

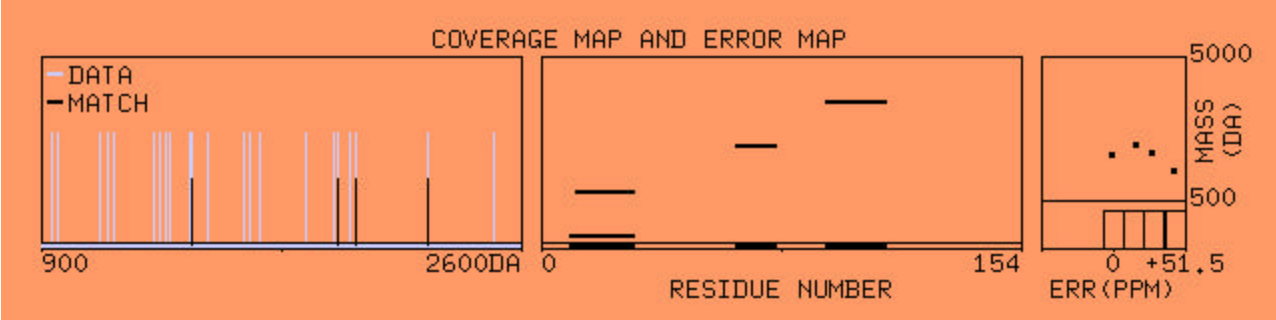
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404123109-1928-192168001107

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 235 [Pass:0]
Measured peptides : 23
Matched peptides : 4
Min. sequence coverage: 36%



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
1430.849	M	1430.788	43	63	76	1 ARFEAAVAVADAIK
1949.968	M	1949.968	-0	92	111	0 LFGSVGTAEVAEAVSNQSGK
2012.160	M	2012.104	28	12	30	0 VENLGVLGDIVNVKPGYAR
2269.279	M	2269.242	16	10	30	1 EKVENLGVLGDIVNVKPGYAR

Unmatched Monoisotopic Masses:

935.478 962.444 1109.620 1140.661 1158.685 1302.768 1323.717 1340.757 1357.777
1424.789 1434.802 1493.779 1621.954 1638.911 1674.971 1838.957 1940.935 1994.033
2501.236

Search again using unmatched masses:

Search again in All taxa