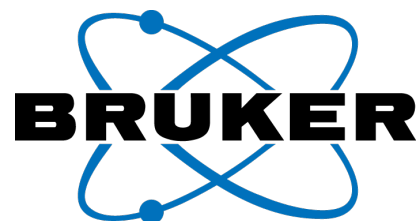


Bruker MALDI Biotyper Identification Results



Run Info:

Run Identifier: 210107-1120-10184
Comment:
Operator: maldi@FLEX-PC
Run Creation Date/Time: 2021-01-07T11:35:38.162
Number of Tests: 23
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
A1 (+++)(A)	* A 116 (Standard)	Klebsiella pneumoniae	2.32	Klebsiella pneumoniae	2.30
A2 (+++)(A)	* A 118 (Standard)	Klebsiella pneumoniae	2.28	Klebsiella pneumoniae	2.28
A3 (+++)(A)	* A 130 (Standard)	Klebsiella pneumoniae	2.28	Klebsiella pneumoniae	2.23
A4 (+++)(A)	* A 261 (Standard)	Klebsiella pneumoniae	2.40	Klebsiella pneumoniae	2.31
A5 (+++)(A)	* A 184 (Standard)	Klebsiella pneumoniae	2.27	Klebsiella pneumoniae	2.22
A6 (+++)(A)	* A 212 (Standard)	Klebsiella variicola	2.31	Klebsiella variicola	2.23
A7 (+++)(A)	* B 077 (Standard)	Klebsiella pneumoniae	2.34	Klebsiella pneumoniae	2.33

Result overview table--continued on next page

Result overview table--continued from previous page					
Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
A8 (+) (B)	* B 135 (Standard)	Enterobacter bugandensis	1.93	Enterobacter bugandensis	1.87
A9 (+++)(A)	* B 157 (Standard)	Klebsiella variicola	2.24	Klebsiella variicola	2.21
A10 (-) (C)	* B 192 (Standard)	No Organism Identification Possible	1.59	No Organism Identification Possible	1.59
A11 (+++)(A)	* B 248 (Standard)	Klebsiella pneumoniae	2.32	Klebsiella pneumoniae	2.23
A12 (+++)(A)	* C 124 (Standard)	Klebsiella pneumoniae	2.16	Klebsiella pneumoniae	2.14
B1 (+++)(A)	* C 130 (Standard)	Klebsiella pneumoniae	2.36	Klebsiella pneumoniae	2.35
B2 (+++)(A)	* C 161 (Standard)	Klebsiella pneumoniae	2.34	Klebsiella pneumoniae	2.34
B3 (+++)(A)	* C 198 (Standard)	Klebsiella pneumoniae	2.11	Klebsiella pneumoniae	2.05
B4 (+++)(A)	* C 228 (Standard)	Klebsiella pneumoniae	2.13	Klebsiella pneumoniae	2.08
B5 (+++)(A)	* C 239 (Standard)	Klebsiella variicola	2.16	Klebsiella variicola	2.15
B6 (+++)(A)	* C 240 (Standard)	Klebsiella pneumoniae	2.26	Klebsiella pneumoniae	2.24
B7 (+++)(A)	* C 243 (Standard)	Klebsiella pneumoniae	2.31	Klebsiella pneumoniae	2.30
B8 (+++)(A)	* C 257 (Standard)	Klebsiella pneumoniae	2.40	Klebsiella pneumoniae	2.31
B9 (+++)(A)	* C 276 (Standard)	Klebsiella pneumoniae	2.36	Klebsiella pneumoniae	2.36
B10 (+++)(A)	* C 303 (Standard)	Klebsiella pneumoniae	2.31	Klebsiella pneumoniae	2.27
B11 (+++)(A)	* C 311 (Standard)	Klebsiella pneumoniae	2.44	Klebsiella pneumoniae	2.36

Matching Hints

Matched Pattern	Comment
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 30008 DSM	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter baumannii ECI RUH_875 HCB	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 ECIII RUH_5875 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 nosocomialis AL_95 HCB	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
Acinetobacter nosocomialis LMG 10626 LMG	Member of the Acinetobacter baumannii /calcoaceticus complex. Extraction must be performed to permit reliable species identification.
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
Enterobacter bugandensis DSM 29888T DSM	is a member of Enterobacter cloacae complex
Enterobacter cloacae 20105_2 CHB	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 kobei DSM 13645T DSM	is a member of Enterobacter cloacae complex
Klebsiella aerogenes DSM 12058 DSM	synonym of Enterobacter aerogenes
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
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
<i>Matching Hints table--continued on next page</i>	

<i>Matching Hints table--continued from previous page</i>	
Matched Pattern	Comment
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_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

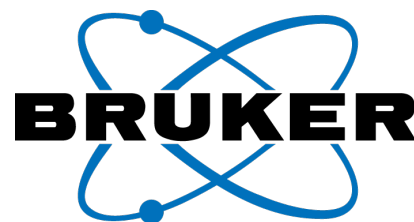
Meaning of Score Values

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: A1
Sample Description:
Sample ID: * A 116
Sample Creation Date/Time: 2021-01-07T11:24:58.364
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 RV_BA_03_B_LBK	2.30	573
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.27	574
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.26	574
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.25	573
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.20	72407
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.17	72407
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.09	573
9 (+++)	Klebsiella variicola NY_1064 MCW	2.02	244366
10 (+++)	Klebsiella variicola 150519_03 SAH	2.00	244366

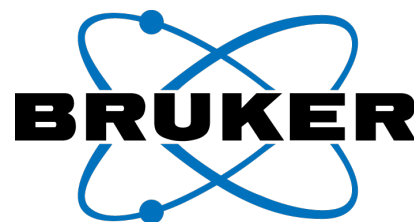
Sample 2



Sample Name: A2
Sample Description:
Sample ID: * A 118
Sample Creation Date/Time: 2021-01-07T11:24:58.365
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	2.28	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.28	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.23	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.22	574
5 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.21	573
6 (+++)	Klebsiella pneumoniae 37585 PFM	2.19	573
7 (+++)	Klebsiella variicola 150416_01 BRUS	2.13	244366
8 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.12	72407
9 (+++)	Klebsiella variicola CC_2053 MCW	2.09	244366
10 (+++)	Klebsiella variicola 150519_03 SAH	2.01	244366

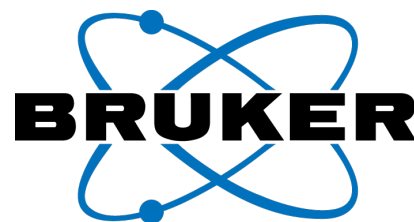
Sample 3



Sample Name: A3
Sample Description:
Sample ID: * A 130
Sample Creation Date/Time: 2021-01-07T11:24:58.366
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 HAM	2.28	574
2 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.23	573
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.23	72407
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.22	72407
5 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.22	72407
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.19	574
7 (+++)	Klebsiella pneumoniae 37585 PFM	2.12	573
8 (+++)	Klebsiella variicola DSM 15968T DSM	2.07	244366
9 (+++)	Klebsiella variicola 150519_03 SAH	2.02	244366
10 (+)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	1.99	39831

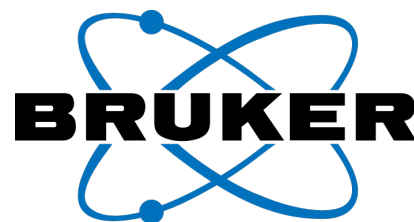
Sample 4



Sample Name: A4
Sample Description:
Sample ID: * A 261
Sample Creation Date/Time: 2021-01-07T11:24:58.366
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

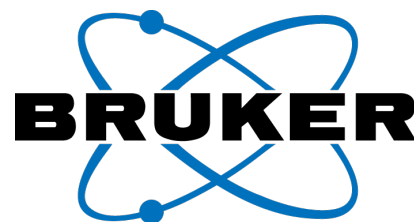
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.40	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.31	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.30	574
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.30	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.29	574
6 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.27	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.24	72407
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.20	244366
9 (+++)	Klebsiella pneumoniae 37595 PFM	2.15	573
10 (+++)	Klebsiella variicola 150519_03 SAH	2.09	244366

Sample 5



Sample Name: A5
Sample Description:
Sample ID: * A 184
Sample Creation Date/Time: 2021-01-07T11:24:58.368
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

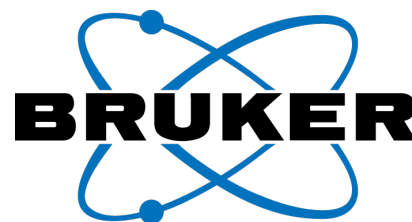
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.27	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.22	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.19	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.16	72407
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.12	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.10	574
7 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.09	573
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.07	573
9 (+)	Klebsiella variicola CC_2053 MCW	1.93	244366
10 (+)	Klebsiella variicola 150519_03 SAH	1.93	244366

Sample 6

Sample Name: A6
Sample Description:
Sample ID: * A 212
Sample Creation Date/Time: 2021-01-07T11:24:58.368
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	2.31	244366
2 (+++)	Klebsiella variicola 37924 PFM	2.23	244366
3 (+++)	Klebsiella variicola CC_2053 MCW	2.22	244366
4 (+++)	Klebsiella variicola 150416_01 BRUS	2.21	244366
5 (+++)	Klebsiella variicola 150519_03 SAH	2.21	244366
6 (+++)	Klebsiella variicola TC_4131 MCW	2.18	244366
7 (+++)	Klebsiella variicola NY_1064 MCW	2.14	244366
8 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.13	574
9 (+++)	Klebsiella variicola DSM 15968T HAM	2.13	244366
10 (+++)	Klebsiella variicola NY_1125 MCW	2.13	244366

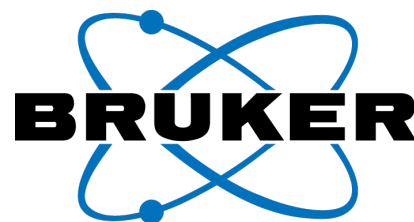
Sample 7



Sample Name: A7
Sample Description:
Sample ID: * B 077
Sample Creation Date/Time: 2021-01-07T11:24:58.369
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	2.34	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.33	573
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.31	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.24	574
5 (+++)	Klebsiella pneumoniae 37585_PFM	2.24	573
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.22	72407
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.21	574
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	2.06	39831
9 (+++)	Klebsiella variicola 150519_03_SAH	2.00	244366
10 (+++)	Klebsiella pneumoniae 37595_PFM	2.00	573

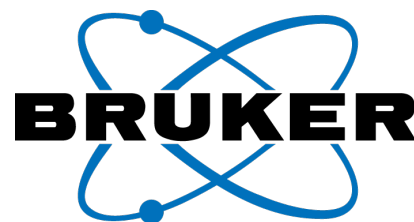
Sample 8



Sample Name: A8
Sample Description:
Sample ID: * B 135
Sample Creation Date/Time: 2021-01-07T11:24:58.369
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 bugandensis DSM 101090 DSM	1.93	547
2 (+)	Enterobacter bugandensis DSM 101086 DSM	1.87	547
3 (+)	Enterobacter bugandensis DSM 101087 DSM	1.86	547
4 (+)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	1.78	72407
5 (+)	Enterobacter bugandensis DSM 101089 DSM	1.77	547
6 (+)	Enterobacter cloacae 20105_2 CHB	1.77	550
7 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.76	72407
8 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	1.76	72407
9 (+)	Enterobacter bugandensis DSM 29888T DSM	1.75	547
10 (+)	Enterobacter kobei DSM 13645T DSM	1.73	208224

Sample 9



Sample Name: A9
Sample Description:
Sample ID: * B 157
Sample Creation Date/Time: 2021-01-07T11:24:58.370
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	2.24	244366
2 (+++)	Klebsiella variicola CC 2053 MCW	2.21	244366
3 (+++)	Klebsiella variicola 150416_01 BRUS	2.13	244366
4 (+++)	Klebsiella variicola TC 4131 MCW	2.08	244366
5 (+++)	Klebsiella variicola NY 1064 MCW	2.08	244366
6 (+++)	Klebsiella variicola CC 2123 MCW	2.04	244366
7 (+++)	Klebsiella variicola 150519_03 SAH	2.04	244366
8 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.02	574
9 (+)	Klebsiella variicola 37924 PFM	1.99	244366
10 (+)	Klebsiella variicola NY 1125 MCW	1.99	244366

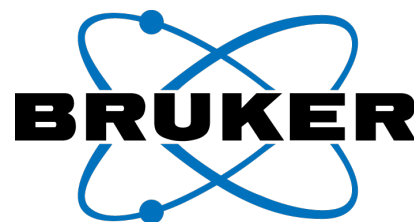
Sample 10



Sample Name: A10
Sample Description:
Sample ID: * B 192
Sample Creation Date/Time: 2021-01-07T11:24:58.370
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): C
Applied Taxonomy Tree: Bruker Taxonomy

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (-)	Acinetobacter baumannii GEIH_2000 HCB	1.59	470
2 (-)	Acinetobacter baumannii ECI RUH_875 HCB	1.59	470
3 (-)	Acinetobacter nosocomialis LMG_10626 LMG	1.56	72607
4 (-)	Acinetobacter baumannii DSM_30007T HAM	1.55	470
5 (-)	Acinetobacter baumannii LMG_994 HAM	1.54	470
6 (-)	Acinetobacter baumannii ECII RUH_134 HCB	1.51	470
7 (-)	Acinetobacter baumannii DSM_30008 DSM	1.48	470
8 (-)	Acinetobacter baumannii ECIII RUH_5875 HCB	1.47	470
9 (-)	Acinetobacter baumannii B389 UFL	1.46	470
10 (-)	Acinetobacter nosocomialis AL_95 HCB	1.35	72607

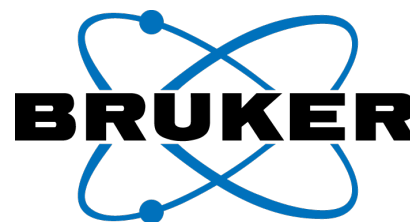
Sample 11



Sample Name: A11
Sample Description:
Sample ID: * B 248
Sample Creation Date/Time: 2021-01-07T11:24:58.371
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	2.32	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.23	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.18	72407
4 (+++)	Klebsiella variicola 150416_01 BRUS	2.11	244366
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.08	574
6 (+++)	Klebsiella variicola TC_4131 MCW	2.03	244366
7 (+++)	Klebsiella variicola CC_2053 MCW	2.02	244366
8 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.00	573
9 (+++)	Klebsiella variicola NY_1064 MCW	2.00	244366
10 (+)	Klebsiella pneumoniae 37585 PFM	1.97	573

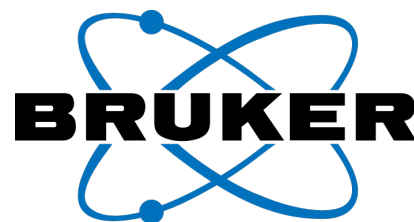
Sample 12



Sample Name: A12
Sample Description:
Sample ID: * C 124
Sample Creation Date/Time: 2021-01-07T11:24:58.371
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	2.16	573
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.14	574
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.13	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.11	72407
5 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.10	72407
6 (+++)	Klebsiella pneumoniae 37595_PFM	2.00	573
7 (+)	Klebsiella pneumoniae 37585_PFM	1.98	573
8 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.95	72407
9 (+)	Klebsiella variicola DSM 15968T HAM	1.93	244366
10 (+)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	1.90	39831

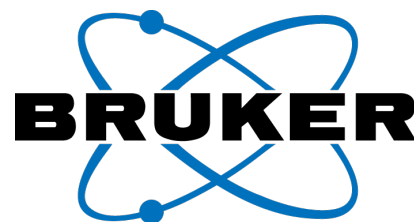
Sample 13



Sample Name: B1
Sample Description:
Sample ID: * C 130
Sample Creation Date/Time: 2021-01-07T11:24:58.372
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	2.36	573
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.35	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.32	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.27	574
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.23	574
6 (+++)	Klebsiella pneumoniae 37585_PFM	2.15	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.14	72407
8 (+++)	Klebsiella variicola NY_1064_MCW	2.04	244366
9 (+++)	Klebsiella variicola CC_2053_MCW	2.01	244366
10 (+++)	Klebsiella variicola 150519_03_SAH	2.01	244366

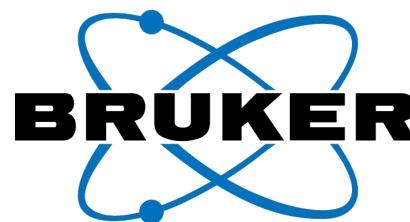
Sample 14



Sample Name: B2
Sample Description:
Sample ID: * C 161
Sample Creation Date/Time: 2021-01-07T11:24:58.372
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	2.34	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.34	72407
3 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.25	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.24	574
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.23	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.21	574
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.21	72407
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.01	573
9 (+++)	Klebsiella variicola CC 2053 MCW	2.00	244366
10 (+)	Klebsiella variicola 150519_03 SAH	1.99	244366

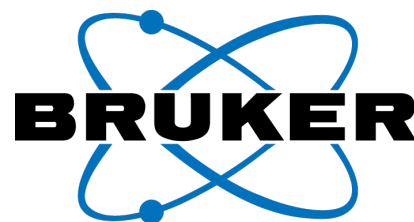
Sample 15



Sample Name: B3
Sample Description:
Sample ID: * C 198
Sample Creation Date/Time: 2021-01-07T11:24:58.373
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	2.11	573
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.05	574
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.03	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.01	72407
5 (+)	Klebsiella variicola NY_1064_MCW	1.89	244366
6 (+)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	1.88	72407
7 (+)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	1.88	39831
8 (+)	Klebsiella variicola 150519_03_SAH	1.87	244366
9 (+)	Klebsiella aerogenes DSM 12058 DSM	1.85	28451
10 (+)	Klebsiella pneumoniae 37595_PFM	1.84	573

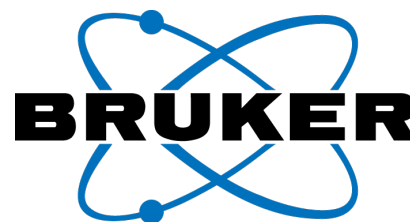
Sample 16



Sample Name: B4
Sample Description:
Sample ID: * C 228
Sample Creation Date/Time: 2021-01-07T11:24:58.373
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	2.13	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.08	72407
3 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.08	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.06	574
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.04	574
6 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.97	72407
7 (+)	Klebsiella pneumoniae 37585 PFM	1.96	573
8 (+)	Klebsiella pneumoniae 37595 PFM	1.87	573
9 (+)	Klebsiella variicola TC 4131 MCW	1.80	244366
10 (+)	Klebsiella variicola DSM 15968T HAM	1.80	244366

Sample 17



Sample Name: B5
Sample Description:
Sample ID: * C 239
Sample Creation Date/Time: 2021-01-07T11:24:58.374
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	2.16	244366
2 (+++)	Klebsiella variicola 150416_01 BRUS	2.15	244366
3 (+++)	Klebsiella variicola TC_4131 MCW	2.14	244366
4 (+++)	Klebsiella variicola CC_2053 MCW	2.12	244366
5 (+++)	Klebsiella variicola NY_1064 MCW	2.09	244366
6 (+++)	Klebsiella variicola 150519_03 SAH	2.08	244366
7 (+++)	Klebsiella variicola NY_1125 MCW	2.05	244366
8 (+++)	Klebsiella variicola 150519_01 SAH	2.04	244366
9 (+++)	Klebsiella variicola 37924 PFM	2.03	244366
10 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	1.97	574

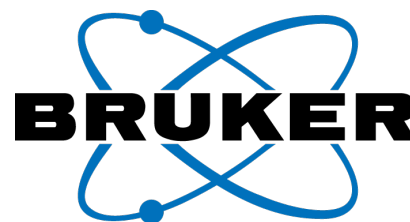
Sample 18



Sample Name: B6
Sample Description:
Sample ID: * C 240
Sample Creation Date/Time: 2021-01-07T11:24:58.374
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

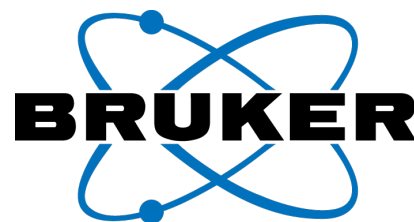
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.26	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.24	72407
3 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.23	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.20	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.13	72407
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.08	574
7 (+++)	Klebsiella pneumoniae 37585 PFM	2.02	573
8 (+)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	1.99	39831
9 (+)	Klebsiella variicola 150416_01 BRUS	1.99	244366
10 (+)	Klebsiella pneumoniae 37595 PFM	1.88	573

Sample 19



Sample Name: B7
Sample Description:
Sample ID: * C 243
Sample Creation Date/Time: 2021-01-07T11:24:58.375
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

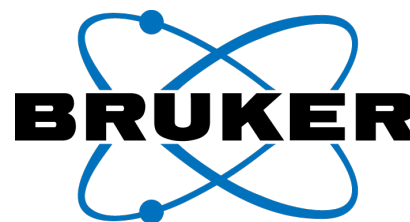
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.31	574
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.30	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.25	574
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.23	573
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.23	72407
6 (+++)	Klebsiella pneumoniae 37585 PFM	2.13	573
7 (+++)	Klebsiella pneumoniae 37595 PFM	2.06	573
8 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.04	72407
9 (+)	Klebsiella variicola 150416_01 BRUS	1.92	244366
10 (+)	Klebsiella variicola 150519_03 SAH	1.86	244366

Sample 20

Sample Name: B8
Sample Description:
Sample ID: * C 257
Sample Creation Date/Time: 2021-01-07T11:24:58.375
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

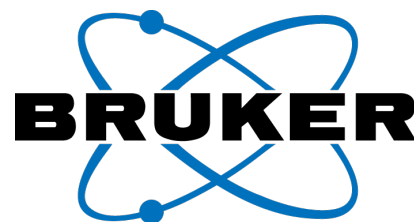
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.40	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.31	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.27	574
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.21	574
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.20	573
6 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.19	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.18	72407
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.09	573
9 (+)	Klebsiella variicola 150519_03 SAH	1.99	244366
10 (+)	Klebsiella variicola 150416_01 BRUS	1.97	244366

Sample 21



Sample Name: B9
Sample Description:
Sample ID: * C 276
Sample Creation Date/Time: 2021-01-07T11:24:58.376
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

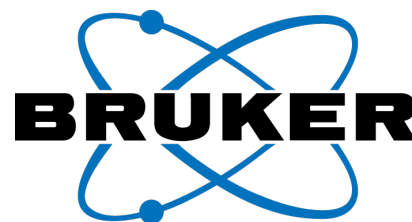
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.36	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.36	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.34	72407
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.33	573
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.32	72407
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.31	574
7 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.30	573
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.17	573
9 (+++)	Klebsiella variicola NY_1064 MCW	2.05	244366
10 (+++)	Klebsiella variicola 37924 PFM	2.03	244366

Sample 22

Sample Name: B10
Sample Description:
Sample ID: * C 303
Sample Creation Date/Time: 2021-01-07T11:24:58.376
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 ozaenae DSM 16358T DSM	2.27	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.21	72407
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.16	573
5 (+++)	Klebsiella variicola 150416_01 BRUS	2.08	244366
6 (+++)	Klebsiella pneumoniae 37585 PFM	2.07	573
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.06	574
8 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.99	72407
9 (+)	Klebsiella variicola 150519_03 SAH	1.95	244366
10 (+)	Klebsiella variicola CC_2053 MCW	1.95	244366

Sample 23



Sample Name: B11
Sample Description:
Sample ID: * C 311
Sample Creation Date/Time: 2021-01-07T11:24:58.377
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	2.44	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.36	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.26	574
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.26	573
5 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.23	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.16	574
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.12	72407
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.10	573
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.00	244366
10 (+)	Klebsiella variicola 150519_03 SAH	1.99	244366