

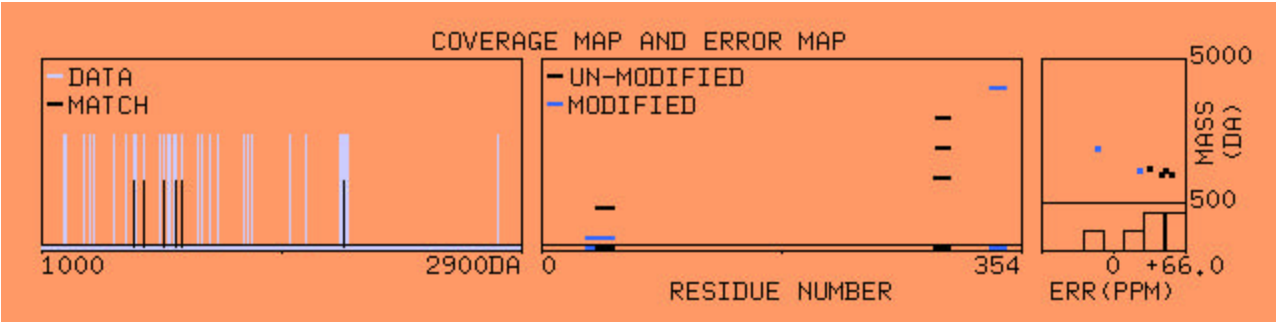
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060330144942-16EC-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 165 [Pass:0]
Measured peptides : 37
Matched peptides : 6
Min. sequence coverage: 14%



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
1367.813	M	1367.751	45	41	54	0	VNSPVILQGSAGAR
1404.744	M	1404.667	55	291	302	0	FLAENPAEFDPR
1489.870	M	1489.832	26	332	343	0	IKPISLEIMFQR
							(1)+O@M;
1532.835	M	1532.762	48	291	303	1	FLAENPAEFDPRK
1560.822	M	1560.768	35	290	302	1	RFLAENPAEFDPR
2199.100	M	2199.131	-14	33	54	1	AVMEAADKVNNSPVILQGSAGAR
							(1)+O@M;

Unmatched Monoisotopic Masses:

1088.620	1101.617	1171.315	1191.566	1208.746	1288.194	1337.941	1374.069	1470.210
1487.076	1500.467	1508.964	1523.389	1531.005	1619.201	1635.039	1666.925	1701.825
1702.484	1800.179	1816.863	1833.741	1834.461	1983.253	2047.857	2180.125	2189.296
2206.577	2214.180	2216.948	2806.394					

Search again using unmatched masses:

Search again in All taxa