

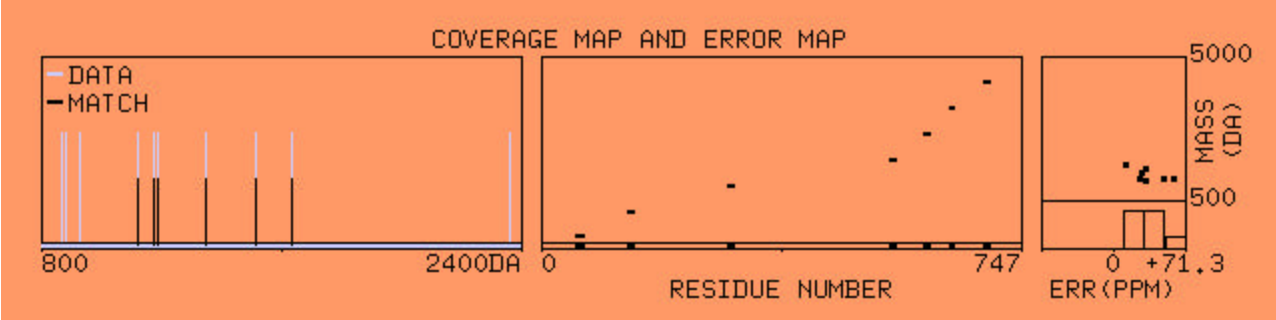
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404123855-1D2C-192168001107

1. FTTSGH1722 1584927 1582687 [-1 L=2241 r=-1.207] (FTT1526c 1584927 1582687 -)
====>ref|ZP_00314716.1| COG2838: Monomeric isocitrate dehydrogenase [Microbulbifer
degradans2-40] Score = 953 bits (2463), Expect = 0.0====>ref|YP_170456.1| isocitrate
dehydrogenase [Francisella tularensis subsp. tularensisSchu 4] Score = 1476 bits (3821),
Expect = 0.0

Sample ID : Schu4 237 [Pass:0]
Measured peptides : 10
Matched peptides : 6
Min. sequence coverage: 9%



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
1123.707	M	1123.671	32	136	146	0	VLGSAVNPVLR
1172.709	M	1172.651	50	544	554	0	NTISVTGNVLR
1172.709	M	1172.640	59	635	645	0	VLAETLSQANK
1187.662	M	1187.629	28	292	301	1	EFKELGVNPR
1349.798	M	1349.759	29	689	699	1	AKFEPIYIELK
1515.827	M	1515.779	32	596	607	0	HVEQLIEENHLR
1638.844	M	1638.825	12	55	68	0	ILANFN DYLTAEQK

Unmatched Monoisotopic Masses:

868.527 882.550 927.556 2361.198

Search again using unmatched masses:

Search again in

All taxa