

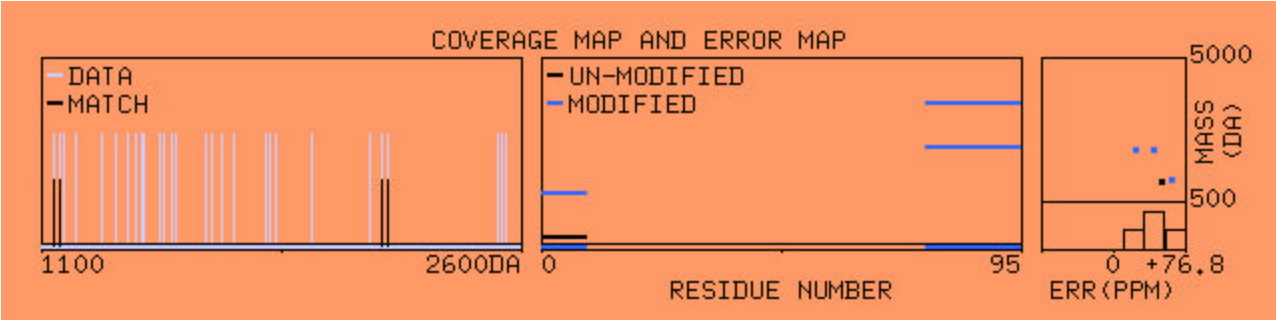
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330160314-1D10-192168001107

1. FTTSGH1924 1763981 1764265 [+2 L= 285 r=-1.205] (FTT1695 1763981 1764265 +)
====>ref|NP_954379.1| chaperonin, 10 kDa [Geobacter sulfurreducens PCA]gb|AAR36729.1|
chaperonin, 10 kDa [Geobacter sulfurreducens PCA] Score = 130 bits (328), Expect =
1e-29====>ref|YP_170600.1| Chaperonin protein, groES [Francisella tularensis subsp.
tularensisSchu 4] Score = 186 bits (471), Expect = 3e-46====>emb|CAA67359.1| groES
[Francisella tularensis]sp|P94797|CH10_FRATU 10 kDa chaperonin (Protein Cpn10) (groES
protein) Score = 184 bits (468), Expect = 6e-46

Sample ID : Schu4 188 [Pass:0]
Measured peptides : 29
Matched peptides : 4
Min. sequence coverage: 28%



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
1141.661	M	1141.602	52	1	9	0	MNIRPLQDR
1157.671	M	1157.597	64	1	9	0	MNIRPLQDR
							(1)+O@M;
2167.075	M	2167.020	26	77	95	1	VGDETLMMREEDIMGIIA
							(2)+O@M;
2183.109	M	2183.015	43	77	95	1	VGDETLMMREEDIMGIIA
							(3)+O@M;

Unmatched Monoisotopic Masses:

1174.681	1208.619	1288.066	1337.819	1373.929	1395.810	1417.895	1419.417	1470.057
1486.922	1510.910	1523.190	1618.999	1634.887	1666.780	1701.677	1702.347	1800.044
1816.724	1833.554	1945.999	2129.707	2528.295	2538.669	2555.418		

Search again using unmatched masses:

Search again in All taxa