

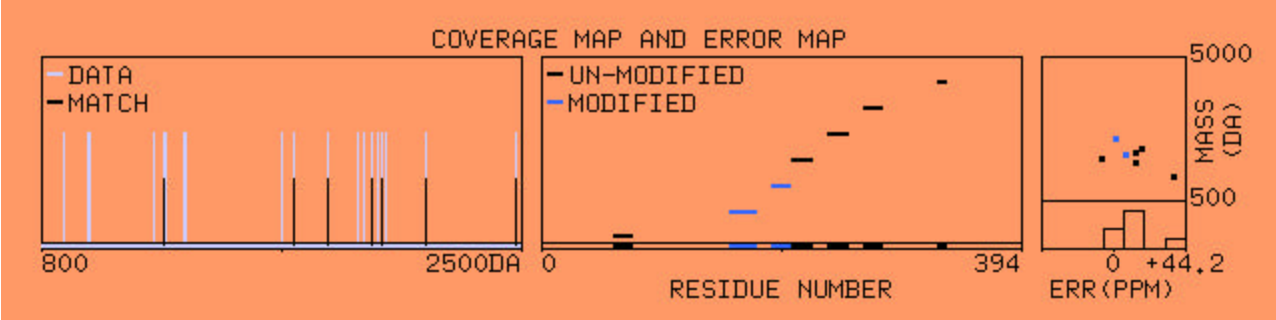
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404153651-1A78-192168001107

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 246 [Pass:1]
Measured peptides : 19
Matched peptides : 7
Min. sequence coverage: 30%



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
1232.654	M	1232.609	37	326	334	0	GYRPQFYFR
1695.949	M	1695.926	14	265	280	1	LLDRGEAGDNVGILVR
1815.863	M	1815.874	-6	60	75	0	GITINTSHVEYESPNR
1973.992	M	1973.976	8	189	205	0	IVELVQAMDDYIPAPER
							(1)+O@M;
2008.123	M	2008.094	14	235	253	0	GVVNIGDEVEVVGIRPTQK
2162.162	M	2162.125	17	206	224	0	DTEKPFILPIEDVFSISGR
2481.176	M	2481.172	2	156	177	0	ELLDQYEFPGDDTPVIMGSALR
							(1)+O@M;

Unmatched Monoisotopic Masses:

882.520 963.562 975.069 1199.706 1243.624 1309.634 1311.640 1652.918 1920.532
1940.934 1993.976 2024.025

Search again using unmatched masses:

Search again

in

All taxa