

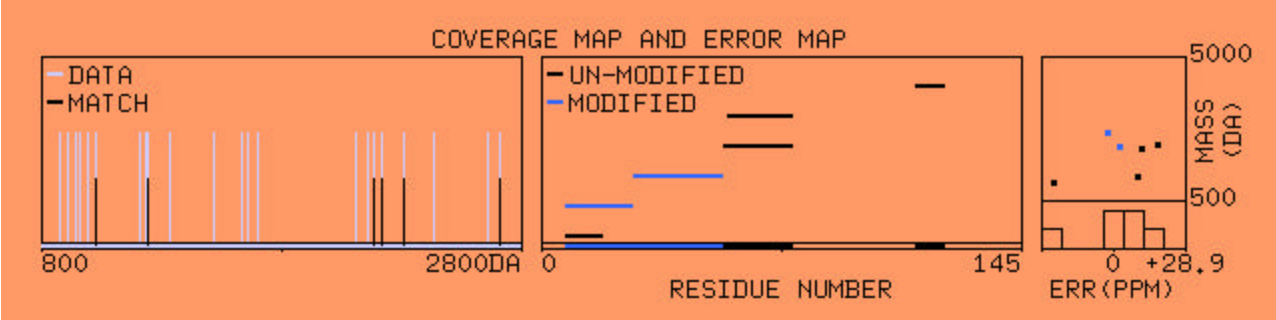
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404164807-1F10-192168001107

1. FTTSGH0070 66188 66622 [+2 L= 435 r=-1.227] (FTT0065 66188 66622 +)
====>ref|ZP_00459476.1| H+-transporting two-sector ATPase, delta/epsilon
subunit[Burkholderia cenocepacia HI2424] Score = 96.3 bits (238), Expect =
3e-19====>ref|YP_169140.1| ATP synthase epsilon chain [Francisella tularensis
subsp.tularensis Schu 4] Score = 281 bits (719), Expect = 4e-75

Sample ID : Schu4 269 [Pass:0]
Measured peptides : 24
Matched peptides : 6
Min. sequence coverage: 54%



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
1028.536	M	1028.561	-24	114	122	1	LDIEAANKR
1245.709	M	1245.696	10	8	19	0	VDVVSPLGSVFK
2185.125	M	2185.100	11	57	76	0	DQHTDVLVYVSGGIVEVTPTR
2220.151	M	2220.145	3	8	28	1	VDVVSPLGSVFKGEADMVSLR
							(1)+O@M;
2313.237	M	2313.195	18	56	76	1	KDQHTDVLVYVSGGIVEVTPTR
2714.363	M	2714.368	-2	29	55	0	GSAGEMGIAYGHTELLSTLPAGVVNVR
							(1)+O@M;

Unmatched Monoisotopic Masses:

882.484 912.473 943.488 962.449 996.547 1216.706 1239.699 1240.683 1338.813
1524.902 1636.828 1641.915 1660.933 1703.527 2110.161 2166.694 2439.117 2660.630

Search again using unmatched masses:

Search again

 in

All taxa