

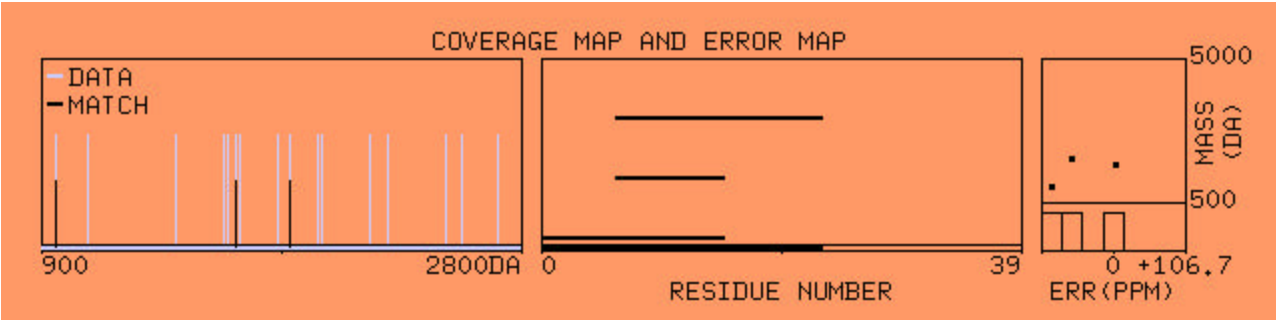
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404145401-0F3C-192168001107

1. FTTSGH1206 1086581 1086465 [-3 L= 117 r=-1.319] [OlapWith _1236 L=59 S=98] (None identical in .gff) =====>Hypothetical ORF FTTSGH1206

Sample ID : Schu4 239 [Pass:0]
Measured peptides : 16
Matched peptides : 3
Min. sequence coverage: 59%



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
961.437	M	961.523	-89	7	15	0	IFIGQGEAK
1673.960	M	1673.953	4	1	15	1	LINMLKIFIGQGEAK
1885.947	M	1886.062	-61	7	23	1	IFIGQGEAKLVQTDVLR

Unmatched Monoisotopic Masses:

1082.644	1434.779	1621.941	1638.865	1687.886	1838.927	1993.976	2013.116	2198.137
2271.255	2501.213	2567.291	2705.100					

Search again using unmatched masses:

Search again

 in

All taxa