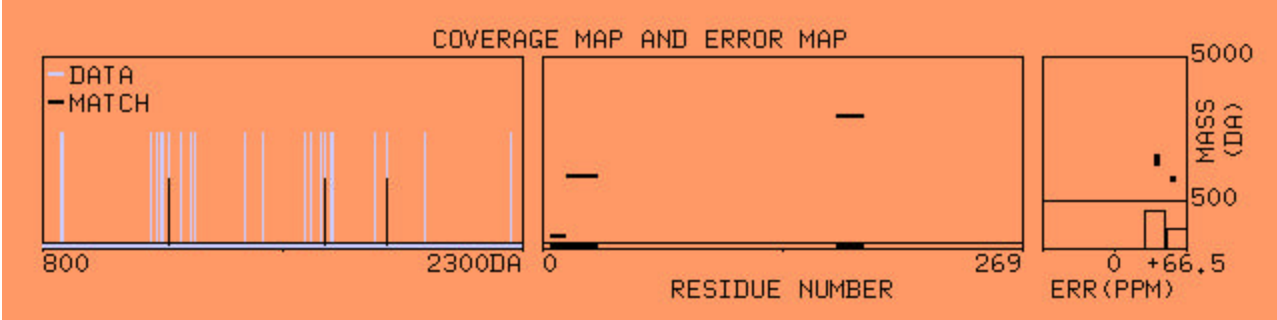



ProFound - Search Result Details



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Details for rank 1 candidate in search 20060330152551-16EC-192168001107

1. FTTSGH2019 1862587 1861781 [-2 L= 807 r=-1.224] (FTT1772c 1862587 1861781 -)  
====>gb|AAO11393.1| Tryptophan synthase alpha chain [Vibrio vulnificus  
CMCP6]ref|NP\_761866.1| Tryptophan synthase alpha chain [Vibrio vulnificus CMCP6] Score =  
356 bits (914), Expect = 4e-97====>ref|YP\_170662.1| tryptophan synthase alpha chain  
[Francisella tularensis subsp.tularensis Schu 4] Score = 526 bits (1355), Expect = e-148  
Sample ID : Schu4 174 [Pass:0]  
Measured peptides : 22  
Matched peptides : 3  
Min. sequence coverage: 16%



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
1198.689	M	1198.623	55	5	14	0	YTTLFANLEK
1685.929	M	1685.862	40	166	180	0	QISELGSGYTYLLSR
1874.041	M	1873.968	39	15	31	1	RNEGAFIPFVTIGDPNK

Unmatched Monoisotopic Masses:

856.517 864.422 1140.730 1158.696 1174.777 1179.673 1234.746 1263.768 1277.787  
1434.837 1493.806 1621.952 1638.920 1669.907 1702.167 1707.838 1838.984 1994.039  
2263.171

Search again using unmatched masses:

Search again in All taxa