

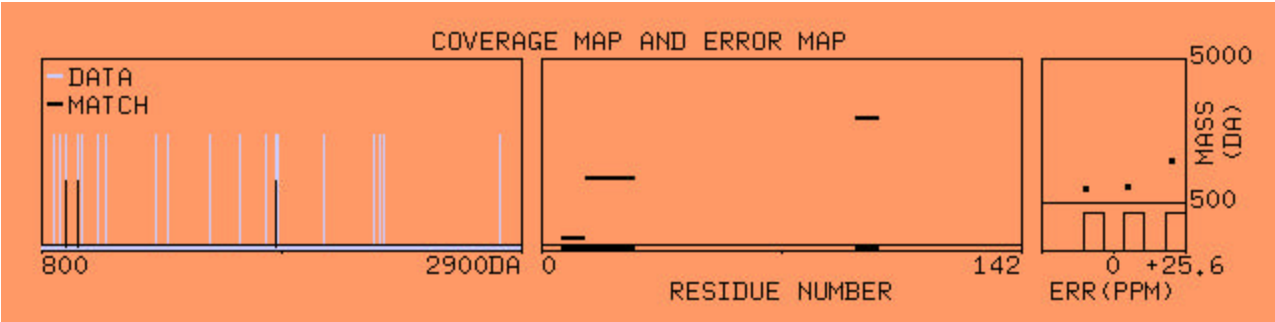
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404164606-1D0C-192168001107

1. FTTSGH2047 1882631 1883056 [+2 L= 426 r=-1.197] (FTT1794 1882631 1883056 +)
====>ref|YP_247020.1| Small heat shock protein [Rickettsia felis
URRWXCa2]gb|AAY61855.1| Small heat shock protein [Rickettsia felis URRWXCa2] Score = 116
bits (291), Expect = 2e-25====>ref|YP_170678.1| heat shock protein [Francisella tularensis
subsp. tularensis Schu4] Score = 287 bits (734), Expect = 8e-77

Sample ID : Schu4 268 [Pass:1]
Measured peptides : 20
Matched peptides : 3
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
909.450	M	909.459	-10	7	13	0	YNPFELK
961.434	M	961.429	5	94	100	0	YYGEFQR
1826.901	M	1826.862	21	14	28	0	HSINDLFDNFFSFPK

Unmatched Monoisotopic Masses:

855.385 882.529 978.450 1050.569 1086.586 1305.694 1352.695 1535.915 1674.937
1784.831 1835.608 2041.137 2252.243 2255.512 2286.298 2298.280 2807.302

Search again using unmatched masses:

Search again

 in

All taxa