

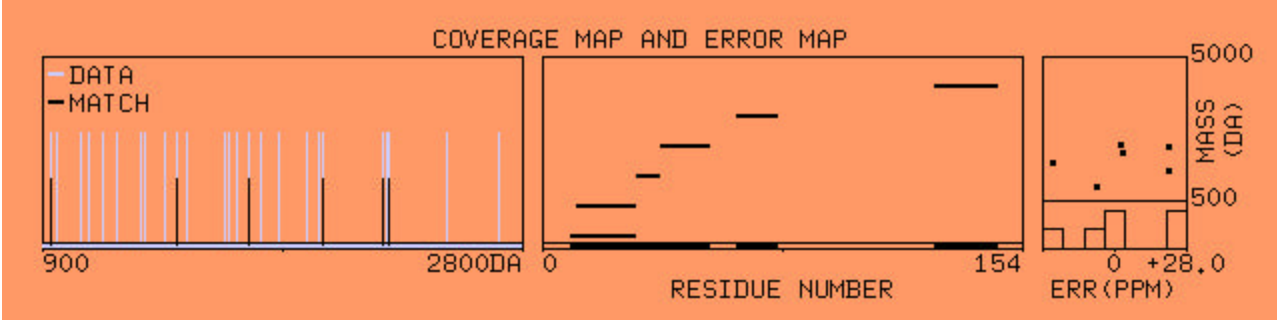
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404151141-1E60-192168001107

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 242 [Pass:0]
Measured peptides : 27
Matched peptides : 6
Min. sequence coverage: 51%



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
934.521	M	934.527	-7	31	38	0	NFLIPFGK
1430.818	M	1430.788	21	63	76	1	ARFEAAVAVADAIK
1716.875	M	1716.915	-23	39	54	1	AVQATQANIKAFEAQK
2012.111	M	2012.104	3	12	30	0	VENLGVLGDIVNVKPGYAR
2250.152	M	2250.104	21	127	146	0	SIGEFELTVHVYTDVDADIK
2269.248	M	2269.242	3	10	30	1	EKVENLGVLGDIVNVKPGYAR

Unmatched Monoisotopic Masses:

962.436	1057.622	1082.626	1087.566	1140.667	1199.681	1288.717	1308.721	1383.812
1434.781	1470.792	1621.922	1638.873	1674.939	1765.934	1838.944	1951.964	1994.006
2262.156	2501.251	2705.046						

Search again using unmatched masses:

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