

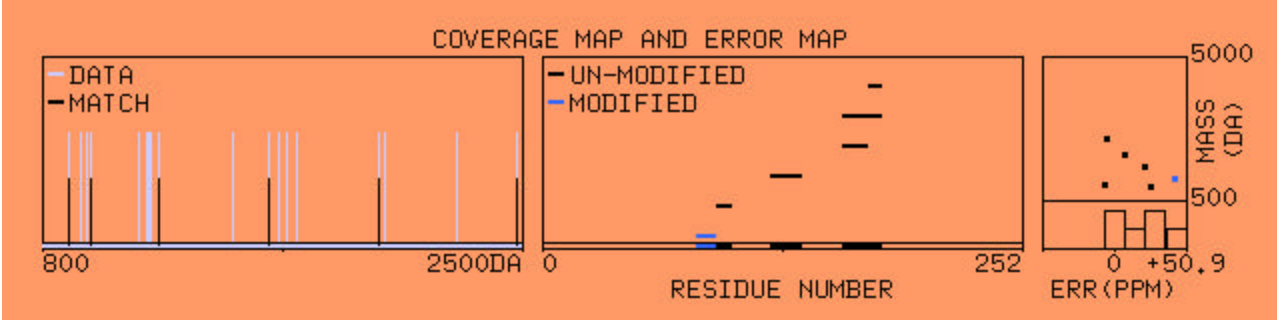
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404161937-1E00-192168001107

1. FTTSGH1267 1137666 1138421 [+3 L= 756 r=-1.232] (FTT1127 1137666 1138421 +)
====>ref|ZP_00521456.1| Rhodanese-like [Solibacter usitatus Ellin6076]gb|EAM59572.1|
Rhodanese-like [Solibacter usitatus Ellin6076] Score = 140 bits (352), Expect =
5e-32====>ref|YP_170100.1| Rhodanese-like family protein [Francisella tularensis
subsp.tularensis Schu 4] Score = 509 bits (1310), Expect = e-143

Sample ID : Schu4 259 [Pass:0]
Measured peptides : 18
Matched peptides : 6
Min. sequence coverage: 23%



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
895.390	M	895.367	26	172	178	0	EDSEFN
971.570	M	971.576	-6	92	100	1	SALVADKLR
1215.535	M	1215.484	42	82	91	0	MYFYCGGGFR
							(1)+O@M;
1603.855	M	1603.820	22	158	171	0	INSQELDGIVFDVR
1993.020	M	1993.005	8	121	137	0	GYPTVSPNQFRPNEFLK
2481.162	M	2481.176	-6	158	178	1	INSQELDGIVFDVREDSEFN

Unmatched Monoisotopic Masses:

935.505 962.459 1144.613 1174.750 1178.641 1182.539 1479.785 1638.919 1667.925
1703.527 2013.129 2270.265

Search again using unmatched masses:

Search again in All taxa