

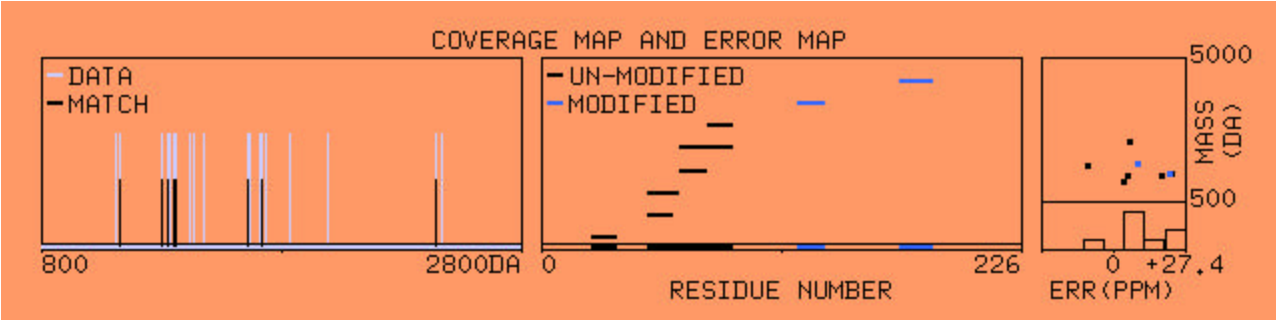
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404163001-1E00-192168001107

1. FTTSGH1526 1399353 1398676 [-1 L= 678 r=-1.261] (FTT1358c 1400863 1399322 -)
(FTT1357c 1399308 1398676 -) =====>gb|AAY51447.1| 23 kDa protein [uncultured
proteobacterium]gb|AAY51446.1| 23 kDa protein [uncultured proteobacterium] Score = 62.0
bits (149), Expect = 1e-08=====>ref|YP_170617.1| intracellular growth locus, subunit C
[Francisella tularensissubsp. tularensis Schu 4] Score = 409 bits (1050), Expect =
e-113=====>gb|AAP58964.1| IgIC [Francisella tularensis subsp. novicida]Length = 211 Score =
405 bits (1040), Expect = e-112

Sample ID : Schu4 263 [Pass:0]
Measured peptides : 21
Matched peptides : 8
Min. sequence coverage: 36%



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
1131.593	M	1131.588	4	79	90	0	ADSATAAAASVIR
1305.725	M	1305.717	6	51	62	0	LFIDSLTIAGEK
1329.702	M	1329.677	19	66	78	0	NIVAIDGGEDVTK
1353.730	M	1353.699	23	25	36	0	QQVTSGETIHVR
1365.710	M	1365.680	22	122	134	0	VSSILQASATDMK
							(1)+O@M;
1661.907	M	1661.923	-10	51	65	1	LFIDSLTIAGEKLDK
1723.886	M	1723.869	10	170	185	0	AFSISIEPTELMGVSK
							(1)+O@M;
2443.270	M	2443.254	6	66	90	1	NIVAIDGGEDVTKADSATAAAASVIR

Unmatched Monoisotopic Masses:

803.514 1115.593 1337.709 1419.114 1434.807 1475.761 1670.469 1713.028 1740.872
1834.899 1994.007 2472.269 2800.474

Search again using unmatched masses:

Search again

in

All taxa