

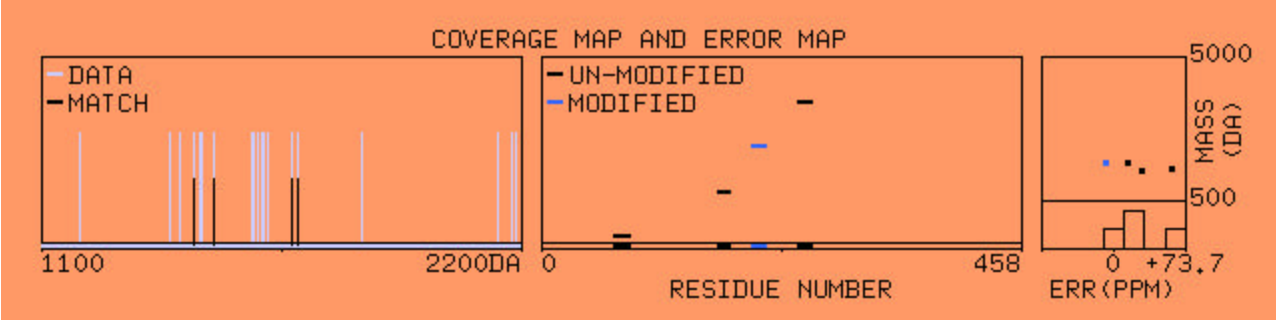
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330144250-1D10-192168001107

1. FTTSGH0069 64799 66172 [+2 L=1374 r=-1.243] (FTT0064 64799 66172 +)
====>ref|ZP_00565839.1| ATP synthase F1, beta subunit [Methylobacillus flagellatus
KT]gb|EAN02168.1| ATP synthase F1, beta subunit [Methylobacillus flagellatus KT] Score =
756 bits (1951), Expect = 0.0====>ref|YP_169139.1| ATP synthase beta chain [Francisella
tularensis subsp. tularensisSchu 4] Score = 894 bits (2309), Expect = 0.0

Sample ID : Schu4 163 [Pass:0]
Measured peptides : 19
Matched peptides : 4
Min. sequence coverage: 13%



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
1449.826	M	1449.782	30	246	259	0	YTLAGTEVSALLGR
1493.781	M	1493.689	61	168	181	0	EHSGYSVFAGVGER
1675.786	M	1675.798	-7	201	215	0	VSLVYGQMNEPPGNR
							(1)+O@M;
1688.920	M	1688.895	15	69	85	0	NTNAPISVPVGHGTLGR

Unmatched Monoisotopic Masses:

1191.443 1395.865 1417.953 1464.372 1470.139 1582.912 1586.001 1595.421 1605.152
1612.186 1619.024 1834.327 2149.564 2179.554 2189.172

Search again using unmatched masses:

Search again

 in

All taxa