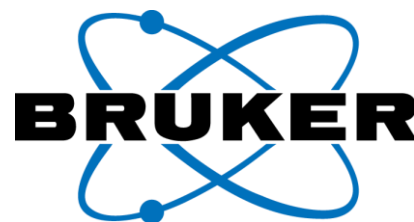


## Sample 54



**Sample Name:** E9  
**Sample Description:**  
**Sample ID:** 55  
**Sample Creation Date/Time:** 2024 07-08T11:40:32.316  
**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\_smart  
**Consistency Category (based on 2 best matches):** A  
**Applied MSP Library(ies):** BDAL / contains 11897 MSPs / ff6e1dd8-c047-4b48-bb64-1b3eae491247 / 2022-11-25T15:39:48.900

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Porphyromonas pogonae DSM 101539 DSM	<u>2.19</u>	<u>836</u>
2 (-)	<u>Pseudomonas cedrina ssp cedrina CIP 105541T HAM</u>	<u>1.25</u>	<u>76762</u>
3 (-)	Achromobacter insolitus LMG 6003T HAM	<u>1.22</u>	<u>217204</u>
4 (-)	Pseudomonas caricapapayae LMG 2152T HAM	<u>1.14</u>	<u>46678</u>
5 (-)	<u>Neisseria meningitidis 24086406 MLD</u>	<u>1.13</u>	<u>487</u>
6 (-)	Robbsia andropogonis DSM 9511T HAM	<u>1.13</u>	<u>28092</u>
7 (-)	Pseudomonas chlororaphis ssp aurantiaca CIP 106718T HAM	<u>1.13</u>	<u>333</u>
8 (-)	Brevundimonas diminuta DSM 7234T HAM	<u>1.10</u>	<u>293</u>
9 (-)	Sphingobium aromaticiconvertens DSM 12677T DSM	<u>1.10</u>	<u>365341</u>
10 (-)	Dysgonomonas capnocytophagoides CCUG 46664 CCUG	<u>1.09</u>	<u>45254</u>

## Sample 55



**Sample Name:** E10  
**Sample Description:**  
**Sample ID:** 56  
**Sample Creation Date/Time:** 2024 07-08T11:40:32.317  
**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\_smart  
**Consistency Category (based on 2 best matches):** A  
**Applied MSP Library(ies):** BDAL / contains 11897 MSPs / ff6e1dd8-c047-4b48-bb64-1b3eae491247 / 2022-11-25T15:39:48.900

Rank (Quality)	Matched Pattern	Score Value	NCBI Identifier
1 (+++)	Porphyromonas pogonae DSM 101539 DSM	<u>2.14</u>	<u>836</u>
2 (-)	Brevundimonas intermedia DSM 4732T HAM	<u>1.29</u>	<u>74315</u>
3 (-)	Paeniclostridium sordellii 1070_ATCC 9714T BOG	<u>1.28</u>	<u>1505</u>
4 (-)	<u>Flavobacterium hibernum DSM 12611T HAM</u>	<u>1.27</u>	<u>37752</u>
5 (-)	<u>Phocaeicola massiliensis ENR_0154 ENR</u>	<u>1.26</u>	<u>204516</u>
6 (-)	<u>Phocaeicola massiliensis ENR_0631 ENR</u>	<u>1.26</u>	<u>204516</u>
7 (-)	Bacteroides fragilis MB_9009_05 THL	<u>1.25</u>	<u>817</u>
8 (-)	Malassezia furfur CCUG 59936 CCUG	<u>1.23</u>	<u>55194</u>
9 (-)	Achromobacter insolitus LMG 6003T HAM	<u>1.22</u>	<u>217204</u>
10 (-)	Staphylococcus xylosus DSM 20267 DSM	<u>1.21</u>	<u>1288</u>

