

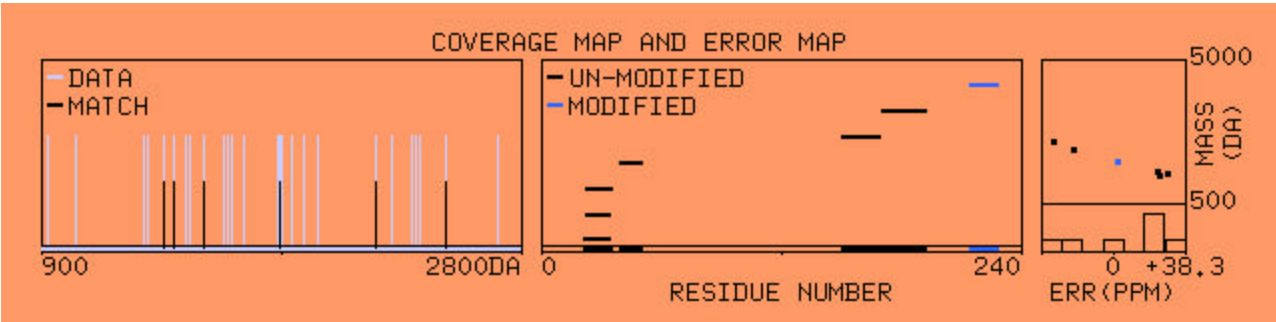
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404162736-1F0C-192168001107

1. FTTSGH1358 1220376 1219657 [-1 L= 720 r=-1.229] (FTT1201 1220376 1219657 -)
====>gb|AAF96951.1| oxidoreductase, short-chain dehydrogenase/reductase family
[Vibrio cholerae O1 biovar eltor str. N16961] Score = 337 bits (864), Expect =
2e-91====>ref|YP_170160.1| Oxidoreductase, short-chain dehydrogenase family
protein[Francisella tularensis subsp. tularensis Schu 4] Score = 463 bits (1192), Expect =
e-129

Sample ID : Schu4 262 [Pass:0]
Measured peptides : 28
Matched peptides : 6
Min. sequence coverage: 30%



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
1384.780	M	1384.746	25	23	35	0	FSEAGHSLLLVGR
1423.844	M	1423.803	29	40	51	0	IEELNLPNTLIR
1540.883	M	1540.847	24	23	36	1	FSEAGHSLLLVGR
1540.883	M	1540.847	24	22	35	1	RFSEAGHSLLLVGR
1844.858	M	1844.854	2	215	229	0	TAMFMYSQPQNINIR
(2)+O@M:							
2221.051	M	2221.097	-21	151	170	1	FAVSSISSETVREEVANDNVR
2500.210	M	2500.290	-32	171	193	1	IMTICPGAVETELLSHTTSEKIK

Unmatched Monoisotopic Masses:

930.556 1036.617 1307.703 1320.658 1383.740 1473.859 1487.795 1621.839 1638.897
1657.825 1703.315 1707.802 1838.945 1851.941 1893.068 1940.956 1993.968 2286.131
2369.002 2383.923 2398.987 2705.111

Search again using unmatched masses:

Search again in All taxa