


ProFound - Search Result Details



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Details for rank 1 candidate in search 20060404161735-1E90-192168001107

1. FTTSGH1816 1672528 1673349 [+1 L= 822 r=-1.210] (FTT1606 1672528 1673349 +)
=====>gb|AAN33522.1| septum site-determining protein MinD [Brucella suis
1330]ref|NP_699517.1| septum site-determining protein MinD [Brucella suis 1330] Score = 330
bits (845), Expect = 4e-89=====>ref|YP_170522.1| septum site-determining protein MinD
[Francisella tularensis subsp.tularensis Schu 4] Score = 528 bits (1360), Expect = e-149
Sample ID : Schu4 258 [Pass:0]
Measured peptides : 23
Matched peptides : 7
Min. sequence coverage: 26%



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
955.470	M	955.440	32	242	249	0	AYFDAVDR
1188.715	M	1188.686	24	61	70	0	VVYDLINVVR
1232.713	M	1232.676	30	39	49	0	TVVIDFDVGLR
1344.822	M	1344.787	26	60	70	1	RVVYDLINVVR
1503.845	M	1503.793	35	85	97	0	IDDLYIIPASQTR
1596.901	M	1596.862	25	174	186	1	EGEFKEIHLLLNLR
1706.835	M	1706.818	10	116	131	0	NSFDIVLCDSPAGIEK

Unmatched Monoisotopic Masses:

882.473 962.476 1307.719 1338.739 1493.786 1559.779 1604.867 1621.899 1638.906
1838.931 1994.009 2013.104 2270.254 2300.199 2501.245 2705.219

Search again using unmatched masses:

Search again

 in

All taxa