

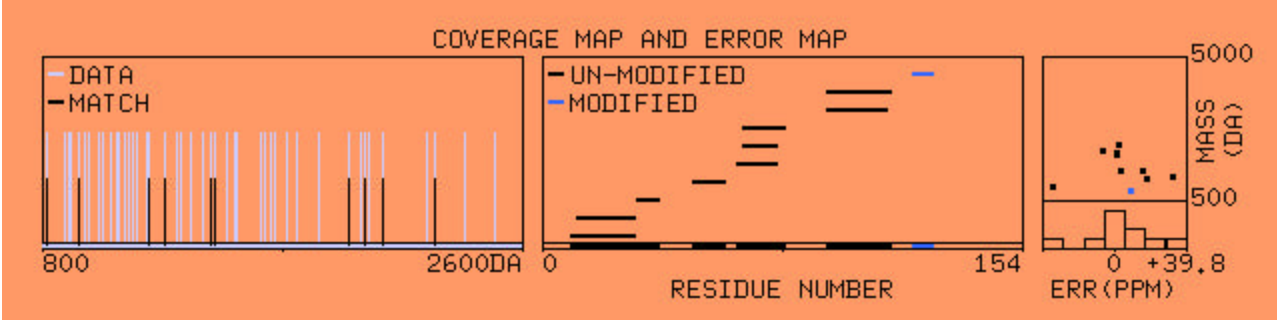
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404155246-1E00-192168001107

1. FTTSGH1181 1070076 1069615 [-1 L= 462 r=-1.186] (FTT1060c 1070067 1069615 -)  
====>ref|NP\_253619.1| 50S ribosomal protein L9 [Pseudomonas aeruginosa  
PAO1]gb|AAG08317.1| 50S ribosomal protein L9 [Pseudomonas aeruginosa PAO1] Score = 157  
bits (396), Expect = 1e-37====>ref|YP\_170041.1| 50S ribosomal protein L9 [Francisella  
tularensis subsp. tularensisSchu 4] Score = 287 bits (734), Expect = 8e-77

Sample ID : Schu4 250 [Pass:0]  
Measured peptides : 46  
Matched peptides : 10  
Min. sequence coverage: 55%



Note: click on the  symbol to change column format.

Measured	Avg/	Computed	Error	 Residues	Missed		
Mass(M)	Mono	Mass	 (ppm)	Start	To	Cut	Peptide sequence
816.423	M	816.416	9	120	126	0	MPEGVIR
							(1)+O@M;
934.496	M	934.527	-33	31	38	0	NFLIPFGK
1203.671	M	1203.649	18	65	76	0	FEAAVAVADAIK
1262.691	M	1262.650	32	49	59	1	AFEAQKAELEK
1430.811	M	1430.788	16	63	76	1	ARFEAAVAVADAIK
1446.776	M	1446.771	3	65	78	1	FEAAVAVADAIKDK
1949.971	M	1949.968	1	92	111	0	LFGSVGTAEVAEAVSNQSGK
2012.107	M	2012.104	1	12	30	0	VENLGVLGDIVNVKPGYAR
2078.051	M	2078.063	-6	92	112	1	LFGSVGTAEVAEAVSNQSGKK
2269.249	M	2269.242	3	10	30	1	EKVENLGVLGDIVNVKPGYAR

Unmatched Monoisotopic Masses:

882.494	903.437	910.462	962.433	973.538	1012.556	1030.597	1057.563	1082.605
1087.594	1108.583	1125.618	1140.612	1158.668	1194.662	1302.724	1304.703	1323.664
1357.717	1405.707	1490.863	1493.769	1523.825	1533.796	1621.904	1638.877	1657.849
1674.926	1716.880	1758.905	1838.956	1993.998	2025.041	2240.150	2383.953	2501.218

Search again using unmatched masses:

Search again

 in 

All taxa