

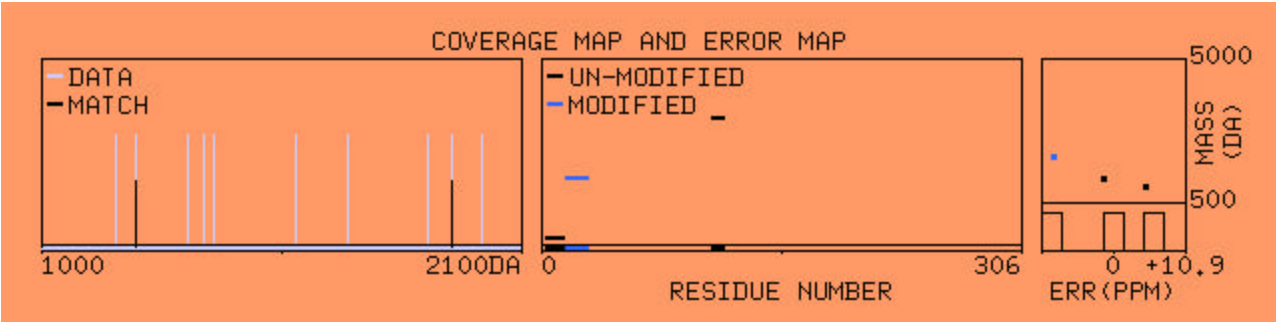
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330151914-16EC-192168001107

1. FTTSGH1548 1420127 1421044 [+2 L= 918 r=-1.243] (FTT1374 1420127 1421044 +)
=====>gb|AAU38480.1| FabD protein [Mannheimia succiniciproducens
MBEL55E]ref|YP_089065.1| FabD protein [Mannheimia succiniciproducens MBEL55E] Score =
302 bits (774), Expect = 8e-81=====>gb|AAT77114.1| probable S-malonyltransferase
[Francisella tularensis subsp.tularensis] Score = 597 bits (1540), Expect = e-169

Sample ID : Schu4 172 [Pass:0]
Measured peptides : 11
Matched peptides : 3
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
1001.555	M	1001.550	5	109	117	0	DALQLVSTR
1217.638	M	1217.640	-1	4	15	0	TAVVFPQGGSQK
1940.843	M	1940.861	-9	16	30	0	LGMLQDYENFETFR

(1)+O@M;

Unmatched Monoisotopic Masses:

1171.205 1337.833 1373.972 1395.850 1584.732 1701.712 1887.692 2009.075

Search again using unmatched masses:

Search again in All taxa