

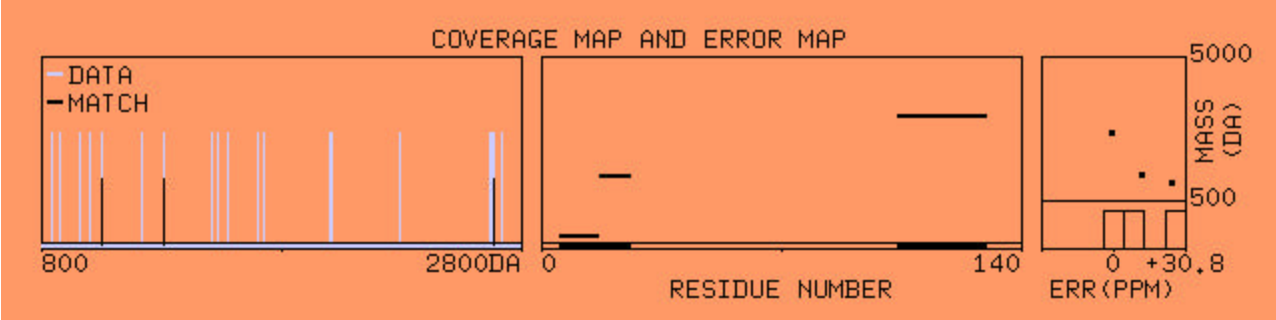
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404164316-1B98-192168001107

1. FTTSGH0402 373851 373432 [-1 L= 420 r=-1.195] (FTT0373c 373851 373432 -)
=====>gb|AAO90766.1| nucleoside diphosphate kinase [Coxiella burnetii RSA 493]ref|NP_820252.1| nucleoside diphosphate kinase [Coxiella burnetii RSA 493] Score = 214 bits (546), Expect = 5e-55=====>ref|YP_169420.1| nucleoside diphosphate kinase [Francisella tularensis subsp.tularensis Schu 4] Score = 273 bits (699), Expect = 9e-73

Sample ID : Schu4 267 [Pass:1]
Measured peptides : 20
Matched peptides : 3
Min. sequence coverage: 34%



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
1049.577	M	1049.550	26	18	26	0	NVIGEIYSR
1312.776	M	1312.760	13	6	17	0	TLSIKPDAVEK
2688.187	M	2688.189	-1	105	130	0	ADFADSIDANAVHGSDAEDTAAQEIR

Unmatched Monoisotopic Masses:

848.475 882.526 962.427 1006.545 1224.654 1509.834 1535.913 1577.731 1703.587
1729.816 1732.265 2001.997 2013.144 2299.145 2667.097 2679.152 2721.199

Search again using unmatched masses:

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