

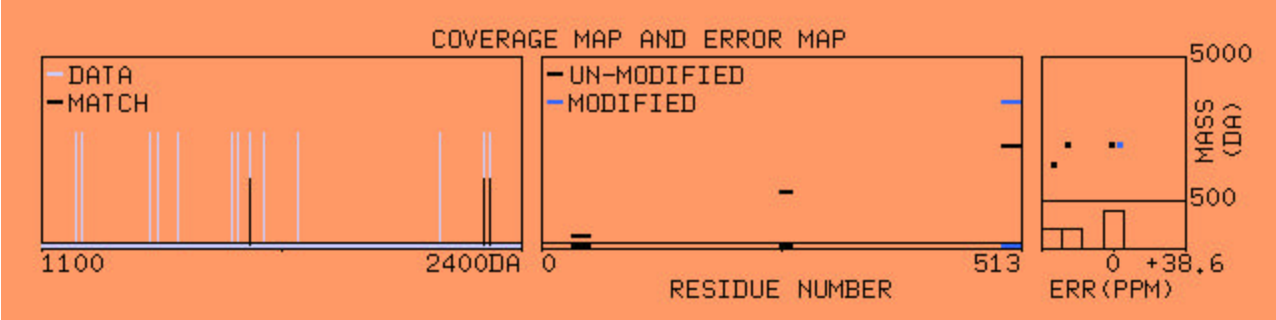
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330140252-1CBC-192168001107

1. FTTSGH0212 213991 212453 [-2 L=1539 r=-1.235] (FTT0195 213991 212453 -)
=====>ref|XP_392523.2| PREDICTED: similar to ENSANGP00000014913 [Apis mellifera]Length = 626 Score = 440 bits (1132), Expect = e-122=====>ref|YP_169256.1| L-glutaminase [Francisella tularensis subsp. tularensis Schu 4]emb|CAG44828.1| L-glutaminase [Francisella tularensis subsp. tularensis SCHU S4] Score = 1042 bits (2695), Expect = 0.0

Sample ID : Schu4 151 [Pass:0]
Measured peptides : 14
Matched peptides : 4
Min. sequence coverage: 11%



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
1665.739	M	1665.793	-32	256	269	1	ETADARNYALAHFMK
2296.154	M	2296.208	-24	493	512	0	KPLDDAIMNNNISIIELLDR
2312.212	M	2312.204	4	493	512	0	KPLDDAIMNNNISIIELLDR
							(1)+O@M;
2313.205	M	2313.205	-0	34	54	1	TDILDILVESGISNDDPRITK

Unmatched Monoisotopic Masses:

1192.221 1208.600 1395.844 1417.934 1470.102 1619.088 1634.896 1702.350 1794.828 2180.023

Search again using unmatched masses:

Search again

 in

All taxa