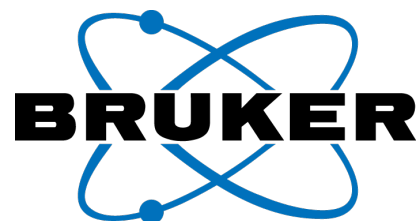


Bruker MALDI Biotyper Identification Results



Run Info:

Run Identifier: 210115-1234-10184
Comment:
Operator: maldi@FLEX-PC
Run Creation Date/Time: 2021-01-15T12:52:19.497
Number of Tests: 26
Type: Standard
BTS-QC: not present
BTS-QC Position:
Instrument ID: 269944.01064
Server Version: 4.1.100 (PYTH) 174 2019-06-158_01-16-09

Result Overview

Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
D5 (+++)(A)	* A 713 (Standard)	Burkholderia gladioli	2.35	Burkholderia gladioli	2.34
D6 (+++)(A)	* A 573 (Standard)	Klebsiella pneumoniae	2.51	Klebsiella pneumoniae	2.43
D7 (+++)(A)	* A 775 (Standard)	Klebsiella pneumoniae	2.24	Klebsiella pneumoniae	2.18
D8 (+++)(A)	* A 855 (Standard)	Klebsiella pneumoniae	2.41	Klebsiella pneumoniae	2.38
D9 (+++)(A)	* B 662 (Standard)	Klebsiella pneumoniae	2.37	Klebsiella pneumoniae	2.35
D10 (+++)(A)	* B 679 (Standard)	Klebsiella pneumoniae	2.32	Klebsiella pneumoniae	2.25
D11 (+++)(A)	* B 723 (Standard)	Klebsiella pneumoniae	2.31	Klebsiella pneumoniae	2.25

Result overview table--continued on next page

<i>Result overview table--continued from previous page</i>					
Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
D12 (+++)(A)	* B 743 (Standard)	Klebsiella pneumoniae	2.30	Klebsiella pneumoniae	2.25
E1 (+++)(A)	* B 797 (Standard)	Klebsiella pneumoniae	2.41	Klebsiella pneumoniae	2.32
E2 (+++)(A)	* C 840 (Standard)	Klebsiella pneumoniae	2.22	Klebsiella pneumoniae	2.10
E3 (+++)(A)	* C 898 (Standard)	Klebsiella pneumoniae	2.11	Klebsiella variicola	1.99
E4 (+++)(A)	* C 984 (Standard)	Klebsiella variicola	2.27	Klebsiella variicola	2.06
E5 (+++)(A)	* C 913 (Standard)	Klebsiella pneumoniae	2.31	Klebsiella pneumoniae	2.30
E6 (+++)(A)	* C 871 (Standard)	Klebsiella variicola	2.26	Klebsiella variicola	2.24
E7 (+++)(A)	* C 863 (Standard)	Klebsiella variicola	2.42	Klebsiella variicola	2.39
E8 (+)(B)	* C 935 (Standard)	Enterobacter kobei	1.97	Enterobacter bugandensis	1.94
E9 (+++)(A)	* C 926 (Standard)	Klebsiella pneumoniae	2.15	Klebsiella pneumoniae	2.11
E10 (+)(B)	* C 980 (Standard)	Acinetobacter baumannii	1.92	Acinetobacter baumannii	1.85
E11 (+++)(A)	* C 1001 (Standard)	Klebsiella pneumoniae	2.29	Klebsiella pneumoniae	2.20
E12 (+++)(A)	* C 1008 (Standard)	Klebsiella pneumoniae	2.27	Klebsiella pneumoniae	2.18
F1 (+++)(A)	* C 1003 (Standard)	Klebsiella pneumoniae	2.45	Klebsiella pneumoniae	2.41
F2 (+++)(A)	* C 1039 (Standard)	Klebsiella pneumoniae	2.51	Klebsiella pneumoniae	2.39
F3 (+++)(A)	* C 1046 (Standard)	Klebsiella variicola	2.19	Klebsiella variicola	2.13
<i>Result overview table--continued on next page</i>					

<i>Result overview table--continued from previous page</i>					
Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
F4 (+++)(A)	* C 1077 (Standard)	Klebsiella variicola	2.31	Klebsiella variicola	2.28
F5 (+++)(A)	* C 1075 (Standard)	Klebsiella pneumoniae	2.31	Klebsiella pneumoniae	2.26
F6 (+++)(A)	* C 117 (Standard)	Klebsiella pneumoniae	2.22	Klebsiella pneumoniae	2.15

Matching Hints

Matched Pattern	Comment
Acinetobacter baumannii 13101_1 CHB	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii B389 UFL	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii DSM 30007T HAM	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii DSM 30007T_QC DSM	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii DSM 30008 DSM	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii DSM 30011 DSM	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii ECII RUH_134 HCB	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii GEIH_2000 HCB	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii LMG 994 HAM	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter pittii 20 HCB	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Burkholderia cenocepacia LMG 12614 HAM	is a member of Burkholderia cepacia complex
Enterobacter asburiae DSM 17506T DSM	is a member of Enterobacter cloacae complex
Enterobacter bugandensis DSM 101086 DSM	is a member of Enterobacter cloacae complex
Enterobacter bugandensis DSM 101087 DSM	is a member of Enterobacter cloacae complex
Enterobacter bugandensis DSM 101089 DSM	is a member of Enterobacter cloacae complex
Enterobacter bugandensis DSM 101090 DSM	is a member of Enterobacter cloacae complex
<i>Matching Hints table--continued on next page</i>	

<i>Matching Hints table--continued from previous page</i>	
Matched Pattern	Comment
Enterobacter bugandensis DSM 101092 DSM	is a member of Enterobacter cloacae complex
Enterobacter bugandensis DSM 29888T DSM	is a member of Enterobacter cloacae complex
Enterobacter cloacae MB11506_1 CHB	is a member of Enterobacter cloacae complex
Enterobacter cloacae MB_5277_05 THL	is a member of Enterobacter cloacae complex
Enterobacter kobei DSM 13645T DSM	is a member of Enterobacter cloacae complex
Klebsiella pneumoniae 37585 PFM	closely related to Klebsiella variicola
Klebsiella pneumoniae 37595 PFM	closely related to Klebsiella variicola
Klebsiella pneumoniae RV_BA_03_B LBK	closely related to Klebsiella variicola
Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	closely related to Klebsiella variicola
Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	closely related to Klebsiella variicola
Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	closely related to Klebsiella variicola
Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	closely related to Klebsiella variicola
Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	closely related to Klebsiella variicola
Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	closely related to Klebsiella variicola
Klebsiella variicola 150416_01 BRUS	closely related to Klebsiella pneumoniae
<i>Matching Hints table--continued on next page</i>	

<i>Matching Hints table--continued from previous page</i>	
Matched Pattern	Comment
Klebsiella variicola 150519_01 SAH	closely related to Klebsiella pneumoniae
Klebsiella variicola 150519_03 SAH	closely related to Klebsiella pneumoniae
Klebsiella variicola 37924 PFM	closely related to Klebsiella pneumoniae
Klebsiella variicola CC_2053 MCW	closely related to Klebsiella pneumoniae
Klebsiella variicola CC_2123 MCW	closely related to Klebsiella pneumoniae
Klebsiella variicola DSM 15968T DSM	closely related to Klebsiella pneumoniae
Klebsiella variicola DSM 15968T HAM	closely related to Klebsiella pneumoniae
Klebsiella variicola NY_1036 MCW	closely related to Klebsiella pneumoniae
Klebsiella variicola NY_1064 MCW	closely related to Klebsiella pneumoniae
Klebsiella variicola NY_1125 MCW	closely related to Klebsiella pneumoniae
Klebsiella variicola TC_4131 MCW	closely related to Klebsiella pneumoniae

Meaning of Score Values

Range	Interpretation	Symbols	Color
2.00 - 3.00	High-confidence identification	(+++)	green
1.70 - 1.99	Low-confidence identification	(+)	yellow
0.00 - 1.69	No Organism Identification Possible	(-)	red

Meaning of Consistency Categories (A - C)

Category	Interpretation
(A)	High consistency: The best match is a high-confidence identification. The second-best match is (1) a high-confidence identification in which the species is identical to the best match, (2) a low-confidence identification in which the species or genus is identical to the best match, or (3) a non-identification.
(B)	Low consistency: The requirements for high consistency are not met. The best match is a high- or low-confidence identification. The second-best match is (1) a high- or low-confidence identification in which the genus is identical to the best match or (2) a non-identification.
(C)	No consistency: The requirements for high or low consistency are not met.

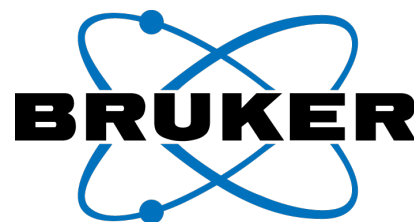
Sample 1



Sample Name: D5
Sample Description:
Sample ID: * A 713
Sample Creation Date/Time: 2021-01-15T12:38:13.789
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

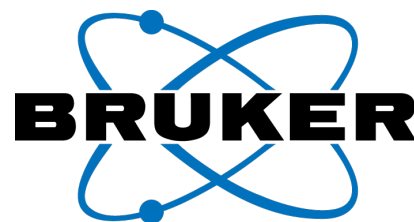
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Burkholderia gladioli DSM 4285T DSM	<u>2.35</u>	<u>28095</u>
2 (+++)	Burkholderia gladioli LMG 2121 LMG	<u>2.34</u>	<u>28095</u>
3 (+++)	Burkholderia gladioli DSM 8361 DSM	<u>2.32</u>	<u>28095</u>
4 (+++)	Burkholderia gladioli DSM 11318 DSM	<u>2.30</u>	<u>28095</u>
5 (+++)	Burkholderia gladioli Wv22575 CHB	<u>2.30</u>	<u>28095</u>
6 (+++)	Burkholderia gladioli LMG 6880 LMG	<u>2.10</u>	<u>28095</u>
7 (+)	Burkholderia gladioli LMG 18159 LMG	<u>1.96</u>	<u>28095</u>
8 (+)	Burkholderia glumae DSM 9512T HAM	<u>1.94</u>	<u>337</u>
9 (+)	Burkholderia gladioli BK1959 EVM	<u>1.88</u>	<u>28095</u>
10 (+)	<u>Burkholderia cenocepacia LMG 12614 HAM</u>	<u>1.86</u>	<u>95486</u>

Sample 2



Sample Name: D6
Sample Description:
Sample ID: * A 573
Sample Creation Date/Time: 2021-01-15T12:38:13.790
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

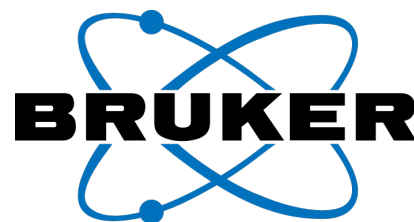
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.51	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.43	72407
3 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.33	573
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.29	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.26	574
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.19	574
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.17	72407
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.13	573
9 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	2.03	39831
10 (+)	Klebsiella variicola 150519_03 SAH	1.94	244366

Sample 3

Sample Name: D7
Sample Description:
Sample ID: * A 775
Sample Creation Date/Time: 2021-01-15T12:38:13.790
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.24	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.18	72407
3 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.11	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.04	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.03	72407
6 (+)	Klebsiella pneumoniae 37585 PFM	1.97	573
7 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	1.97	574
8 (+)	Klebsiella variicola DSM 15968T DSM	1.94	244366
9 (+)	Klebsiella pneumoniae 37595 PFM	1.88	573
10 (+)	Klebsiella variicola 150519_03 SAH	1.87	244366

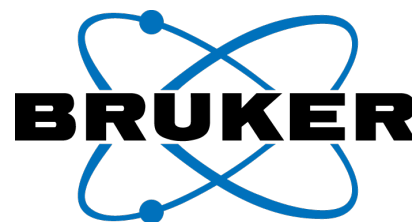
Sample 4



Sample Name: D8
Sample Description:
Sample ID: * A 855
Sample Creation Date/Time: 2021-01-15T12:38:13.791
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.41	574
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.38	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.34	72407
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.28	72407
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.28	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.28	574
7 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.27	573
8 (+++)	Klebsiella variicola 150519_03 SAH	2.09	244366
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.08	244366
10 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	2.05	39831

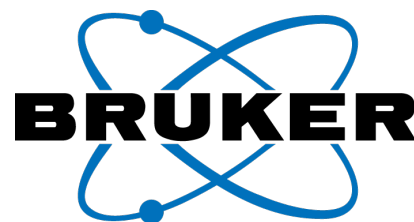
Sample 5



Sample Name: D9
Sample Description:
Sample ID: * B 662
Sample Creation Date/Time: 2021-01-15T12:38:13.791
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.37	574
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.35	72407
3 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.26	573
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.25	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.25	574
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.24	72407
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.22	72407
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	2.19	39831
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.14	244366
10 (+++)	Klebsiella variicola 37924 PFM	2.09	244366

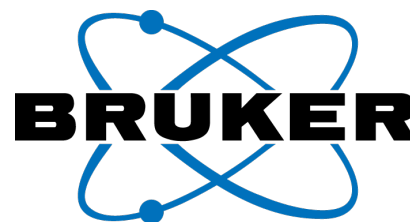
Sample 6



Sample Name: D10
Sample Description:
Sample ID: * B 679
Sample Creation Date/Time: 2021-01-15T12:38:13.792
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

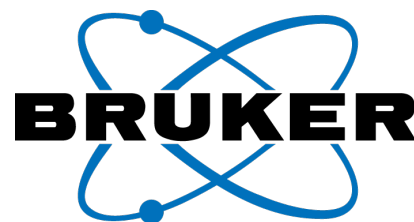
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.32	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.25	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.24	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.19	574
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.16	573
6 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.15	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.11	72407
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.07	244366
9 (+++)	Klebsiella pneumoniae 37595 PFM	2.02	573
10 (+++)	Klebsiella variicola 150519_03 SAH	2.02	244366

Sample 7



Sample Name: D11
Sample Description:
Sample ID: * B 723
Sample Creation Date/Time: 2021-01-15T12:38:13.792
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.31	574
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.25	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.24	72407
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.09	573
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.09	573
6 (+++)	Klebsiella pneumoniae 37595 PFM	2.01	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.01	72407
8 (+++)	Klebsiella variicola CC_2053 MCW	2.00	244366
9 (+)	Klebsiella variicola NY_1064 MCW	1.99	244366
10 (+)	Klebsiella variicola TC_4131 MCW	1.97	244366

Sample 8

Sample Name: D12
Sample Description:
Sample ID: * B 743
Sample Creation Date/Time: 2021-01-15T12:38:13.793
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.30	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.25	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.23	72407
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.22	573
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.20	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.19	574
7 (+++)	Klebsiella pneumoniae 37595 PFM	2.17	573
8 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.14	72407
9 (+++)	Klebsiella variicola 150519_03 SAH	2.04	244366
10 (+++)	Klebsiella variicola 150416_01 BRUS	2.04	244366

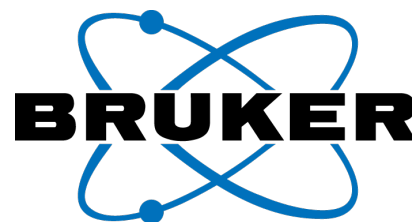
Sample 9



Sample Name: E1
Sample Description:
Sample ID: * B 797
Sample Creation Date/Time: 2021-01-15T12:38:13.793
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.41	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.32	573
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.30	72407
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.20	72407
5 (+++)	Klebsiella pneumoniae 37585_PFM	2.14	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.14	574
7 (+++)	Klebsiella variicola CC_2053_MCW	2.11	244366
8 (+++)	Klebsiella variicola 150416_01_BRUS	2.08	244366
9 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.08	574
10 (+++)	Klebsiella variicola 150519_03_SAH	2.08	244366

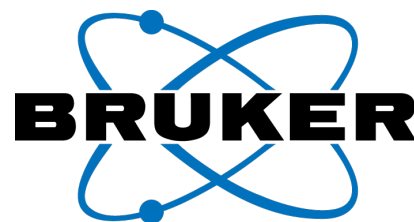
Sample 10



Sample Name: E2
Sample Description:
Sample ID: * C 840
Sample Creation Date/Time: 2021-01-15T12:38:13.794
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.22	574
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.10	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.04	72407
4 (+)	Klebsiella pneumoniae RV_BA_03_B LBK	1.94	573
5 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	1.93	574
6 (+)	Klebsiella pneumoniae 37585 PFM	1.90	573
7 (+)	Klebsiella variicola 150416_01 BRUS	1.88	244366
8 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.87	72407
9 (+)	Klebsiella variicola 150519_03 SAH	1.84	244366
10 (+)	Klebsiella variicola CC_2053 MCW	1.84	244366

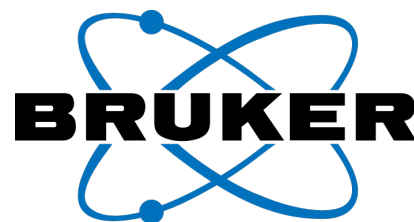
Sample 11



Sample Name: E3
Sample Description:
Sample ID: * C 898
Sample Creation Date/Time: 2021-01-15T12:38:13.794
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.11	573
2 (+)	Klebsiella variicola DSM 15968T HAM	1.99	244366
3 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	1.93	574
4 (+)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	1.89	72407
5 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	1.88	72407
6 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	1.85	72407
7 (+)	Klebsiella variicola DSM 15968T_DSM	1.79	244366
8 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	1.78	574
9 (+)	Klebsiella pneumoniae 37585_PFM	1.70	573
10 (-)	Klebsiella variicola 150519_03_SAH	1.64	244366

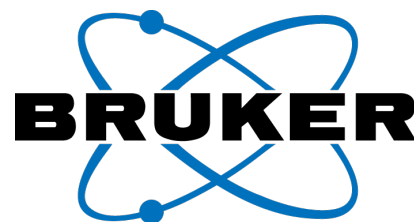
Sample 12



Sample Name: E4
Sample Description:
Sample ID: * C 984
Sample Creation Date/Time: 2021-01-15T12:38:13.795
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola 150416_01 BRUS	2.27	244366
2 (+++)	Klebsiella variicola CC_2053 MCW	2.06	244366
3 (+++)	Klebsiella variicola DSM_15968T DSM	2.04	244366
4 (+++)	Klebsiella variicola NY_1064 MCW	2.04	244366
5 (+++)	Klebsiella variicola TC_4131 MCW	2.03	244366
6 (+++)	Klebsiella variicola NY_1125 MCW	2.02	244366
7 (+++)	Klebsiella variicola CC_2123 MCW	2.01	244366
8 (+++)	Klebsiella variicola 150519_03 SAH	2.01	244366
9 (+)	Klebsiella pneumoniae ssp pneumoniae DSM_30104T_QC DSM	1.98	72407
10 (+)	Klebsiella pneumoniae ssp ozaenae DSM_16358T DSM	1.94	574

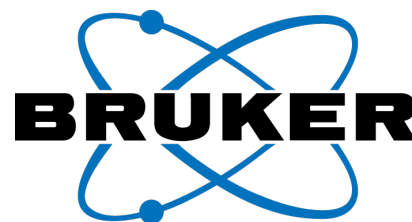
Sample 13



Sample Name: E5
Sample Description:
Sample ID: * C 913
Sample Creation Date/Time: 2021-01-15T12:38:13.795
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.31	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.30	72407
3 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.29	573
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.17	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.16	574
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.12	574
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.06	72407
8 (+)	Klebsiella variicola 150519 03 SAH	1.98	244366
9 (+)	Klebsiella variicola DSM 15968T HAM	1.96	244366
10 (+)	Klebsiella variicola NY 1125 MCW	1.95	244366

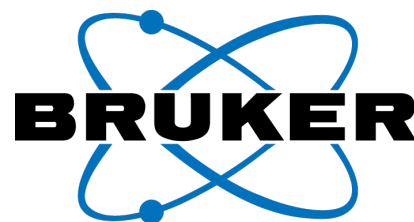
Sample 14



Sample Name: E6
Sample Description:
Sample ID: * C 871
Sample Creation Date/Time: 2021-01-15T12:38:13.796
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola DSM 15968T DSM	2.26	244366
2 (+++)	Klebsiella variicola CC 2053 MCW	2.24	244366
3 (+++)	Klebsiella variicola 150519_03 SAH	2.21	244366
4 (+++)	Klebsiella variicola 150416_01 BRUS	2.18	244366
5 (+++)	Klebsiella variicola TC 4131 MCW	2.13	244366
6 (+++)	Klebsiella variicola NY 1125 MCW	2.11	244366
7 (+++)	Klebsiella variicola NY 1064 MCW	2.11	244366
8 (+++)	Klebsiella variicola 37924 PFM	2.10	244366
9 (+++)	Klebsiella variicola CC 2123 MCW	2.06	244366
10 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.02	72407

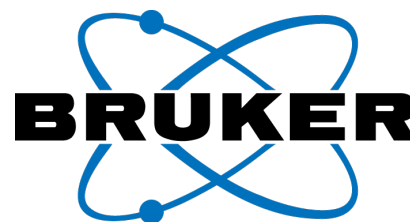
Sample 15



Sample Name: E7
Sample Description:
Sample ID: * C 863
Sample Creation Date/Time: 2021-01-15T12:38:13.796
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola CC_2053 MCW	2.42	244366
2 (+++)	Klebsiella variicola DSM_15968T DSM	2.39	244366
3 (+++)	Klebsiella variicola 150519_03 SAH	2.36	244366
4 (+++)	Klebsiella variicola TC_4131 MCW	2.35	244366
5 (+++)	Klebsiella variicola 150416_01 BRUS	2.35	244366
6 (+++)	Klebsiella variicola NY_1064 MCW	2.34	244366
7 (+++)	Klebsiella variicola CC_2123 MCW	2.31	244366
8 (+++)	Klebsiella variicola NY_1036 MCW	2.27	244366
9 (+++)	Klebsiella variicola NY_1125 MCW	2.27	244366
10 (+++)	Klebsiella variicola 37924 PFM	2.27	244366

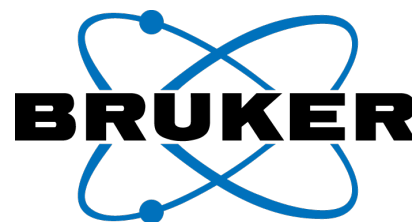
Sample 16



Sample Name: E8
Sample Description:
Sample ID: * C 935
Sample Creation Date/Time: 2021-01-15T12:38:13.797
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): B
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+)	Enterobacter kobei DSM 13645T DSM	1.97	208224
2 (+)	Enterobacter bugandensis DSM 101090 DSM	1.94	547
3 (+)	Enterobacter bugandensis DSM 29888T DSM	1.93	547
4 (+)	Enterobacter bugandensis DSM 101087 DSM	1.91	547
5 (+)	Enterobacter bugandensis DSM 101086 DSM	1.90	547
6 (+)	Enterobacter bugandensis DSM 101089 DSM	1.87	547
7 (+)	Enterobacter cloacae MB_5277_05 THL	1.86	550
8 (+)	Enterobacter asburiae DSM 17506T DSM	1.85	61645
9 (+)	Enterobacter bugandensis DSM 101092 DSM	1.85	547
10 (+)	Enterobacter cloacae MB11506_1 CHB	1.84	550

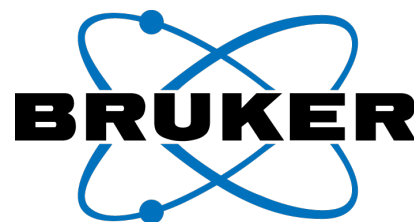
Sample 17



Sample Name: E9
Sample Description:
Sample ID: * C 926
Sample Creation Date/Time: 2021-01-15T12:38:13.798
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.15	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.11	573
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.08	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.07	72407
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.02	72407
6 (+++)	Klebsiella pneumoniae 37585_PFM	2.00	573
7 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	1.99	574
8 (+)	Klebsiella variicola DSM 15968T_HAM	1.93	244366
9 (+)	Klebsiella variicola DSM 15968T DSM	1.91	244366
10 (+)	Klebsiella variicola 150416_01_BRUS	1.87	244366

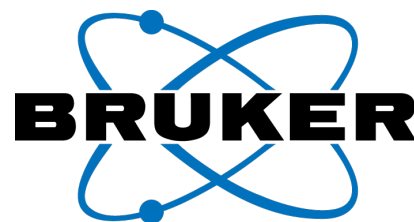
Sample 18



Sample Name: E10
Sample Description:
Sample ID: * C 980
Sample Creation Date/Time: 2021-01-15T12:38:13.798
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): B
Applied Taxonomy Tree: Bruker Taxonomy

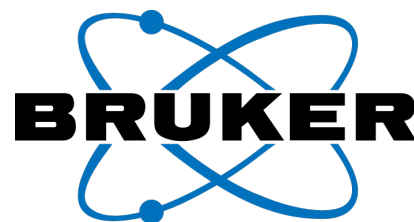
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+)	Acinetobacter baumannii DSM 30007T HAM	1.92	470
2 (+)	Acinetobacter baumannii LMG 994 HAM	1.85	470
3 (+)	Acinetobacter baumannii B389 UFL	1.77	470
4 (+)	Acinetobacter baumannii GEIH_2000 HCB	1.77	470
5 (+)	Acinetobacter baumannii DSM 30008 DSM	1.77	470
6 (+)	Acinetobacter baumannii 13101_1 CHB	1.77	470
7 (-)	Acinetobacter baumannii DSM 30011 DSM	1.65	470
8 (-)	Acinetobacter baumannii DSM 30007T_QC DSM	1.51	470
9 (-)	Acinetobacter baumannii ECII RUH_134 HCB	1.50	470
10 (-)	Acinetobacter pittii 20 HCB	1.49	48296

Sample 19



Sample Name: E11
Sample Description:
Sample ID: * C 1001
Sample Creation Date/Time: 2021-01-15T12:38:13.799
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

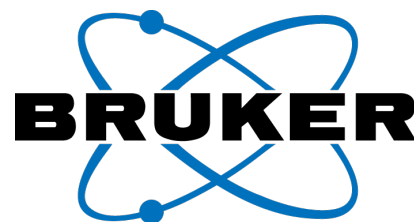
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.29	573
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.20	574
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.19	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.15	72407
5 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.14	72407
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.08	72407
7 (+)	Klebsiella pneumoniae 37595_PFM	1.98	573
8 (+)	Klebsiella variicola DSM 15968T DSM	1.97	244366
9 (+)	Klebsiella variicola DSM 15968T HAM	1.96	244366
10 (+)	Klebsiella variicola 37924_PFM	1.93	244366

Sample 20

Sample Name: E12
Sample Description:
Sample ID: * C 1008
Sample Creation Date/Time: 2021-01-15T12:38:13.799
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.27	573
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.18	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.14	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.11	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.05	72407
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.02	72407
7 (+++)	Klebsiella variicola DSM 15968T HAM	2.02	244366
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	2.00	39831
9 (+)	Klebsiella variicola 150519_03_SAH	1.93	244366
10 (+)	Klebsiella pneumoniae 37585_PFM	1.93	573

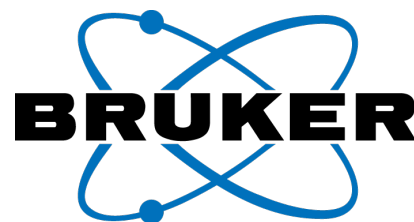
Sample 21



Sample Name: F1
Sample Description:
Sample ID: * C 1003
Sample Creation Date/Time: 2021-01-15T12:38:13.800
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.45	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.41	72407
3 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.29	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.27	574
5 (+++)	Klebsiella variicola CC 2053 MCW	2.27	244366
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.26	574
7 (+++)	Klebsiella pneumoniae 37585 PFM	2.24	573
8 (+++)	Klebsiella variicola TC 4131 MCW	2.22	244366
9 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.20	72407
10 (+++)	Klebsiella variicola 150519_03 SAH	2.19	244366

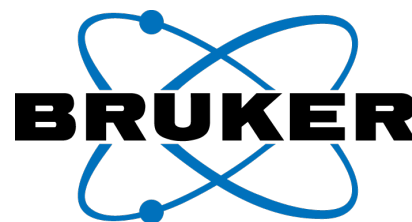
Sample 22



Sample Name: F2
Sample Description:
Sample ID: * C 1039
Sample Creation Date/Time: 2021-01-15T12:38:13.800
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.51	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.39	72407
3 (+++)	Klebsiella pneumoniae 37585 PFM	2.36	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.34	574
5 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.30	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.29	574
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.20	72407
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.19	573
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.13	244366
10 (+++)	Klebsiella variicola DSM 15968T DSM	2.03	244366

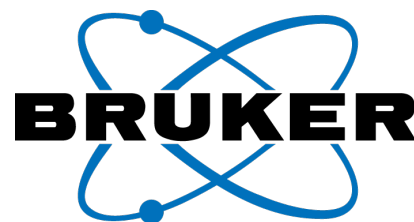
Sample 23



Sample Name: F3
Sample Description:
Sample ID: * C 1046
Sample Creation Date/Time: 2021-01-15T12:38:13.801
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola CC_2053 MCW	2.19	244366
2 (+++)	Klebsiella variicola DSM_15968T DSM	2.13	244366
3 (+++)	Klebsiella variicola 150519_03 SAH	2.07	244366
4 (+++)	Klebsiella variicola TC_4131 MCW	2.05	244366
5 (+++)	Klebsiella variicola NY_1125 MCW	2.04	244366
6 (+++)	Klebsiella variicola CC_2123 MCW	2.04	244366
7 (+++)	Klebsiella variicola NY_1064 MCW	2.02	244366
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.00	244366
9 (+)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	1.99	72407
10 (+)	Klebsiella variicola NY_1036 MCW	1.96	244366

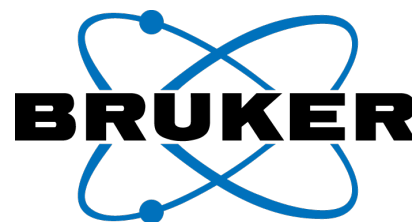
Sample 24



Sample Name: F4
Sample Description:
Sample ID: * C 1077
Sample Creation Date/Time: 2021-01-15T12:38:13.801
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

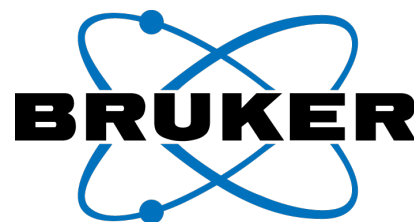
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola 150416_01 BRUS	2.31	244366
2 (+++)	Klebsiella variicola CC 2053 MCW	2.28	244366
3 (+++)	Klebsiella variicola DSM 15968T DSM	2.28	244366
4 (+++)	Klebsiella variicola 150519_03 SAH	2.27	244366
5 (+++)	Klebsiella variicola NY 1064 MCW	2.22	244366
6 (+++)	Klebsiella variicola TC 4131 MCW	2.19	244366
7 (+++)	Klebsiella variicola 37924 PFM	2.17	244366
8 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.16	574
9 (+++)	Klebsiella variicola 150519_01 SAH	2.11	244366
10 (+++)	Klebsiella variicola CC 2123 MCW	2.11	244366

Sample 25



Sample Name: F5
Sample Description:
Sample ID: * C 1075
Sample Creation Date/Time: 2021-01-15T12:38:13.802
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoXecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.31	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.26	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.21	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.21	574
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.20	573
6 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.18	573
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.16	574
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.08	573
9 (+)	Klebsiella variicola 150416_01 BRUS	1.98	244366
10 (+)	Klebsiella variicola 150519_03 SAH	1.91	244366

Sample 26

Sample Name: F6
Sample Description:
Sample ID: * C 117
Sample Creation Date/Time: 2021-01-15T12:38:13.802
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
AutoExecute Method: MBT_AutoX
Consistency Category (based on 2 best matches): A
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.22	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.15	573
3 (+++)	Klebsiella pneumoniae 37585 PFM	2.14	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.10	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.04	72407
6 (+)	Klebsiella pneumoniae 37595 PFM	1.99	573
7 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.93	72407
8 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	1.93	574
9 (+)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	1.91	39831
10 (+)	Klebsiella variicola 150519_03 SAH	1.76	244366