

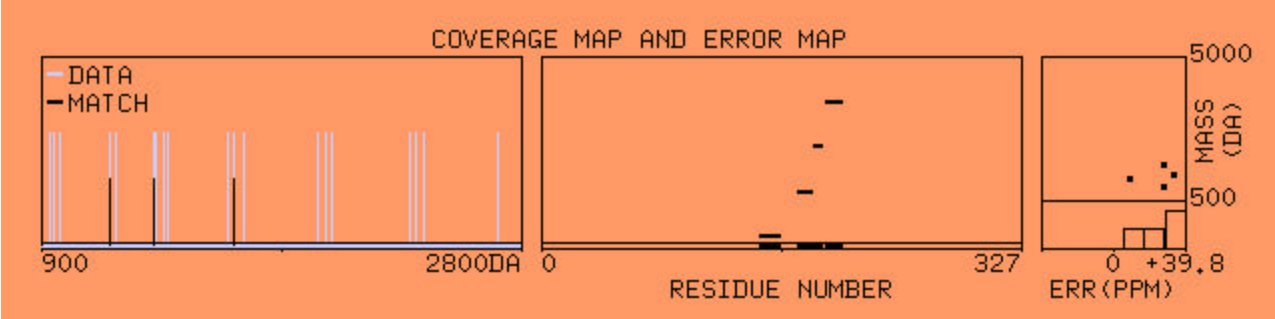
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404160651-1E60-192168001107

1. FTTSGH1646 1510900 1509920 [-2 L= 981 r=-1.256] (FTT1459c 1510888 1509920 -)
====>gb|AAM27817.1| wbpP [Pseudomonas aeruginosa]Length = 341 Score = 310 bits (795),
Expect = 3e-83====>ref|YP_170396.1| NAD dependent epimerase [Francisella tularensis
subsp. tularensisSchu 4] Score = 669 bits (1726), Expect = 0.0

Sample ID : Schu4 254 [Pass:0]
Measured peptides : 21
Matched peptides : 4
Min. sequence coverage: 14%



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
901.470	M	901.444	29	186	192	0	YFNVFGR
1169.619	M	1169.607	10	176	185	0	LYGLDTYGLR
1342.732	M	1342.688	33	194	206	0	QDPNGAYAAVIPK
1666.877	M	1666.831	28	149	163	1	EGREGNVLSPTYAFTK

Unmatched Monoisotopic Masses:

939.538 948.511 975.199 1193.662 1356.779 1387.759 1403.741 1638.959 1703.479
1994.044 2031.999 2049.005 2362.227 2383.955 2417.177 2419.188 2706.307

Search again using unmatched masses:

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All taxa