

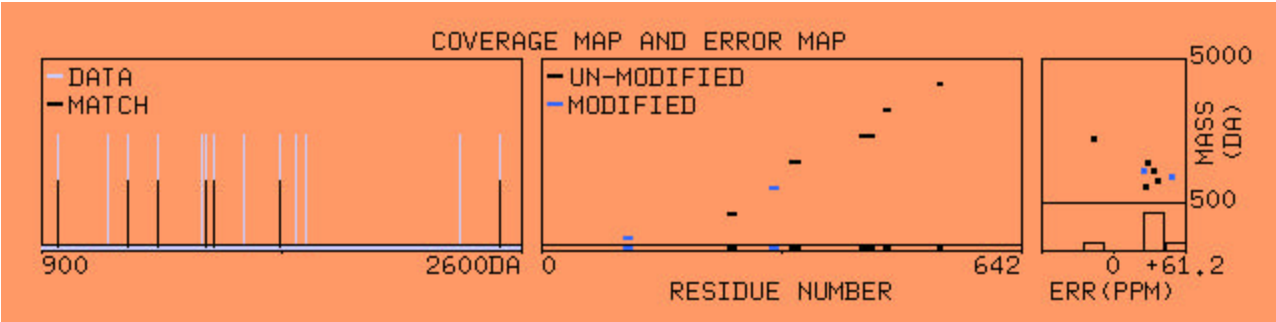
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404124128-1D40-192168001107

1. FTTSGH1433 1292668 1290743 [-2 L=1926 r=-1.202] (FTT1269c 1292668 1290743 -)
====>sp|P48205|DNAK_FRATU Chaperone protein dnaK (Heat shock protein 70) (Heat shock
70 kDaprotein) (HSP70) Score = 1232 bits (3188), Expect = 0.0====>ref|YP_170225.1|
Chaperone protein dnaK (heat shock protein family 70 protein)[Francisella tularensis subsp.
tularensis Schu 4] Score = 1239 bits (3207), Expect = 0.0

Sample ID : Schu4 238 [Pass:0]
Measured peptides : 13
Matched peptides : 7
Min. sequence coverage: 15%



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
957.529	M	957.503	27	531	538	0	FHDLVTAR
1210.680	M	1210.634	38	459	469	0	FDLADIPPAPR
1312.747	M	1312.680	51	112	123	0	MAPPQVSAEVL
							(1)+O@M;
1486.851	M	1486.798	36	334	347	0	SDITEVLLVGGQTR
1509.825	M	1509.786	26	305	317	1	AKFESLVSDLV
							(1)+O@M;
1748.967	M	1748.916	29	249	263	1	EQGIDLHNDKLALQR
2525.210	M	2525.250	-16	425	447	0	SQVFSTAEDNQPAVTIHVLQGER

Unmatched Monoisotopic Masses:

1140.746 1470.888 1619.894 1800.947 1835.298 2386.848

Search again using unmatched masses:

Search again in All taxa