

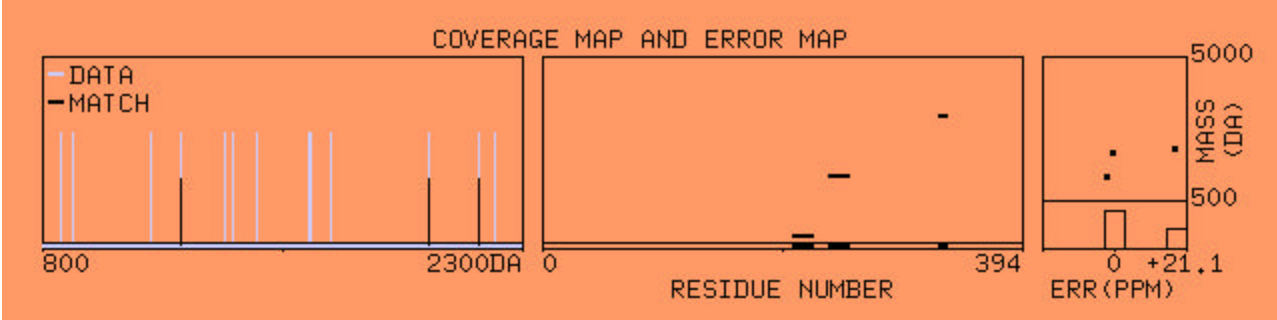
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330152319-184C-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 173 [Pass:0]  
Measured peptides : 13  
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
1232.606	M	1232.609	-2	326	334	0 GYRPQFYFR
2008.094	M	2008.094	-0	235	253	0 GVVNIGDEVEVVGIRPTQK
2162.163	M	2162.125	18	206	224	0 DTEKPFILPIEDVFSISGR

Unmatched Monoisotopic Masses:

856.520 899.527 1139.825 1373.655 1395.482 1469.771 1634.489 1638.823 1701.907  
2215.078

Search again using unmatched masses:

Search again

in

All taxa