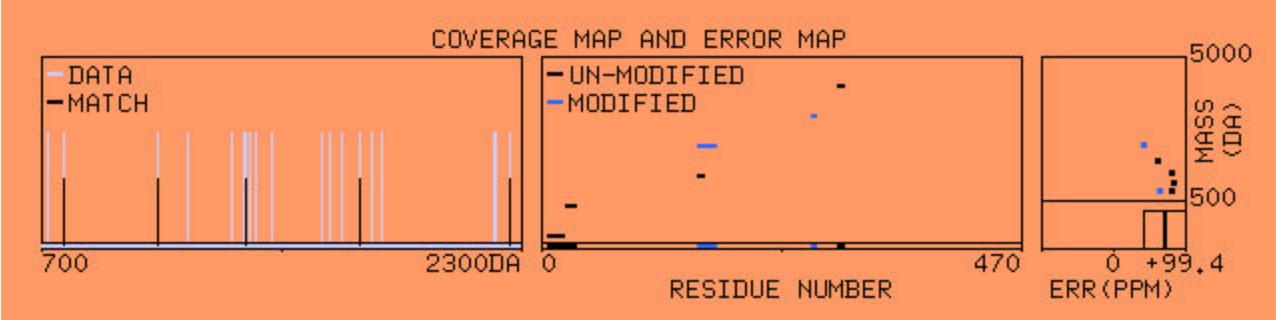



ProFound - Search Result Details



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Details for rank 1 candidate in search 20060330143315-1C90-192168001107

1. FTTSGH1671 1534332 1532923 [-1 L=1410 r=-1.210] (FTT1483c 1534332 1532923 -)
====>gb|AAO90013.1| dihydrolipoamide dehydrogenase [Coxiella burnetii RSA
493]ref|NP_819499.1| dihydrolipoamide dehydrogenase [Coxiella burnetii RSA 493] Score = 619
bits (1596), Expect = e-176====>ref|YP_170418.1| dihydrolipoamide dehydrogenase
[Francisella tularensis subsp.tularensis Schu 4] Score = 920 bits (2379), Expect = 0.0
Sample ID : Schu4 161 [Pass:0]
Measured peptides : 19
Matched peptides : 5
Min. sequence coverage: 12%



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
774.491	M	774.427	83	291	297	0	GFIPVDK
774.491	M	774.442	64	265	271	0	VLMAIGR
(1)+O@M;							
1086.624	M	1086.534	83	153	161	0	LPFVPEDDR
1382.887	M	1382.776	80	24	36	0	AADLGLEVV LVER
1765.024	M	1764.915	62	6	23	0	TQVVVLGSGPGGYSA AFR
2261.223	M	2261.124	44	153	172	1	LPFVPEDDRIIDSTGALEMK
(1)+O@M;							

Unmatched Monoisotopic Masses:

720.162	1191.471	1337.883	1373.997	1395.894	1417.998	1470.142	1634.994	1666.863
1702.442	1800.162	1834.408	2207.661	2217.037				

Search again using unmatched masses:

Search again

 in

All taxa