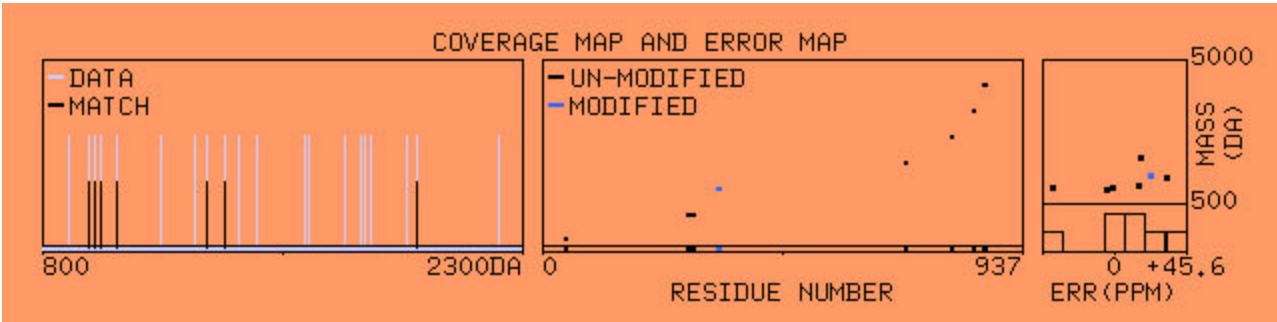



ProFound - Search Result Details



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Details for rank 1 candidate in search 20060404123540-1E90-192168001107

1. FTTSGH0092 88770 91580 [+3 L=2811 r=-1.220] (FTT0087 88770 91580 +)
====>ref|YP_004349.1| aconitate hydratase [Thermus thermophilus HB27]gb|AAS80722.1|
aconitate hydratase [Thermus thermophilus HB27] Score = 1062 bits (2746), Expect =
0.0====>ref|YP_169161.1| aconitate hydratase [Francisella tularensis subsp. tularensis Schu
4]emb|CAG44720.1| aconitate hydratase [Francisella tularensis subsp. tularensis SCHU Score =
1862 bits (4824), Expect = 0.0
Sample ID : Schu4 236 [Pass:0]
Measured peptides : 23
Matched peptides : 7
Min. sequence coverage: 8%



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
944.394	M	944.398	-5	706	713	0	DFNSYGSR
967.519	M	967.556	-38	841	848	0	NLNNIKPR
983.575	M	983.576	-1	44	51	0	VLLNQLR
1036.535	M	1036.518	16	796	804	0	AVIAESYER
1317.711	M	1317.667	33	860	871	0	TAHTTTFEALAR
1371.717	M	1371.685	23	341	350	1	EMYKEQLLFR
							(1)+O@M;
1972.974	M	1972.941	17	284	300	0	FVEYYGEGLESLSLPDR

Unmatched Monoisotopic Masses:

882.518 982.552 1172.678 1276.716 1374.713 1418.719 1470.866 1619.849 1631.851
1746.869 1794.835 1797.859 1811.891 1826.835 1940.954 2230.212

Search again using unmatched masses:

Search again

 in

All taxa