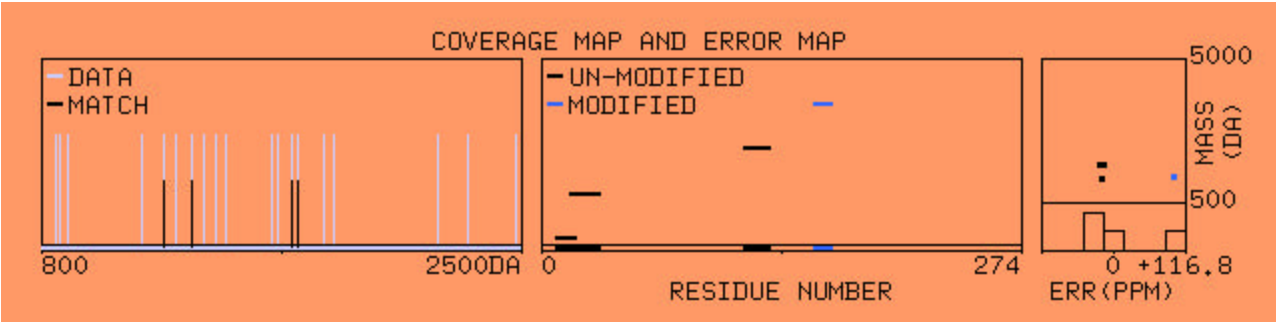



ProFound - Search Result Details



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Details for rank 1 candidate in search 20060330150508-1238-192168001107

1. FTTSGH1816 1672528 1673349 [+1 L= 822 r=-1.210] (FTT1606 1672528 1673349 +)
====>gb|AAN33522.1| septum site-determining protein MinD [Brucella suis
1330]ref|NP_699517.1| septum site-determining protein MinD [Brucella suis 1330] Score = 330
bits (845), Expect = 4e-89====>ref|YP_170522.1| septum site-determining protein MinD
[Francisella tularensis subsp.tularensis Schu 4] Score = 528 bits (1360), Expect = e-149
Sample ID : Schu4 168 [Pass:1]
Measured peptides : 20
Matched peptides : 4
Min. sequence coverage: 20%



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
1233.685	M	1233.708	-18	9	21	1 VFVVTSGKGGVGK
1336.795	M	1336.665	97	156	167	1 DSDRILGMLSSK
						(1)+O@M;
1684.851	M	1684.878	-16	17	34	1 GGVGKTTSAAVAYAFK
1706.779	M	1706.818	-23	116	131	0 NSFDIVLCDSPAGIEK

Unmatched Monoisotopic Masses:

856.529 868.528 893.580 1158.663 1277.731 1375.635 1417.897 1457.777 1621.886
1638.859 1800.810 1834.261 1834.872 2206.352 2315.172 2483.123

Search again using unmatched masses:

Search again

 in

All taxa