

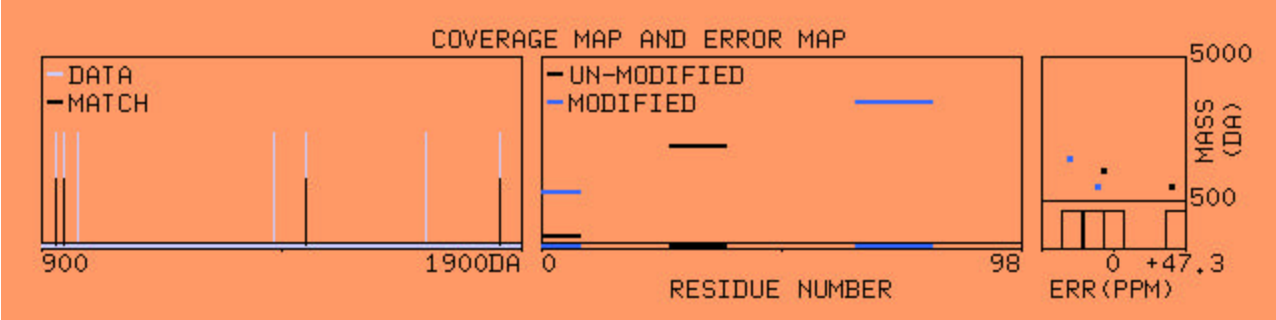
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404165959-1DBC-192168001107

1. FTTSGH1445 1301196 1300903 [-1 L= 294 r=-1.217] (FTT1281c 1301196 1300903 -)
=====>emb|CAA34495.1| ORF102 [Pseudomonas putida]pir||T01754 hypothetical protein 102
- Pseudomonas putida Score = 107 bits (266), Expect = 1e-22====>ref|YP_170237.1|
Sigma-54 modulation protein [Francisella tularensis subsp.tularensis Schu 4] Score = 198 bits
(504), Expect = 4e-50

Sample ID : Schu4 273 [Pass:0]
Measured peptides : 7
Matched peptides : 4
Min. sequence coverage: 37%



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
931.527	M	931.490	39	1	8	0	MNIQITGR
947.476	M	947.485	-10	1	8	0	MNIQITGR
							(1)+O@M;
1452.679	M	1452.688	-6	27	38	0	VEHYFDNITSTK
1854.803	M	1854.855	-28	65	80	1	AEDKDLAAIDMLEDK
							(1)+O@M;

Unmatched Monoisotopic Masses:

975.295 1387.741 1703.606

Search again using unmatched masses:

Search again in All taxa