

[illegible]

logo

			LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	
Acr00151	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00134	type II-A		LIKEFESASLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00150	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00133	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00145	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN		125
Acr00143	type II-A		LIKEFECANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00135	type II-A		LIKEFECANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00155	type II-A		LIKEFECANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00153	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00131	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00129	type II-A		LIKEFECANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00144	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN	KKDVCKALENAITVLKEKKNELI	134
Acr00149	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN	KKDVCKALENAITVLKEKKNELI	149
Acr00154	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALNNIAVLKEKKEELL	119
Acr00139	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKKTLLNNAIAVLKEKKEELL	149
Acr00130	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN	KKDVCKALENAITVLKEKKNELI	149
Acr00146	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN		125
Acr00141	type II-A		LIKEFESTNLEVLPPF	TFNRFENETHVN	KKDVCKALENAITVLKEKKNELI	149
Acr00137	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN	EKDVCKALENAITVLKEKKNELI	149
Acr00147	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVN	KKDVCKALENAITVLKEKKNELI	149
Acr00158	type II-A		LIKEFESANLEVLPPF	TFNRFENETHVDIEKDV	RKALENAITVLKEKKNELI	149
Acr00171	type II-A		YINIVETKDIIDLEI	ATD	EDKEKHDKQEVIIKSELN	123
Acr00177	type II-A		YINIVETKDIIDLEI	ATD	EDMKSDNQEIILKSELK	123
Acr00081	type I-F		IESIEANLEFIKD	EGAKFNTENAKIAQLFLEA		95
Acr00082	type I-F		IESIEANLEFIKD	EGAKFNTENAKIAQLFLEA		91
Acr00202	type II-A		LLKWLKNNRDDVIGG	TGRMYTASGNY	IANAYLEVALESSSLGSGSYMLQMRFKDYSGGQEPISGRQNRLEWIIENNLENIR	183
Acr00203	type II-A		LLKWLKNNRDDVIGG	TGRMYTASGNY	IANAYLEVALESSSLGSGSYMIQMRFKDYSGGQEPISGRQNRLEWIIENNLENIR	183
Acr00201	type II-A		LLKWLKNNRDDVIGG	TGRMYTASGNY	IANAYLEVALESSSLGSGSYMLQMRFKDYSGGQEPISGRQNRLEWIIENNLENIR	183
Acr00076	type I-F		ISDDEIKTHLQELKS	KDAHFMNESLQIAKQFCSKCET		94
Acr00075	type I-F		GMTADDLEAQLFLSA	AGAVFSESVAVDHGLKLAAADTE		95
Acr00079	type I-F		VNKRRTFKDELKFLRE	LGIKFADQALRYAVYLSGILDYQSSKLKWL		99
Acr00215	type II-A		NIMWLMKNIDENWSQSPYGES	FYSSKDIDWGYKPEGSLRVSDHWNFGQDGEHCPTAEPVDGWAVCKFENGKYHLIKKF		182
Acr00024	type I-D		DIWLEYNKVSYGG	ETTYEVKNVEEINTTEEFMNLFF		129
Acr00204	type II-A		DRITKFTAYLCIKAADGTIVNGWLASQTDMLANDWVIVE			103
consensus			*** * ** *	** * * *		

- non conserved
- similar
- ≥ 50% conserved