

Scores for complete sequences (score includes all domains):

Sequence	Description	Score	E-value	N
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XC-78	putative salivary p	120.6	1.9e-30	1
gi 1575479 gb AAB09540.1	FS-H precursor	115.9	4.7e-29	1
gi 1575479 gb AAB09540.1	FS-H precursor	115.9	4.7e-29	1
XC-58	similar to CF saliv	109.0	5.5e-27	1
XC-54	similar to fungal p	107.1	2.2e-26	1
cluster-169	putative secreted s	105.5	6.3e-26	1
XC-80	putative salivary s	98.6	7.7e-24	1
XC-7-90-90-50-CLU	putative salivary s	98.1	1.1e-23	1
XC-63salivary	secreted peptide of	97.0	2.2e-23	1
cluster-145	putative secreted s	90.7	1.9e-21	1
XC-44	putative salivary s	86.2	4.2e-20	1
XC-7-90-90-51	putative salivary s	81.3	1.2e-18	1
cluster-119	putative secreted s	80.7	1.9e-18	1
XC-92	putative salivary s	78.9	6.6e-18	1
XC-7-90-90-49	putative salivary s	70.4	2.4e-15	1
cluster-133	putative secreted s	64.5	1.4e-13	1
gi 4336703 gb AAD17905.1	salivary antigen 1	61.8	8.9e-13	2
gi 4336703 gb AAD17905.1	salivary antigen 1	61.8	8.9e-13	2
gi 14423664 sp Q94424 CTF1_CTEFE	Salivary antigen 1	61.8	8.9e-13	2
gi 3805687 gb AAC69105.1	FS-I [Ctenocephalid	52.8	4.6e-10	2
gi 3805687 gb AAC69105.1	FS-I [Ctenocephalid	52.8	4.6e-10	2
XC-7-90-90-48	putative salivary s	43.7	2.6e-07	1
XC-2	putative secreted s	8.5	2.2	1
gi 79324430 ref NP_001031491.1	unknown protein [Ar	6.1	3.7	1