

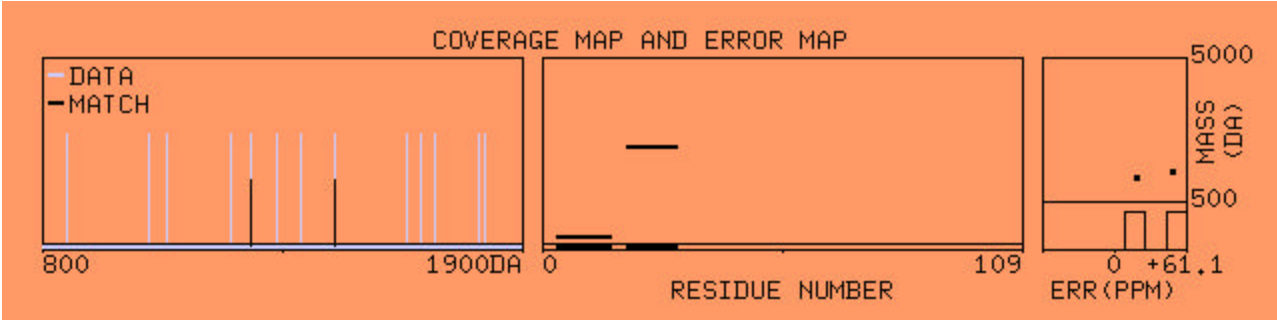
ProFound - Search Result Details


Copyright © 1997-2002 ProteoMetrics LLC

Details for rank 1 candidate in search 20060330155828-184C-192168001107

1. FTTSGH0693 649910 650236 [+2 L= 327 r=-1.257] (FTT0630 649910 650236 +)
=====>gb|AAK02990.1| Hfq [Pasteurella multocida subsp. multocida str.
Pm70]ref|NP_245843.1| Hfq [Pasteurella multocida subsp. multocida str. Pm70] Score = 120
bits (302), Expect = 1e-26=====>ref|YP_169651.1| host factor I for bacteriophage Q beta
replication [Francisellatularensis subsp. tularensis Schu 4] Score = 220 bits (560), Expect =
1e-56

Sample ID : Schu4 186 [Pass:0]
Measured peptides : 14
Matched peptides : 2
Min. sequence coverage: 23%



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
1276.763	M	1276.739	19	20	31	0	VSVSVYLVNGIK
1472.873	M	1472.798	51	4	16	0	ISSLQDPFLNALR

Unmatched Monoisotopic Masses:

856.517 1045.598 1087.695 1234.717 1337.831 1395.810 1470.080 1634.920 1666.778
1702.356 1800.044 1816.742

Search again using unmatched masses:

Search again

 in

All taxa