

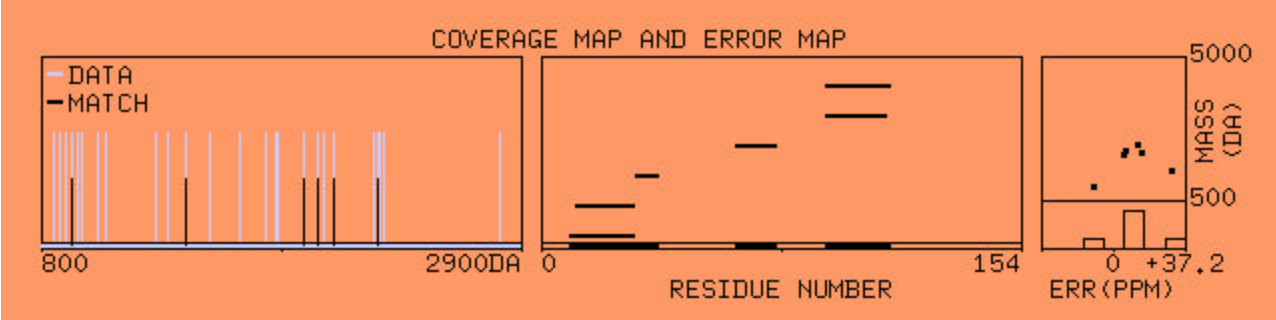
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404164529-1E98-192168001107

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 268 [Pass:0]
Measured peptides : 26
Matched peptides : 6
Min. sequence coverage: 42%



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
934.518	M	934.527	-10	31	38	0	NFLIPFGK
1430.832	M	1430.788	31	63	76	1	ARFEAAVAVADAIK
1949.979	M	1949.968	6	92	111	0	LFGSVGTAEVAEAVSNQSGK
2012.135	M	2012.104	15	12	30	0	VENLGVLGDIVNVKPGYAR
2078.077	M	2078.063	7	92	112	1	LFGSVGTAEVAEAVSNQSGKK
2269.270	M	2269.242	12	10	30	1	EKVENLGVLGDIVNVKPGYAR

Unmatched Monoisotopic Masses:

855.385 882.529 910.458 962.442 978.450 1050.569 1086.586 1305.694 1352.695
1535.915 1674.937 1784.831 1827.909 1835.608 2041.137 2252.243 2255.512 2286.298
2298.280 2807.302

Search again using unmatched masses:

Search again in

All taxa