

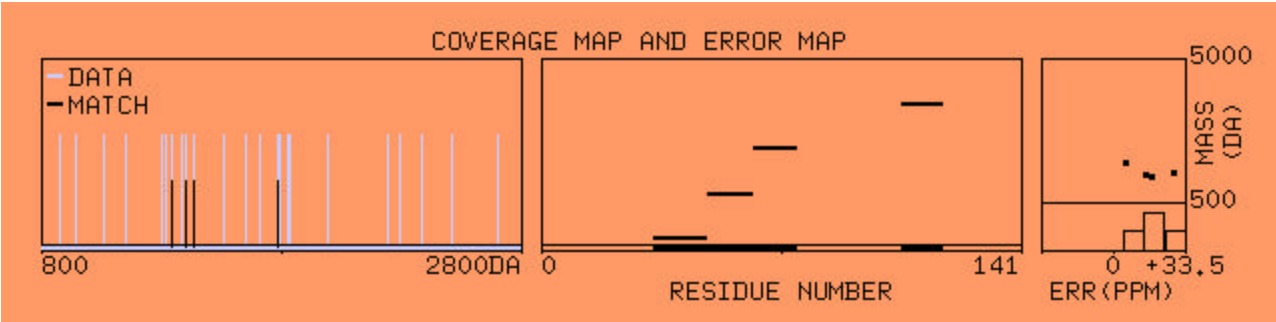
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404165711-1BE4-192168001107

1. FTTSGH1060 962552 962130 [-3 L= 423 r=-1.188] (FTT0949 962552 962130 -)
=====>gb|AAO08511.1| Unknown [Vibrio vulnificus CMCP6]ref|NP_763521.1| hypothetical
protein VV21653 [Vibrio vulnificus CMCP6] Score = 52.0 bits (123), Expect = 6e-06

Sample ID : Schu4 272 [Pass:0]
Measured peptides : 24
Matched peptides : 4
Min. sequence coverage: 38%



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
1345.784	M	1345.760	18	107	118	1	AVADKDLQVFLK
1405.741	M	1405.719	16	63	75	0	VTQDLQGSNGFIK
1433.739	M	1433.699	28	50	62	0	NTTTEQVLQASDK
1789.920	M	1789.909	6	34	49	0	NNSEPILEIVSFETAK

Unmatched Monoisotopic Masses:

882.536 948.483 1060.613 1066.528 1157.616 1307.723 1323.689 1383.736 1563.825
1657.840 1716.875 1794.865 1828.907 1838.015 1994.025 2249.105 2299.153 2383.973
2510.134 2705.166

Search again using unmatched masses:

Search again

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All taxa