylococcus warneri		100.0	0.43	MLT 95		
762702	**Low discrimination**					**762702**
Staphylococcus simulans		57.26	0.27	VPT 4		
Staphylococcus warneri		42.74	0.34	GAL 8		
762703	**Low discrimination**					**762703**
					PYR	
Staphylococcus simulans		79.61	0.37	VPT 4	70	
Staphylococcus hyicus		18.97	0.17	GAL 8	1	
Staphylococcus petrasii subsp. petrasii		1.42	-.04		99	
762710	**Acceptable identification**					**762710**
Staphylococcus warneri		93.73	0.41	MLT 95		
Staphylococcus pasteuri		6.27	0.02			
762712	**Low discrimination**					**762712**
Staphylococcus simulans		88.10	0.39	VPT 4		
Staphylococcus warneri		11.90	0.31	GAL 8		
762713	**Good identification**					**762713**
					NOV	
Staphylococcus simulans		99.56	0.50	VPT 4	1	
Staphylococcus xylosus		0.44	-.02	GAL 8	99	
762740	**Low discrimination**					**762740**
					PYR	
Staphylococcus warneri		88.44	0.70		8	
Staphylococcus petrasii subsp. croceilyticus		4.63	0.18		90	
Staphylococcus haemolyticus		4.17	0.35		98	
762741	**Acceptable identification**					**762741**
					PYR	
Staphylococcus petrasii subsp. petrasii		92.20	0.43	GLR 1	99	
Staphylococcus warneri		7.28	0.35		8	
Staphylococcus haemolyticus		0.52	0.04		98	
762742	**Low discrimination**					**762742**
					PYR	
Staphylococcus warneri		86.39	0.60		8	
Staphylococcus haemolyticus		6.87	0.30		98	
Staphylococcus petrasii subsp. petrasii		4.01	0.18		99	
762743	**Low discrimination**					**762743**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					PYR		
Staphylococcus petrasii subsp. petrasii		76.25	0.37	GLR 1	99		
Staphylococcus simulans		19.38	0.30		70		
Staphylococcus warneri		3.73	0.25		8		
762750	**Low discrimination**						**762750**
					NOV	PYR	
Staphylococcus warneri		87.74	0.67		1	8	
Staphylococcus saprophyticus subsp. bovis		6.63	0.34		99	99	
Staphylococcus haemolyticus		3.35	0.31		2	98	
762752	**Low discrimination**						**762752**
					PYR		
Staphylococcus warneri		75.20	0.57		8		
Staphylococcus simulans		13.16	0.32		70		
Staphylococcus haemolyticus		4.85	0.25		98		
762753	**Low discrimination**						**762753**
					NOV		
Staphylococcus simulans		74.64	0.43	VPT 4	1		
Staphylococcus xylosus		9.79	0.21	GAL 8	99		
Staphylococcus petrasii subsp. petrasii		8.16	0.18		1		
762760	**Good identification**						**762760**
Staphylococcus warneri		100.0	0.29	XYL 1			
762771	**Good identification**						**762771**
Staphylococcus xylosus		100.0	0.33	VPT 14			
762773	**Good identification**						**762773**
					NOV		
Staphylococcus xylosus		99.34	0.43	VPT 14	99		
Staphylococcus simulans		0.67	0.02	ARG 1	1		
763453	**Good identification**						**763453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.27	ARG 1			
763613	**Good identification**						**763613**
Staphylococcus simulans		100.0	0.31	VPT 4			
763642	**Low discrimination**						**763642**
Staphylococcus warneri		89.68	0.29	ESL 1			
Staphylococcus simulans		10.32	0.00				
763650	**Low discrimination**						**763650**
Staphylococcus warneri		59.65	0.36	ESL 1			
Staphylococcus pasteuri		40.35	0.18				
763740	**Good identification**						**763740**
Staphylococcus warneri		100.0	0.29	ESL 1			
763773	**Low discrimination**						**763773**
Staphylococcus xylosus		89.62	0.30	VPT 14			
Staphylococcus equorum		10.38	0.12	ARG 1			
764413	**Good identification**						**764413**
					NOV		

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus cohnii subsp. urealyticum		95.88	0.39	ARG 1	99	
Staphylococcus simulans		4.12	0.05		1	
764443	**Good identification**					**764443**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	ARG 1		
764450	**Low discrimination**					**764450**
					NOV	
Staphylococcus kloosii		55.68	0.27	VPT 10	99	
Staphylococcus warneri		44.32	0.17	ARG 1	1	
764451	**Good identification**					**764451**
Staphylococcus cohnii subsp. urealyticum		100.0	0.34	ARG 1		
764452	**Low discrimination**					**764452**
Staphylococcus kloosii		58.10	0.28	VPT 10		
Staphylococcus cohnii subsp. urealyticum		41.90	0.13	ARG 1		
764453	**Very good identification**					**764453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.54	ARG 1		
764513	**Good identification**					**764513**
Staphylococcus cohnii subsp. urealyticum		100.0	0.31	ARG 1		
764550	**Good identification**					**764550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
764552	**Low discrimination**					**764552**
Staphylococcus kloosii		86.26	0.34	VPT 10		
Staphylococcus cohnii subsp. urealyticum		13.74	0.06	ARG 1		
764553	**Good identification**					**764553**
					NOV	
Staphylococcus cohnii subsp. urealyticum		97.32	0.47	ARG 1	99	
Staphylococcus caprae		2.68	-.02		1	
764613	**Good identification**					**764613**
					NOV	
Staphylococcus simulans		97.09	0.33	VPT 4	1	
Staphylococcus cohnii subsp. urealyticum		2.91	0.08	NIT 96	99	
764640	**Low discrimination**					**764640**
					PYR	
Staphylococcus warneri		85.08	0.47	PHS 1	8	
Staphylococcus petrasii subsp. croceilyticus		14.92	0.05		90	
764642	**Acceptable identification**					**764642**
Staphylococcus warneri		93.65	0.37	PHS 1		
Staphylococcus simulans		6.35	0.03			
764650	**Low discrimination**					**764650**
Staphylococcus warneri		89.41	0.44	PHS 1		
Staphylococcus pasteuri		10.59	0.10			
764652	**Low discrimination**					**764652**
					NOV	
Staphylococcus warneri		54.92	0.34	PHS 1	1	

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		20.57	0.15		1	
Staphylococcus saprophyticus subsp. saprophyticus		16.54	0.02		99	
764740	**Good identification**					**764740**
Staphylococcus warneri		100.0	0.36	PHS 1		
764750	**Good identification**					**764750**
Staphylococcus warneri		100.0	0.33	PHS 1		
765413	**Good identification**					**765413**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ARG 1		
765450	**Good identification**					**765450**
Staphylococcus kloosii		100.0	0.27	VPT 10		
765452	**Low discrimination**					**765452**
Staphylococcus kloosii		78.10	0.28	VPT 10		
Staphylococcus cohnii subsp. urealyticum		21.91	0.05	ARG 1		
765453	**Good identification**					**765453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.46	ARG 1		
765550	**Good identification**					**765550**
Staphylococcus kloosii		100.0	0.33	VPT 10		
765552	**Good identification**					**765552**
Staphylococcus kloosii		100.0	0.34	VPT 10		
765553	**Good identification**					**765553**
Staphylococcus cohnii subsp. urealyticum		100.0	0.38	ARG 1		
766101	**Good identification**					**766101**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.90	0.43	GLR 1	99	
Staphylococcus schleiferi subsp. schleiferi		3.10	0.08	LAC 86	1	
766103	**Very good identification**					**766103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	GLR 1		
766111	**Good identification**					**766111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	GLR 1		
766113	**Very good identification**					**766113**
					COA	
Staphylococcus schleiferi subsp. coagulans		99.57	0.58	GLR 1	99	
Staphylococcus carnosus subsp. carnosus		0.43	0.05		1	
766213	**Good identification**					**766213**
					COA	
Staphylococcus simulans		96.60	0.35	VPT 4	1	
Staphylococcus schleiferi subsp. coagulans		3.41	0.07	TRE 95	99	
766243	**Low discrimination**					**766243**
Staphylococcus epidermidis		83.61	0.31	GLR 1		

Code	Taxon	ID score	T index	tests against	Differentiation tests	Code
Staphylococcus simulans		16.39	0.16			
766253	**Low discrimination**					**766253**
Staphylococcus simulans		84.62	0.28	VPT 4		
Staphylococcus capitis subsp. ureolyticus		15.38	0.00	TRE 95		
766301	**Good identification**					**766301**
Staphylococcus schleiferi subsp. coagulans		95.99	0.32	GLR 1		
Staphylococcus hyicus		4.01	-.04	LAC 86		
766303	**Acceptable identification**					**766303**
					COA	
Staphylococcus schleiferi subsp. coagulans		91.53	0.49	GLR 1	99	
Staphylococcus hyicus		7.15	0.17		24	
Staphylococcus epidermidis		0.79	0.05		1	
766311	**Good identification**					**766311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	GLR 1		
766313	**Good identification**					**766313**
					COA	
Staphylococcus schleiferi subsp. coagulans		96.99	0.48	GLR 1	99	
Staphylococcus simulans		2.42	0.14		1	
Staphylococcus intermedius		0.59	-.01		99	
766340	**Good identification**					**766340**
Staphylococcus epidermidis		100.0	0.31	GLR 1		
766341	**Good identification**					**766341**
Staphylococcus epidermidis		100.0	0.36	GLR 1		
766342	**Good identification**					**766342**
Staphylococcus epidermidis		100.0	0.41	GLR 1		
766343	**Good identification**					**766343**
					COA	
Staphylococcus epidermidis		98.84	0.47	GLR 1	1	
Staphylococcus schleiferi subsp. coagulans		1.16	0.08		99	
766413	**Low discrimination**					**766413**
					NOV	
Staphylococcus simulans		74.38	0.33	VPT 4	1	
Staphylococcus cohnii subsp. urealyticum		12.74	0.23	SUC 96	99	
Staphylococcus carnosus subsp. carnosus		7.68	0.10		1	
766453	**Low discrimination**					**766453**
					NOV	
Staphylococcus cohnii subsp. urealyticum		68.35	0.39	ARG 1	99	
Staphylococcus simulans		31.65	0.26		1	
766602	**Good identification**					**766602**
Staphylococcus simulans		100.0	0.39	VPT 4		
766603	**Acceptable identification**					**766603**
Staphylococcus simulans		93.83	0.49	VPT 4		
Staphylococcus hyicus		5.83	0.17			
Staphylococcus felis		0.33	-.10			

Code	Taxon	ID score	T index	tests against	Differentiation tests		Code
766612	**Good identification**						**766612**
Staphylococcus simulans		100.0	0.51	VPT 4			
766613	**Very good identification**						**766613**
Staphylococcus simulans		99.41	0.62	VPT 4			
Staphylococcus felis		0.59	0.07				
766642	**Low discrimination**						**766642**
Staphylococcus simulans		79.15	0.31	VPT 4			
Staphylococcus warneri		20.85	0.29				
766643	**Good identification**						**766643**
Staphylococcus simulans		98.47	0.42	VPT 4			
Staphylococcus petrasii subsp. petrasii		1.53	0.00				
766650	**Low discrimination**						**766650**
					NOV	PYR	
Staphylococcus warneri		52.78	0.36	PHS 1	1	8	
Staphylococcus pasteuri		35.71	0.18		1	1	
Staphylococcus saprophyticus subsp. bovis		7.79	0.09		99	99	
766652	**Acceptable identification**						**766652**
					YEP		
Staphylococcus simulans		91.98	0.44	VPT 4	1		
Staphylococcus warneri		4.39	0.26		43		
Staphylococcus pasteuri		3.64	0.10		86		
766653	**Very good identification**						**766653**
					COA	NOV	
Staphylococcus simulans		97.66	0.55	VPT 4	1	1	
Staphylococcus aureus subsp. aureus		2.07	0.05		99	4	
Staphylococcus cohnii subsp. urealyticum		0.27	0.08		1	99	
766703	**Low discrimination**						**766703**
					COA	PYR	
Staphylococcus hyicus		82.95	0.39	VPT 1	24	1	
Staphylococcus simulans		10.09	0.28	BGA 10	1	70	
Staphylococcus intermedius		2.56	0.14		99	90	
766712	**Good identification**						**766712**
Staphylococcus simulans		100.0	0.29	VPT 4			
766713	**Low discrimination**						**766713**
Staphylococcus simulans		53.06	0.40	VPT 4			
Staphylococcus condimenti		19.86	0.18	GAL 8			
Staphylococcus felis		12.10	0.18				
766740	**Good identification**						**766740**
Staphylococcus warneri		100.0	0.29	PHS 1			
766753	**Low discrimination**						**766753**
					COA	PYR	
Staphylococcus aureus subsp. aureus		86.00	0.39	BGA 10	99	10	
Staphylococcus simulans		7.19	0.33	GLR 1	1	70	
Staphylococcus intermedius		4.11	0.26		99	90	
766773	**Acceptable identification**						**766773**

Code Taxon		ID score	T index	tests against	Differentiation tests		Code
					COA	NOV	
Staphylococcus xylosus		91.36	0.31	VPT 14	1	99	
Staphylococcus aureus subsp. aureus		8.64	-.02	ARG 1	99	4	
767101	**Good identification**						**767101**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	GLR 1			
767103	**Very good identification**						**767103**
Staphylococcus schleiferi subsp. coagulans		100.0	0.59	GLR 1			
767111	**Good identification**						**767111**
Staphylococcus schleiferi subsp. coagulans		100.0	0.42	GLR 1			
767113	**Very good identification**						**767113**
Staphylococcus schleiferi subsp. coagulans		100.0	0.58	GLR 1			
767301	**Good identification**						**767301**
Staphylococcus schleiferi subsp. coagulans		100.0	0.32	GLR 1			
767303	**Good identification**						**767303**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	GLR 1			
767311	**Good identification**						**767311**
Staphylococcus schleiferi subsp. coagulans		100.0	0.31	GLR 1			
767313	**Good identification**						**767313**
Staphylococcus schleiferi subsp. coagulans		100.0	0.48	GLR 1			
767453	**Good identification**						**767453**
Staphylococcus cohnii subsp. urealyticum		100.0	0.30	ARG 1			
770640	**Low discrimination**						**770640**
					PYR		
Staphylococcus warneri		85.08	0.47	ORN 1	8		
Staphylococcus petrasii subsp. croceilyticus		14.92	0.05		90		
770642	**Good identification**						**770642**
Staphylococcus warneri		100.0	0.37	ORN 1			
770650	**Low discrimination**						**770650**
Staphylococcus warneri		89.41	0.44	ORN 1			
Staphylococcus pasteuri		10.59	0.10				
770652	**Low discrimination**						**770652**
					NOV		
Staphylococcus warneri		69.14	0.34	ORN 1	1		
Staphylococcus saprophyticus subsp. saprophyticus		20.83	0.02		99		
Staphylococcus pasteuri		10.03	0.02		1		
770740	**Good identification**						**770740**
Staphylococcus warneri		100.0	0.36	ORN 1			
770750	**Good identification**						**770750**

Code Taxon		ID score	T index	tests against	Differentiation tests	Code
Staphylococcus warneri		100.0	0.33	ORN 1		
772613	**Good identification**					**772613**
Staphylococcus simulans		100.0	0.31	VPT 4		
772642	**Low discrimination**					**772642**
Staphylococcus warneri		89.68	0.29	ORN 1		
Staphylococcus simulans		10.32	0.00			
772643	**Low discrimination**					**772643**
Staphylococcus lugdunensis		85.43	0.27	ARG 5		
Staphylococcus simulans		9.63	0.11	GLR 2		
Staphylococcus petrasii subsp. petrasii		4.94	0.00			
772650	**Low discrimination**					**772650**
Staphylococcus warneri		59.65	0.36	ORN 1		
Staphylococcus pasteuri		40.35	0.18			
772740	**Good identification**					**772740**
Staphylococcus warneri		100.0	0.29	ORN 1		
772743	**Good identification**					**772743**
Staphylococcus lugdunensis		96.29	0.27	ARG 5		
Staphylococcus petrasii subsp. petrasii		3.71	-.04	GLR 2		
776343	**Good identification**					**776343**
Staphylococcus epidermidis		100.0	0.29	ORN 12		

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