

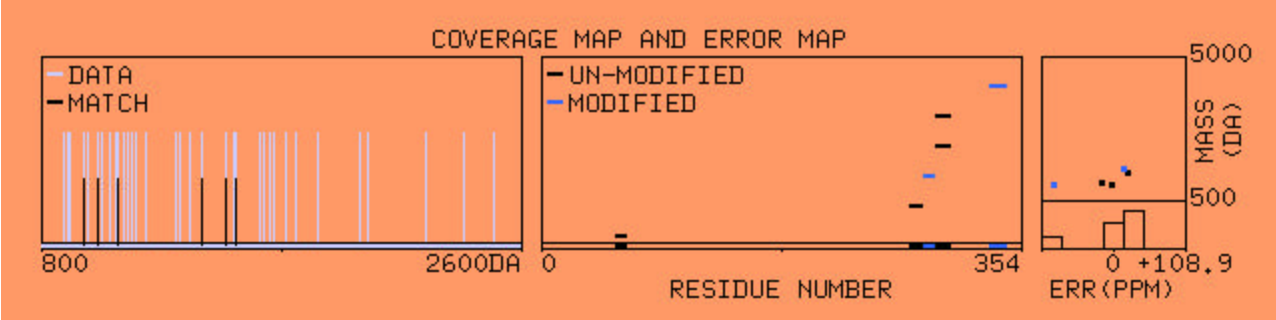
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404155330-1D68-192168001107

1. FTTSGH1540 1411305 1410244 [-1 L=1062 r=-1.228] (FTT1365c 1411305 1410244 -)  
=====>emb|CAA09871.1| fructose-1,6-bisphosphate aldolase [Pseudomonas  
stutzeri]sp|O87796|ALF\_PSEST Fructose-bisphosphate aldolase Score = 593 bits (1529),  
Expect = e-168=====>ref|YP\_170314.1| Fructose-1,6-bisphosphate aldolase [Francisella  
tularensis subsp.tularensis Schu 4] Score = 702 bits (1811), Expect = 0.0

Sample ID : Schu4 250 [Pass:1]  
Measured peptides : 36  
Matched peptides : 6  
Min. sequence coverage: 15%



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
961.425	M	961.512	-91	282	290	1	MAATGAIRR
							(1)+O@M;
1011.548	M	1011.550	-2	55	63	1	KYAGASFIR
1086.586	M	1086.603	-15	273	281	1	KINIDTDLR
1404.699	M	1404.667	23	291	302	0	FLAENPAEFDPR
1489.855	M	1489.832	16	332	343	0	IKPISLEIMFQR
							(1)+O@M;
1532.788	M	1532.762	17	291	303	1	FLAENPAEFDPRK

Unmatched Monoisotopic Masses:

882.494 903.437 910.462 973.538 1030.597 1057.563 1082.605 1108.583 1125.618  
1140.612 1158.668 1194.662 1302.724 1304.703 1323.664 1357.717 1493.769 1523.825  
1621.904 1638.877 1657.849 1674.926 1716.880 1758.905 1838.956 1993.998 2025.041  
2240.150 2383.953 2501.218

Search again using unmatched masses:

Search again in All taxa