

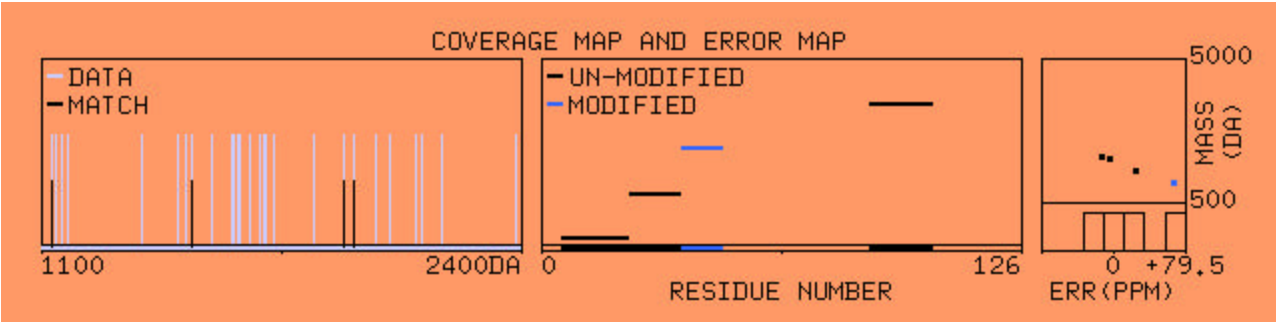
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330160002-1B78-192168001107

1. FTTSGH1506 1368817 1368440 [-2 L= 378 r=-1.214] (FTT1338 1368817 1368440 -)  
====>ref|YP\_254429.1| translation initiation inhibitor [Staphylococcus  
haemolyticusJCSC1435] Score = 145 bits (365), Expect = 5e-34====>ref|YP\_170290.1|  
translation initiation inhibitor [Francisella tularensis subsp.tularensis Schu 4] Score = 246 bits  
(629), Expect = 1e-64

Sample ID : Schu4 187 [Pass:0]  
Measured peptides : 27  
Matched peptides : 4  
Min. sequence coverage: 48%



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
1126.537	M	1126.463	66	38	48	0 ADGTMSGECIK
						(1)+O@M;
1509.832	M	1509.793	26	24	37	0 VGNFVYTSQGIGLR
1917.972	M	1917.979	-3	6	23	0 INTTNAPQAIGSYEQAIK
1944.912	M	1944.936	-12	87	103	0 LIDEVYGSFFNGNFPAR

Unmatched Monoisotopic Masses:

1139.975	1158.670	1174.649	1373.801	1469.957	1492.802	1561.586	1618.901	1622.007
1634.727	1638.894	1666.646	1694.333	1702.164	1707.864	1733.993	1838.959	2008.163
2047.456	2113.044	2129.635	2184.075	2384.020				

Search again using unmatched masses:

Search again in All taxa