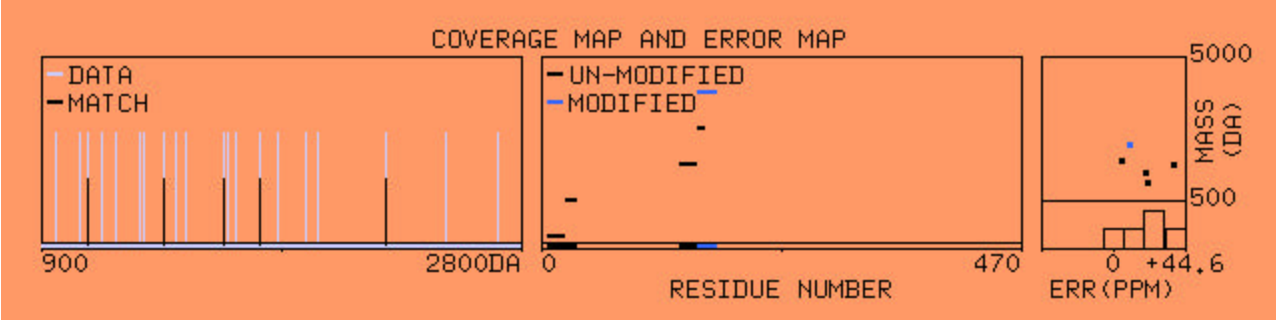



ProFound - Search Result Details



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Details for rank 1 candidate in search 20060404151226-0418-192168001107

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 242 [Pass:1]
Measured peptides : 21
Matched peptides : 5
Min. sequence coverage: 14%



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
1086.558	M	1086.534	22	153	161	0	LPFVPEDDR
1382.804	M	1382.776	20	24	36	0	AADLGLEVV LVER
1620.914	M	1620.854	37	137	152	0	IAFDNCIIAAGSSVIK
1764.926	M	1764.915	6	6	23	0	TQVVVLGSGPGGYSA AFR
2261.148	M	2261.124	11	153	172	1	LPFVPEDDRIIDSTGALEMK

(1)+O@M;

Unmatched Monoisotopic Masses:

962.436 1057.622 1082.626 1140.667 1199.681 1288.717 1308.721 1434.781 1470.792
1638.873 1674.939 1838.944 1951.964 1994.006 2501.251 2705.046

Search again using unmatched masses:

Search again

 in

All taxa