

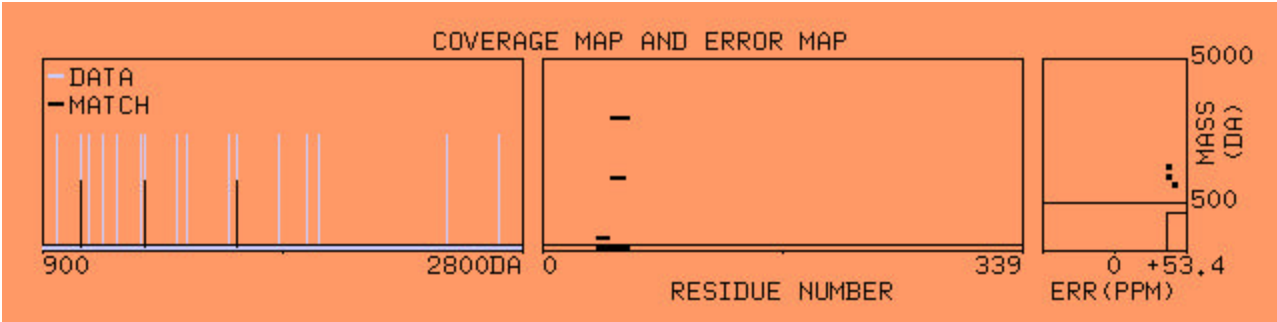
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404151301-0F3C-192168001107

1. FTTSGH0495 476164 477180 [+1 L=1017 r=-1.206] (FTT0459 476164 477180 +)
====>ref|ZP_00636494.1| Peptidase S49 [Shewanella denitrificans OS-217]gb|EAN68982.1|
Peptidase S49 [Shewanella denitrificans OS-217] Score = 302 bits (774), Expect =
1e-80====>ref|YP_169497.1| peptidase family S49 protein [Francisella tularensis
subsp.tularensis Schu 4] Score = 658 bits (1697), Expect = 0.0

Sample ID : Schu4 242 [Pass:2]
Measured peptides : 16
Matched peptides : 3
Min. sequence coverage: 7%



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
1056.614	M	1056.567	44	39	48	1	QEAAANLAKGR
1307.713	M	1307.660	40	49	59	0	LEINEVATEYK
1673.931	M	1673.862	41	49	62	1	LEINEVATEYKHTK

Unmatched Monoisotopic Masses:

962.436 1082.626 1140.667 1199.681 1288.717 1434.781 1470.792 1638.873 1838.944
1951.964 1994.006 2501.251 2705.046

Search again using unmatched masses:

Search again

in

All taxa