

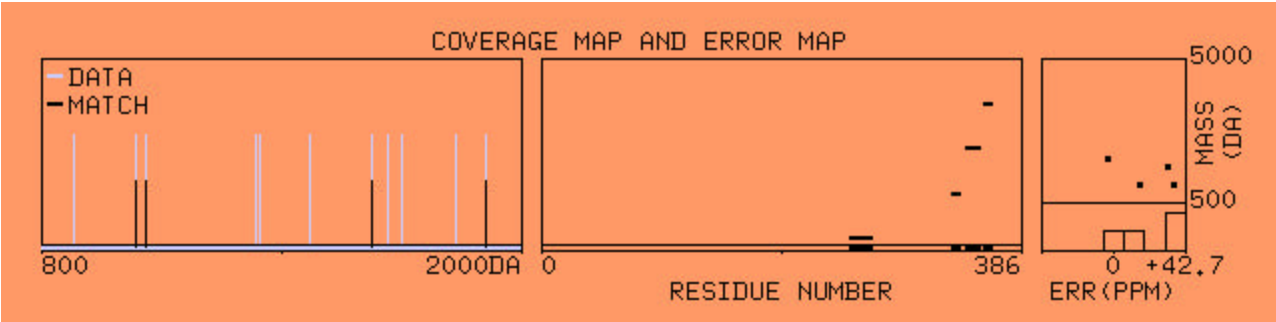
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404154930-1E00-192168001107

1. FTTSGH0162 167315 166158 [-3 L=1158 r=-1.213] (FTT0149c 167315 166158 -)
====>ref|ZP_00315922.1| COG0192: S-adenosylmethionine synthetase [Microbulbifer
degradans2-40] Score = 552 bits (1423), Expect = e-156====>ref|YP_169215.1|
S-adenosylmethionine synthetase [Francisella tularensis subsp.tularensis Schu 4] Score = 766
bits (1978), Expect = 0.0

Sample ID : Schu4 249 [Pass:0]
Measured peptides : 11
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
1037.484	M	1037.468	16	356	364	0	TSNYGHFGR
1060.629	M	1060.591	36	330	338	0	LVAEVFDLR
1626.997	M	1626.945	32	342	354	0	I IENLDLLRPIYR
1913.931	M	1913.938	-3	248	267	0	I IVDTYGGAHHGGGAFSGK

Unmatched Monoisotopic Masses:

883.479 1338.699 1348.832 1471.002 1667.820 1703.396 1835.391

Search again using unmatched masses:

Search again in All taxa