

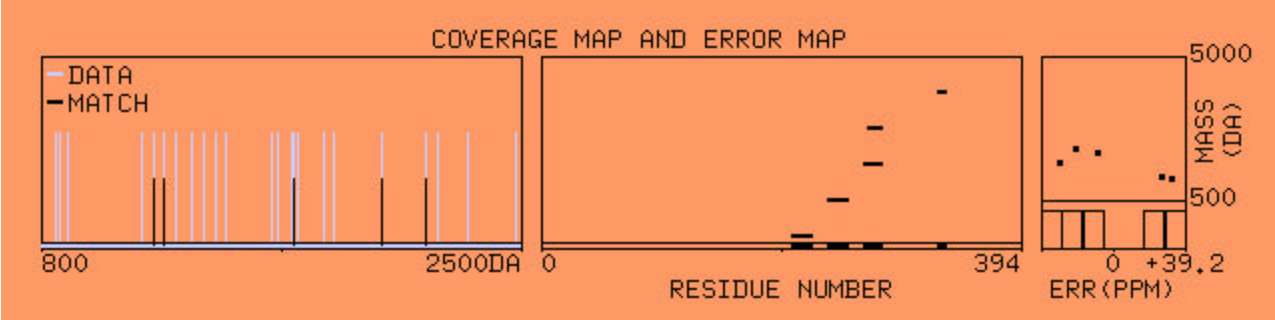
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330150330-1C90-192168001107

1. FTTSGH0150 150045 151226 [+3 L=1182 r=-1.259] (FTT0137 150045 151226 +)  
====>gb|AAM35853.1| elongation factor Tu [Xanthomonas axonopodis pv. citri str. 306]  
gb|AAM35841.1| elongation factor Tu [Xanthomonas axonopodis pv. citri str. 306] Score = 673 bits (1737), Expect = 0.0====>ref|YP\_169203.1| elongation factor Tu (EF-Tu) [Francisella tularensis subsp.tularensis Schu 4] Score = 788 bits (2034), Expect = 0.0

Sample ID : Schu4 168 [Pass:0]  
Measured peptides : 25  
Matched peptides : 5  
Min. sequence coverage: 16%



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
1198.669	M	1198.630	33	269	280	0	GEAGDNVGILVR
1232.641	M	1232.609	26	326	334	0	GYRPQFYFR
1695.876	M	1695.926	-29	265	280	1	LLDRGEAGDNVGILVR
2008.077	M	2008.094	-9	235	253	0	GVVNIGDEVEVVGIRPTQK
2162.082	M	2162.125	-20	206	224	0	DTEKPFILPIEDVFSISGR

Unmatched Monoisotopic Masses:

856.529 868.528 893.580 1158.663 1234.693 1277.731 1337.803 1375.635 1417.897  
1457.777 1621.886 1638.859 1685.859 1707.787 1800.810 1834.261 1834.872 2206.352  
2315.172 2483.123

Search again using unmatched masses:

Search again

in

All taxa