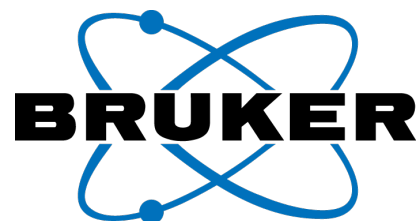


Bruker MALDI Biotyper Identification Results



Run Info:

Run Identifier: 210112-1156-10113
Comment:
Operator: maldi@FLEX-PC
Run Creation Date/Time: 2021-01-12T12:18:41.588
Number of Tests: 35
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
C2 (+++)(A)	* A 366 (Standard)	Klebsiella pneumoniae	2.25	Klebsiella pneumoniae	2.19
C3 (+++)(A)	* A 373 (Standard)	Klebsiella variicola	2.28	Klebsiella variicola	2.27
C4 (+++)(A)	* A 377 (Standard)	Klebsiella pneumoniae	2.43	Klebsiella pneumoniae	2.37
C5 (+++)(A)	* A 413 (Standard)	Klebsiella variicola	2.45	Klebsiella variicola	2.39
C6 (+++)(A)	* A 415 (Standard)	Klebsiella pneumoniae	2.45	Klebsiella pneumoniae	2.39
C7 (+++)(A)	* A 425 (Standard)	Klebsiella variicola	2.42	Klebsiella variicola	2.39
C8 (+++)(A)	* B 301 (Standard)	Klebsiella pneumoniae	2.46	Klebsiella pneumoniae	2.44

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
C9 (+++)(A)	* B 310 (Standard)	Klebsiella pneumoniae	2.48	Klebsiella pneumoniae	2.41
C10 (+++)(A)	* B 328 (Standard)	Klebsiella pneumoniae	2.43	Klebsiella pneumoniae	2.40
C11 (+++)(A)	* B 386 (Standard)	Klebsiella pneumoniae	2.37	Klebsiella pneumoniae	2.28
C12 (+++)(A)	* B 387 (Standard)	Klebsiella pneumoniae	2.35	Klebsiella pneumoniae	2.31
D1 (+++)(A)	* C 379\ (Standard)	Klebsiella pneumoniae	2.40	Klebsiella pneumoniae	2.37
D2 (+++)(A)	* C 473 (Standard)	Klebsiella pneumoniae	2.45	Klebsiella pneumoniae	2.43
D3 (+++)(A)	* C 475 (Standard)	Klebsiella variicola	2.49	Klebsiella variicola	2.38
D4 (+++)(A)	* C 479 (Standard)	Klebsiella variicola	2.51	Klebsiella variicola	2.52
D5 (+++)(A)	* C 480 (Standard)	Klebsiella variicola	2.39	Klebsiella variicola	2.38
D6 (+++)(A)	* C 332 (Standard)	Klebsiella pneumoniae	2.47	Klebsiella pneumoniae	2.40
D7 (+++)(A)	* C 539 (Standard)	Klebsiella pneumoniae	2.45	Klebsiella pneumoniae	2.44
D8 (+++)(A)	* C 549 (Standard)	Klebsiella variicola	2.36	Klebsiella variicola	2.34
D9 (+++)(A)	* C 583 (Standard)	Klebsiella variicola	2.33	Klebsiella variicola	2.24
D10 (+++)(A)	* C 636 (Standard)	Klebsiella pneumoniae	2.37	Klebsiella pneumoniae	2.19
D11 (+++)(A)	* C 647 (Standard)	Klebsiella pneumoniae	2.33	Klebsiella pneumoniae	2.32
D12 (+++)(A)	* A 467 (Standard)	Klebsiella pneumoniae	2.23	Klebsiella pneumoniae	2.15
<i>Result overview table--continued on next page</i>					

Result overview table--continued from previous page					
Sample Name	Sample ID	Organism (best match)	Score Value	Organism (second-best match)	Score Value
E1 (+++)(A)	* A 499 (Standard)	Klebsiella variicola	2.38	Klebsiella variicola	2.37
E2 (+++)(A)	* A 501 (Standard)	Klebsiella pneumoniae	2.34	Klebsiella pneumoniae	2.33
E3 (+++)(A)	* A 504 (Standard)	Klebsiella variicola	2.30	Klebsiella variicola	2.29
E4 (+)(B)	* B 424 (Standard)	Klebsiella pneumoniae	1.95	Klebsiella variicola	1.93
E5 (+++)(A)	* B 454 (Standard)	Klebsiella pneumoniae	2.40	Klebsiella pneumoniae	2.39
E6 (+++)(A)	* B 477 (Standard)	Klebsiella pneumoniae	2.30	Klebsiella pneumoniae	2.24
E7 (+++)(A)	* C 656 (Standard)	Klebsiella pneumoniae	2.33	Klebsiella pneumoniae	2.31
E8 (+++)(A)	* C 713 (Standard)	Klebsiella variicola	2.41	Klebsiella variicola	2.38
E9 (+++)(A)	* C 542 (Standard)	Klebsiella pneumoniae	2.36	Klebsiella pneumoniae	2.33
E10 (+++)(A)	* C 717 (Standard)	Klebsiella variicola	2.32	Klebsiella variicola	2.30
E11 (+++)(A)	* A 434 (Standard)	Acinetobacter lwoffii	2.06	Acinetobacter lwoffii	1.99
E12 (+)(B)	* A 446 (Standard)	Acinetobacter radioresistens	1.84	Acinetobacter radioresistens	1.73

Matching Hints

Matched Pattern	Comment
Acinetobacter lwoffii 11_0038419_001_01 LGL	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii 13 PIM	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii 2_Ring240 MHH	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii 54 PIM	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii B101 UFL	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii B138 UFL	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii DSM 2403T DSM	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii LMG 10596 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii LMG 1325 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter lwoffii V506 MCRF	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens B381 UFL	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens DSM 6976T HAM	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens LMG 10609 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens LMG 10610 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens LMG 10611 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens LMG 10612 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
<i>Matching Hints table--continued on next page</i>	

<i>Matching Hints table--continued from previous page</i>	
Matched Pattern	Comment
Acinetobacter radioresistens LMG 10614 HAM	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter radioresistens LMG 10615 LMG	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
Acinetobacter townieri DSM 14962T HAM	Species of this genus have very similar patterns: Therefore distinguishing their species is difficult.
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
<i>Matching Hints table--continued on next page</i>	

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

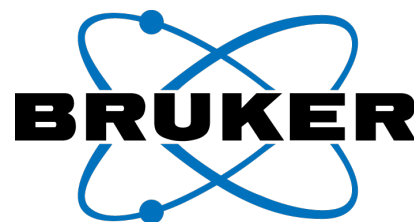
Sample 1



Sample Name: C2
Sample Description:
Sample ID: * A 366
Sample Creation Date/Time: 2021-01-12T12:01:56.747
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	2.35	573
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.19	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.19	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.16	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.09	72407
6 (+++)	Klebsiella variicola DSM 15968T HAM	2.07	244366
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.05	72407
8 (+++)	Klebsiella variicola 37924 PFM	2.04	244366
9 (+++)	Klebsiella variicola DSM 15968T DSM	2.03	244366
10 (+++)	Klebsiella pneumoniae 37585 PFM	2.02	573

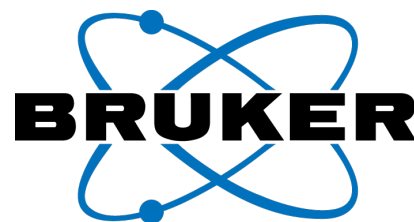
Sample 2



Sample Name: C3
Sample Description:
Sample ID: * A 373
Sample Creation Date/Time: 2021-01-12T12:01:56.747
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 variicola CC_2053 MCW	2.28	244366
2 (+++)	Klebsiella variicola DSM_15968T DSM	2.27	244366
3 (+++)	Klebsiella variicola 150416_01 BRUS	2.22	244366
4 (+++)	Klebsiella variicola TC_4131 MCW	2.21	244366
5 (+++)	Klebsiella variicola 150519_03 SAH	2.16	244366
6 (+++)	Klebsiella variicola NY_1125 MCW	2.15	244366
7 (+++)	Klebsiella variicola 37924 PFM	2.13	244366
8 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.09	72407
9 (+++)	Klebsiella variicola NY_1064 MCW	2.09	244366
10 (+++)	Klebsiella variicola CC_2123 MCW	2.08	244366

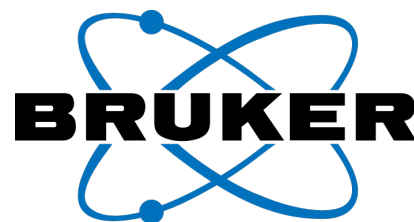
Sample 3



Sample Name: C4
Sample Description:
Sample ID: * A 377
Sample Creation Date/Time: 2021-01-12T12:01:56.748
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	2.43	573
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.37	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.34	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.31	574
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.31	574
6 (+++)	Klebsiella pneumoniae 37585_PFM	2.28	573
7 (+++)	Klebsiella pneumoniae 37595_PFM	2.20	573
8 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.18	72407
9 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	2.14	39831
10 (+++)	Klebsiella variicola DSM 15968T_HAM	2.11	244366

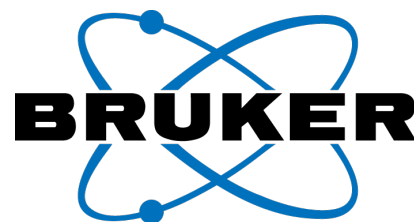
Sample 4



Sample Name: C5
Sample Description:
Sample ID: * A 413
Sample Creation Date/Time: 2021-01-12T12:01:56.748
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	2.45	244366
2 (+++)	Klebsiella variicola 150416_01 BRUS	2.39	244366
3 (+++)	Klebsiella variicola DSM_15968T DSM	2.38	244366
4 (+++)	Klebsiella variicola 150519_03 SAH	2.35	244366
5 (+++)	Klebsiella variicola TC_4131 MCW	2.33	244366
6 (+++)	Klebsiella variicola 37924 PFM	2.30	244366
7 (+++)	Klebsiella variicola CC_2123 MCW	2.29	244366
8 (+++)	Klebsiella variicola NY_1064 MCW	2.29	244366
9 (+++)	Klebsiella variicola NY_1125 MCW	2.28	244366
10 (+++)	Klebsiella variicola 150519_01 SAH	2.22	244366

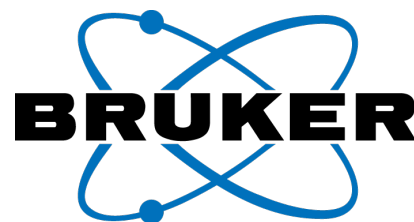
Sample 5



Sample Name: C6
Sample Description:
Sample ID: * A 415
Sample Creation Date/Time: 2021-01-12T12:01:56.749
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	2.45	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.39	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.32	72407
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.27	573
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.26	573
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.18	72407
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.13	574
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.11	244366
9 (+++)	Klebsiella variicola 150519_03 SAH	2.08	244366
10 (+++)	Klebsiella pneumoniae 37595 PFM	2.06	573

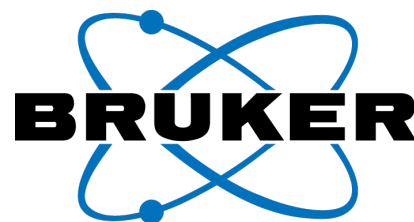
Sample 6



Sample Name: C7
Sample Description:
Sample ID: * A 425
Sample Creation Date/Time: 2021-01-12T12:01:56.749
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

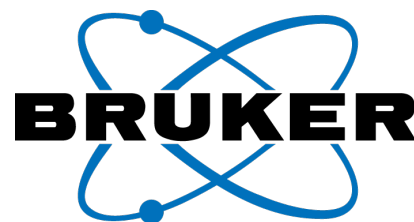
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola CC_2053 MCW	2.42	244366
2 (+++)	Klebsiella variicola 150416_01 BRUS	2.39	244366
3 (+++)	Klebsiella variicola TC_4131 MCW	2.39	244366
4 (+++)	Klebsiella variicola DSM_15968T DSM	2.36	244366
5 (+++)	Klebsiella variicola NY_1064 MCW	2.32	244366
6 (+++)	Klebsiella variicola 150519_03 SAH	2.32	244366
7 (+++)	Klebsiella variicola CC_2123 MCW	2.30	244366
8 (+++)	Klebsiella variicola NY_1125 MCW	2.26	244366
9 (+++)	Klebsiella variicola 37924 PFM	2.22	244366
10 (+++)	Klebsiella variicola 150519_01 SAH	2.20	244366

Sample 7



Sample Name: C8
Sample Description:
Sample ID: * B 301
Sample Creation Date/Time: 2021-01-12T12:01:56.750
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	2.46	573
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.44	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.43	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.40	574
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.38	574
6 (+++)	Klebsiella pneumoniae 37585_PFM	2.31	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.30	72407
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	2.25	39831
9 (+++)	Klebsiella pneumoniae 37595_PFM	2.21	573
10 (+++)	Klebsiella variicola 37924_PFM	2.09	244366

Sample 8

Sample Name: C9
Sample Description:
Sample ID: * B 310
Sample Creation Date/Time: 2021-01-12T12:01:56.750
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	2.48	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.41	573
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.39	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.32	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.29	72407
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.25	574
7 (+++)	Klebsiella pneumoniae 37585_PFM	2.24	573
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	2.11	39831
9 (+++)	Klebsiella pneumoniae 37595_PFM	2.10	573
10 (+++)	Klebsiella variicola TC_4131_MCW	2.07	244366

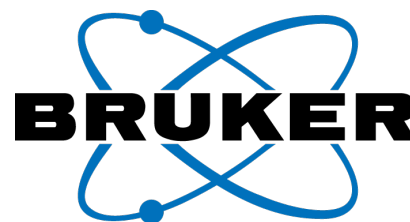
Sample 9



Sample Name: C10
Sample Description:
Sample ID: * B 328
Sample Creation Date/Time: 2021-01-12T12:01:56.751
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	2.43	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.40	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.38	574
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.37	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.35	574
6 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.34	573
7 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.27	72407
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.13	244366
9 (+++)	Klebsiella pneumoniae 37595 PFM	2.12	573
10 (+++)	Klebsiella variicola 150519_03 SAH	2.12	244366

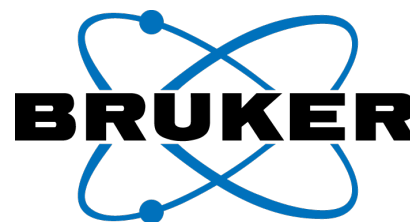
Sample 10



Sample Name: C11
Sample Description:
Sample ID: * B 386
Sample Creation Date/Time: 2021-01-12T12:01:56.751
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	2.37	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.28	573
3 (+++)	Klebsiella pneumoniae 37585 PFM	2.28	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.28	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.25	72407
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.19	72407
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.17	574
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.06	573
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.00	244366
10 (+)	Klebsiella variicola TC_4131 MCW	1.97	244366

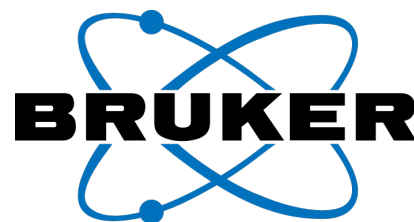
Sample 11



Sample Name: C12
Sample Description:
Sample ID: * B 387
Sample Creation Date/Time: 2021-01-12T12:01:56.752
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	2.35	72407
2 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.31	573
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.25	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.22	72407
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.20	573
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.19	72407
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.13	574
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.08	244366
9 (+++)	Klebsiella variicola 150519_03 SAH	2.04	244366
10 (+)	Klebsiella pneumoniae 37595 PFM	1.99	573

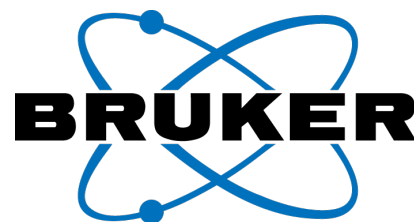
Sample 12



Sample Name: D1
Sample Description:
Sample ID: * C 379\
Sample Creation Date/Time: 2021-01-12T12:01:56.752
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	2.40	573
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.37	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.23	574
4 (+++)	Klebsiella pneumoniae 37585_PFM	2.19	573
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.19	72407
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.09	72407
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.08	574
8 (+++)	Klebsiella pneumoniae 37595_PFM	2.06	573
9 (+)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	1.90	39831
10 (+)	Klebsiella variicola DSM 15968T_DSM	1.84	244366

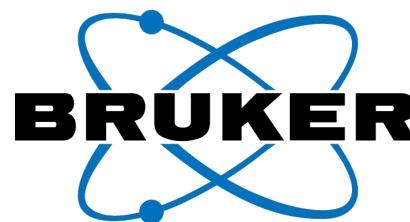
Sample 13



Sample Name: D2
Sample Description:
Sample ID: * C 473
Sample Creation Date/Time: 2021-01-12T12:01:56.753
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 ozaenae DSM 16358T DSM	2.43	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.40	72407
4 (+++)	Klebsiella pneumoniae 37585 PFM	2.32	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.31	574
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.29	72407
7 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.27	573
8 (+++)	Klebsiella variicola 150519_03 SAH	2.15	244366
9 (+++)	Klebsiella variicola CC_2053 MCW	2.15	244366
10 (+++)	Klebsiella variicola 150416_01 BRUS	2.14	244366

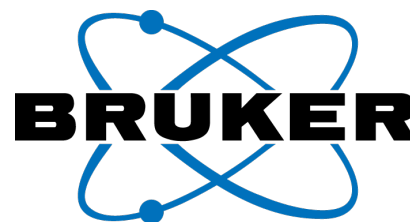
Sample 14



Sample Name: D3
Sample Description:
Sample ID: * C 475
Sample Creation Date/Time: 2021-01-12T12:01:56.753
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	2.49	244366
2 (+++)	Klebsiella variicola 150416_01 BRUS	2.38	244366
3 (+++)	Klebsiella variicola TC_4131 MCW	2.35	244366
4 (+++)	Klebsiella variicola CC_2123 MCW	2.35	244366
5 (+++)	Klebsiella variicola 150519_03 SAH	2.34	244366
6 (+++)	Klebsiella variicola DSM_15968T DSM	2.32	244366
7 (+++)	Klebsiella variicola NY_1064 MCW	2.32	244366
8 (+++)	Klebsiella variicola 37924 PFM	2.30	244366
9 (+++)	Klebsiella variicola 150519_01 SAH	2.29	244366
10 (+++)	Klebsiella variicola NY_1125 MCW	2.28	244366

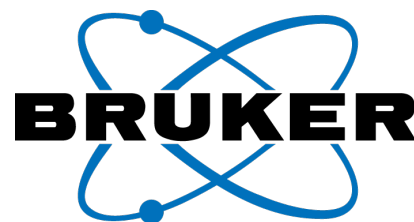
Sample 15



Sample Name: D4
Sample Description:
Sample ID: * C 479
Sample Creation Date/Time: 2021-01-12T12:01:56.754
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	2.54	244366
2 (+++)	Klebsiella variicola CC 2053 MCW	2.52	244366
3 (+++)	Klebsiella variicola 150519_03 SAH	2.48	244366
4 (+++)	Klebsiella variicola 150416_01 BRUS	2.46	244366
5 (+++)	Klebsiella variicola TC 4131 MCW	2.46	244366
6 (+++)	Klebsiella variicola NY 1125 MCW	2.45	244366
7 (+++)	Klebsiella variicola NY 1064 MCW	2.43	244366
8 (+++)	Klebsiella variicola CC 2123 MCW	2.36	244366
9 (+++)	Klebsiella variicola 37924 PFM	2.34	244366
10 (+++)	Klebsiella variicola 150519_01 SAH	2.32	244366

Sample 16



Sample Name: D5
Sample Description:
Sample ID: * C 480
Sample Creation Date/Time: 2021-01-12T12:01:56.754
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 TC_4131 MCW	2.39	244366
2 (+++)	Klebsiella variicola CC_2053 MCW	2.38	244366
3 (+++)	Klebsiella variicola 150416_01 BRUS	2.35	244366
4 (+++)	Klebsiella variicola DSM_15968T DSM	2.33	244366
5 (+++)	Klebsiella variicola 150519_03 SAH	2.31	244366
6 (+++)	Klebsiella variicola NY_1064 MCW	2.30	244366
7 (+++)	Klebsiella variicola CC_2123 MCW	2.29	244366
8 (+++)	Klebsiella variicola 37924 PFM	2.28	244366
9 (+++)	Klebsiella variicola NY_1125 MCW	2.27	244366
10 (+++)	Klebsiella variicola 150519_01 SAH	2.21	244366

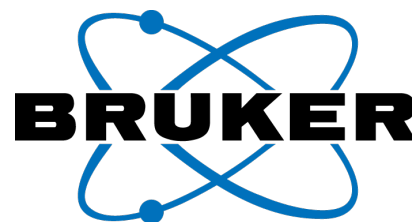
Sample 17



Sample Name: D6
Sample Description:
Sample ID: * C 332
Sample Creation Date/Time: 2021-01-12T12:01:56.755
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	2.47	72407
2 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.40	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.34	72407
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.33	573
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.27	573
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.26	72407
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.21	574
8 (+++)	Klebsiella pneumoniae 37595 PFM	2.08	573
9 (+++)	Klebsiella variicola CC_2053 MCW	2.08	244366
10 (+++)	Klebsiella variicola 150416_01 BRUS	2.07	244366

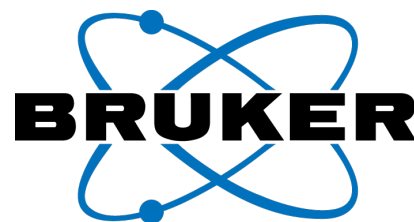
Sample 18



Sample Name: D7
Sample Description:
Sample ID: * C 539
Sample Creation Date/Time: 2021-01-12T12:01:56.755
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 ozaenae DSM 16358T DSM	2.44	574
3 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.43	72407
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.40	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.37	574
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.34	72407
7 (+++)	Klebsiella pneumoniae 37585 PFM	2.33	573
8 (+++)	Klebsiella variicola 150519_03 SAH	2.20	244366
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.20	244366
10 (+++)	Klebsiella pneumoniae 37595 PFM	2.13	573

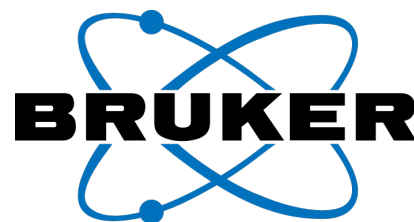
Sample 19



Sample Name: D8
Sample Description:
Sample ID: * C 549
Sample Creation Date/Time: 2021-01-12T12:01:56.756
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	2.36	244366
2 (+++)	Klebsiella variicola 150416_01 BRUS	2.34	244366
3 (+++)	Klebsiella variicola TC_4131 MCW	2.30	244366
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.30	574
5 (+++)	Klebsiella variicola 150519_03 SAH	2.28	244366
6 (+++)	Klebsiella variicola 37924 PFM	2.27	244366
7 (+++)	Klebsiella variicola NY_1064 MCW	2.25	244366
8 (+++)	Klebsiella variicola DSM 15968T DSM	2.24	244366
9 (+++)	Klebsiella variicola 150519_01 SAH	2.24	244366
10 (+++)	Klebsiella variicola NY_1125 MCW	2.21	244366

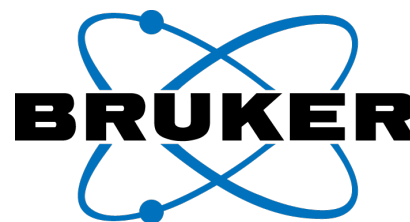
Sample 20



Sample Name: D9
Sample Description:
Sample ID: * C 583
Sample Creation Date/Time: 2021-01-12T12:01:56.756
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 variicola 150416_01 BRUS	2.33	244366
2 (+++)	Klebsiella variicola 37924 PFM	2.34	244366
3 (+++)	Klebsiella variicola CC_2053 MCW	2.21	244366
4 (+++)	Klebsiella variicola TC_4131 MCW	2.20	244366
5 (+++)	Klebsiella variicola DSM 15968T DSM	2.20	244366
6 (+++)	Klebsiella variicola 150519_03 SAH	2.17	244366
7 (+++)	Klebsiella variicola 150519_01 SAH	2.13	244366
8 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.12	574
9 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.10	573
10 (+++)	Klebsiella variicola NY_1064 MCW	2.08	244366

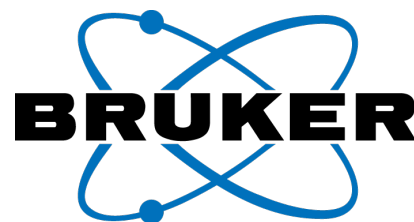
Sample 21



Sample Name: D10
Sample Description:
Sample ID: * C 636
Sample Creation Date/Time: 2021-01-12T12:01:56.757
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	2.37	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.19	72407
3 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.18	573
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.15	574
5 (+++)	Klebsiella pneumoniae 37585 PFM	2.10	573
6 (+++)	Klebsiella variicola NY_1064 MCW	2.05	244366
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.02	574
8 (+++)	Klebsiella variicola TC_4131 MCW	2.02	244366
9 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	1.97	72407
10 (+)	Klebsiella variicola CC_2053 MCW	1.91	244366

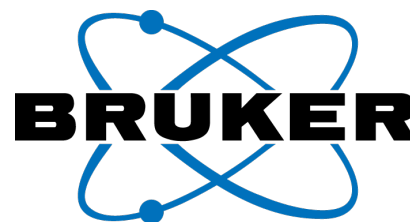
Sample 22



Sample Name: D11
Sample Description:
Sample ID: * C 647
Sample Creation Date/Time: 2021-01-12T12:01:56.757
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	2.33	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.32	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.30	574
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.29	573
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.27	574
6 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.26	72407
7 (+++)	Klebsiella pneumoniae 37585 PFM	2.24	573
8 (+++)	Klebsiella variicola 150416_01 BRUS	2.20	244366
9 (+++)	Klebsiella variicola CC_2053 MCW	2.13	244366
10 (+++)	Klebsiella variicola NY_1064 MCW	2.10	244366

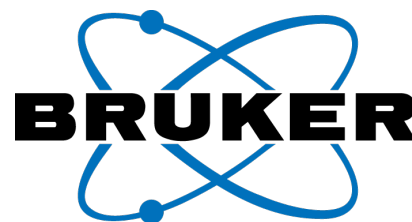
Sample 23



Sample Name: D12
Sample Description:
Sample ID: * A 467
Sample Creation Date/Time: 2021-01-12T12:01:56.758
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	2.33	574
2 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T QC DSM	2.15	72407
3 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.13	573
4 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.10	72407
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.01	574
6 (+)	Klebsiella pneumoniae 37585 PFM	1.94	573
7 (+)	Klebsiella variicola TC_4131_MCW	1.91	244366
8 (+)	Klebsiella pneumoniae 37595 PFM	1.90	573
9 (+)	Klebsiella variicola 150519_03_SAH	1.90	244366
10 (+)	Klebsiella variicola NY_1064_MCW	1.89	244366

Sample 24



Sample Name: E1
Sample Description:
Sample ID: * A 499
Sample Creation Date/Time: 2021-01-12T12:01:56.758
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	2.38	244366
2 (+++)	Klebsiella variicola DSM_15968T DSM	2.37	244366
3 (+++)	Klebsiella variicola TC_4131 MCW	2.34	244366
4 (+++)	Klebsiella variicola 150519_03 SAH	2.29	244366
5 (+++)	Klebsiella variicola NY_1125 MCW	2.29	244366
6 (+++)	Klebsiella variicola NY_1064 MCW	2.27	244366
7 (+++)	Klebsiella variicola 150416_01 BRUS	2.26	244366
8 (+++)	Klebsiella variicola 37924 PFM	2.23	244366
9 (+++)	Klebsiella variicola 150519_01 SAH	2.19	244366
10 (+++)	Klebsiella variicola NY_1036 MCW	2.16	244366

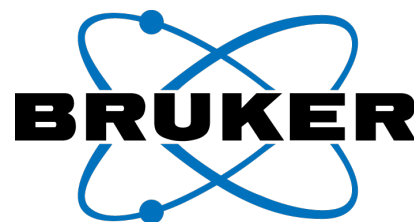
Sample 25



Sample Name: E2
Sample Description:
Sample ID: * A 501
Sample Creation Date/Time: 2021-01-12T12:01:56.759
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 ssp ozaenae DSM 16358T DSM	2.33	574
3 (+++)	Klebsiella pneumoniae 37585 PFM	2.26	573
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC DSM	2.25	72407
5 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.18	573
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.16	574
7 (+++)	Klebsiella pneumoniae 37595 PFM	2.14	573
8 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.13	72407
9 (+++)	Klebsiella variicola 150416_01 BRUS	2.08	244366
10 (+++)	Klebsiella variicola 150519_03 SAH	2.00	244366

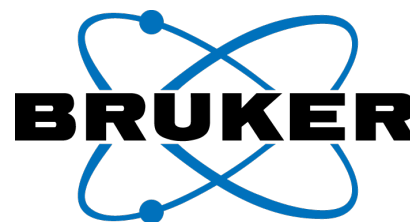
Sample 26



Sample Name: E3
Sample Description:
Sample ID: * A 504
Sample Creation Date/Time: 2021-01-12T12:01:56.759
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

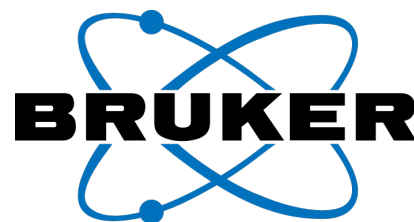
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella variicola DSM 15968T DSM	2.30	244366
2 (+++)	Klebsiella variicola CC 2053 MCW	2.29	244366
3 (+++)	Klebsiella variicola 150416_01 BRUS	2.27	244366
4 (+++)	Klebsiella variicola 37924 PFM	2.22	244366
5 (+++)	Klebsiella variicola 150519_03 SAH	2.21	244366
6 (+++)	Klebsiella variicola TC 4131 MCW	2.18	244366
7 (+++)	Klebsiella variicola NY 1125 MCW	2.15	244366
8 (+++)	Klebsiella variicola NY 1064 MCW	2.10	244366
9 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.10	72407
10 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.10	573

Sample 27



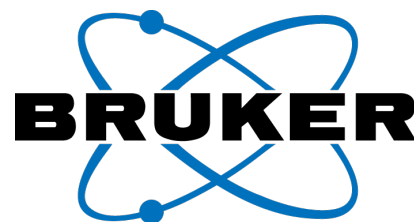
Sample Name: E4
Sample Description:
Sample ID: * B 424
Sample Creation Date/Time: 2021-01-12T12:01:56.760
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
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 (+)	Klebsiella pneumoniae RV_BA_03_B_LBK	1.95	573
2 (+)	Klebsiella variicola DSM 15968T HAM	1.93	244366
3 (+)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	1.87	72407
4 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	1.78	574
5 (+)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	1.76	72407
6 (+)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	1.73	574
7 (-)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	1.67	72407
8 (-)	Klebsiella variicola DSM 15968T_DSM	1.67	244366
9 (-)	Klebsiella variicola 37924_PFM	1.61	244366
10 (-)	Klebsiella pneumoniae 37595_PFM	1.60	573

Sample 28

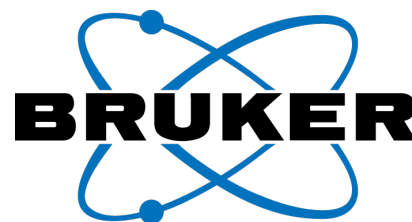
Sample Name: E5
Sample Description:
Sample ID: * B 454
Sample Creation Date/Time: 2021-01-12T12:01:56.760
Sample Type: Standard
Identification Method: MALDI Biotyper MSP Identification Standard Method 1.1
Preprocessing Method: MALDI Biotyper Preprocessing Standard Method 1.1
ACQ Method: D:\Methods\flexControlMethods\MBT_FC.par
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	2.40	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1 CHB	2.39	72407
3 (+++)	Klebsiella pneumoniae 37585 PFM	2.33	573
4 (+++)	Klebsiella pneumoniae RV_BA_03_B LBK	2.33	573
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.30	72407
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.29	574
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.23	574
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T HAM	2.21	39831
9 (+++)	Klebsiella pneumoniae 37595 PFM	2.14	573
10 (+++)	Klebsiella variicola CC_2053 MCW	2.09	244366

Sample 29

Sample Name: E6
Sample Description:
Sample ID: * B 477
Sample Creation Date/Time: 2021-01-12T12:01:56.761
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

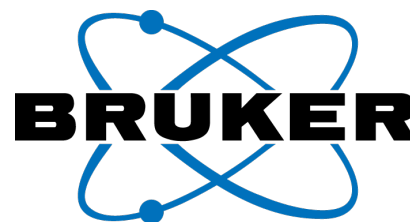
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.30	573
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.24	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.21	72407
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.16	72407
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.13	574
6 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	2.08	39831
7 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.05	574
8 (+++)	Klebsiella pneumoniae 37585_PFM	2.04	573
9 (+++)	Klebsiella variicola DSM 15968T_HAM	2.03	244366
10 (+)	Klebsiella pneumoniae 37595_PFM	1.93	573

Sample 30

Sample Name: E7
Sample Description:
Sample ID: * C 656
Sample Creation Date/Time: 2021-01-12T12:01:56.761
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

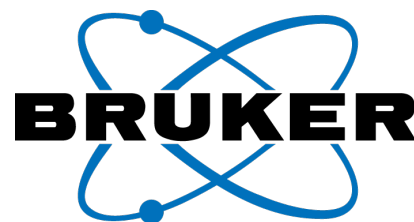
Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Klebsiella pneumoniae RV_BA_03_B_LBK	2.33	573
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295_1_CHB	2.31	72407
3 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_QC_DSM	2.25	72407
4 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_HAM	2.24	574
5 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T_HAM	2.21	72407
6 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T_DSM	2.16	574
7 (+++)	Klebsiella pneumoniae 37585_PFM	2.15	573
8 (+++)	Klebsiella pneumoniae ssp rhinoscleromatis DSM 16231T_HAM	2.15	39831
9 (+++)	Klebsiella variicola DSM 15968T_HAM	2.11	244366
10 (+)	Klebsiella pneumoniae 37595_PFM	1.97	573

Sample 31



Sample Name: E8
Sample Description:
Sample ID: * C 713
Sample Creation Date/Time: 2021-01-12T12:01:56.762
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	2.41	244366
2 (+++)	Klebsiella variicola TC_4131 MCW	2.38	244366
3 (+++)	Klebsiella variicola DSM_15968T DSM	2.32	244366
4 (+++)	Klebsiella variicola 150416_01 BRUS	2.31	244366
5 (+++)	Klebsiella variicola 150519_03 SAH	2.28	244366
6 (+++)	Klebsiella variicola CC_2123 MCW	2.28	244366
7 (+++)	Klebsiella variicola 37924 PFM	2.28	244366
8 (+++)	Klebsiella variicola NY_1125 MCW	2.28	244366
9 (+++)	Klebsiella variicola NY_1064 MCW	2.26	244366
10 (+++)	Klebsiella variicola 150519_01 SAH	2.21	244366

Sample 32

Sample Name: E9
Sample Description:
Sample ID: * C 542
Sample Creation Date/Time: 2021-01-12T12:01:56.763
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	2.36	72407
2 (+++)	Klebsiella pneumoniae ssp pneumoniae 9295 1 CHB	2.33	72407
3 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T DSM	2.32	574
4 (+++)	Klebsiella pneumoniae ssp pneumoniae DSM 30104T HAM	2.30	72407
5 (+++)	Klebsiella pneumoniae ssp ozaenae DSM 16358T HAM	2.30	574
6 (+++)	Klebsiella pneumoniae RV BA 03 B LBK	2.29	573
7 (+++)	Klebsiella pneumoniae 37585 PFM	2.28	573
8 (+++)	Klebsiella variicola 150416 01 BRUS	2.17	244366
9 (+++)	Klebsiella pneumoniae 37595 PFM	2.11	573
10 (+++)	Klebsiella variicola 150519 03 SAH	2.09	244366

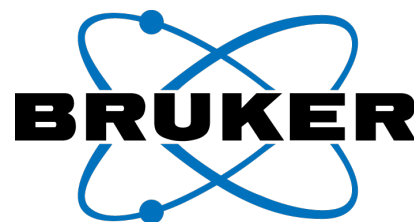
Sample 33



Sample Name: E10
Sample Description:
Sample ID: * C 717
Sample Creation Date/Time: 2021-01-12T12:01:56.763
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	2.32	244366
2 (+++)	Klebsiella variicola 150416_01 BRUS	2.30	244366
3 (+++)	Klebsiella variicola DSM_15968T DSM	2.26	244366
4 (+++)	Klebsiella variicola CC_2123 MCW	2.23	244366
5 (+++)	Klebsiella variicola NY_1125 MCW	2.21	244366
6 (+++)	Klebsiella variicola NY_1064 MCW	2.20	244366
7 (+++)	Klebsiella variicola 150519_03 SAH	2.20	244366
8 (+++)	Klebsiella variicola TC_4131 MCW	2.19	244366
9 (+++)	Klebsiella variicola NY_1036 MCW	2.17	244366
10 (+++)	Klebsiella variicola 37924 PFM	2.14	244366

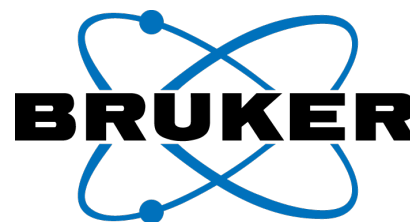
Sample 34



Sample Name: E11
Sample Description:
Sample ID: * A 434
Sample Creation Date/Time: 2021-01-12T12:01:56.764
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 (+++)	Acinetobacter Iwoffii V506 MCRF	2.06	28090
2 (+)	Acinetobacter Iwoffii 11 0038419 001 01 LGL	1.99	28090
3 (+)	Acinetobacter Iwoffii LMG 1325 LMG	1.97	28090
4 (+)	Acinetobacter Iwoffii DSM 2403T DSM	1.84	28090
5 (+)	Acinetobacter Iwoffii 2 Ring240 MHH	1.84	28090
6 (+)	Acinetobacter Iwoffii LMG 10596 LMG	1.78	28090
7 (+)	Acinetobacter Iwoffii B101 UFL	1.77	28090
8 (+)	Acinetobacter Iwoffii B138 UFL	1.72	28090
9 (+)	Acinetobacter Iwoffii 54 PIM	1.70	28090
10 (-)	Acinetobacter Iwoffii 13 PIM	1.61	28090

Sample 35



Sample Name: E12
Sample Description:
Sample ID: * A 446
Sample Creation Date/Time: 2021-01-12T12:01:56.764
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 radioresistens LMG 10614 HAM	1.84	40216
2 (+)	Acinetobacter radioresistens LMG 10611 LMG	1.73	40216
3 (-)	Acinetobacter radioresistens LMG 10610 LMG	1.64	40216
4 (-)	Acinetobacter radioresistens LMG 10609 LMG	1.62	40216
5 (-)	Acinetobacter radioresistens LMG 10615 LMG	1.61	40216
6 (-)	Acinetobacter radioresistens DSM 6976T HAM	1.57	40216
7 (-)	Acinetobacter radioresistens B381 UFL	1.49	40216
8 (-)	Acinetobacter radioresistens LMG 10612 LMG	1.47	40216
9 (-)	Agromyces lapidis HKI 324 HKJ	1.25	279574
10 (-)	Acinetobacter townneri DSM 14962T HAM	1.25	202956