






logo

			
Acr00197	type II-AMNINELIR...EIKNKD.....YTAKLSGTDNSITQLIIRVNNDG..NEYVISSENESEIVEKFIS.....AFKN..GWNQE	66
Acr00198	type II-AMNINELIR...EIKNKD.....YTAKLSGTDNSITQLIIRVNNDG..NEYVISSENESEIVEKFIS.....AFKN..GWNQE	66
Acr00195	type II-AMNINELIR...EIKNKD.....YTAKLSGTDNSITQLIIRVNNDG..NEYVISSENESEIVEKFIS.....AFKN..GWNQE	66
Acr00199	type II-AGSMNINDLIR...EIKNKD.....YTVKLSGTDNSITQLIIRVNNDG..NEYVISSENESEIVEKFIS.....AFKN..GWNQE	68
Acr00192	type II-AMNINELIR...EIKNKD.....YTVKLSGTDNSITQLIIRVNNDG..NEYVISSENESEIVEKFIS.....AFKN..GWNQE	66
Acr00196	type II-AMNISELIR...EIKNKD.....YAVLELCTDDNSITKLIIIDVDNDG..NEYVISSEKNESTIAEKFAS.....TFKN..GWNKE	66
Acr00068	type I-FMSSTISDRISRSVIEAAR.....FIQSWEDADPDNLTESQV..LAASS..FAARLHEGLQATVIQRLVDES...NRDEYREFQAWEEA	77
Acr00069	type I-FMSNTISDRIVARSVIEAAR.....FIQSWEDADPDSLTDQV..LAAAG..FAARLHEGLQATVIQRLVDES...NHEEYREFKAWEEA	77
Acr00001	type I-B	LESKNLRKLLNEYEEIDINEMLNFRSIKNSGTKNDIEIFLHEKAIKFEKSSISTYVVFSEDNELGYFTIANRSLVTPKENFGIL...SKTQQKKLGNAAI	101
Acr00208	type II-A	...MADMTLRQFCERYRKGDFLAKDRETQIEAG..WYDWFCDKALAGRLAKIWLKGLTSYILDNRYVWFKNICPMVGPLYDDVRFEPLDEEQRDELYFGVA	100
Acr00222	type II-CMNVFPIPSLA.....EAYHCAG...WALGAVLGGQL..VALRYVADLGVSVPEGAAG.....VQAVRAWIES	58
Acr00111	type I-FMKYLVKEMQGAQ.....SLRDG..MVVEAKTIR.....WAKC	30
consensus		* * **** * * * * * * * * * * * * * * * * *	

logo

			
Acr00197	type II-A	YEDEE.....EFYNDMQTITLKSELN.....	87
Acr00198	type II-A	YEDEE.....EFYNDMQTITLKSELN.....	87
Acr00195	type II-A	YEDEE.....EFYNDMQTITLKSELN.....	87
Acr00199	type II-A	YEDEE.....EFYNDMQTITLKSELN.....	89
Acr00192	type II-A	YEDEE.....EFYNDMQTITLKSELN.....	87
Acr00196	type II-A	YEDEE.....EFYNDMQSITLKSELN.....	87
Acr00068	type I-F	LLNADGRVTSNPFADW..GWWYRIANVMLATASQNVGVAWGSHVHGRLMAIFQDRFQQHYEDEEC.....	140
Acr00069	type I-F	LLNADGRVASSPFADW..GWWYRIANVMLATASQNVGVTWGSRVHGRLMAIFQDKFKQRYEEQA.....	139
Acr00001	type I-B	LKNGDLMTSSFLGQLGKNVSDIENLITGREILTFAYDLFLKIKELINVKYIWLEQCNEPKLISFYQNFQFKMLESILTSEEGLKVMIMELK	193
Acr00208	type II-A	IDDKRREKKYVIFTAR..NDYENECGFNNVREVRQFINGWEDELKNEEFYKAREKKRQEMEEANNKFAEIMQRADEILWNLKED.....	182
Acr00222	type II-C	PGAAP.....VVRELQALGRVTIG..MCSGWQ.....FVEQ.....	87
Acr00111	type I-F	LAACLQAFR.....GTVMRVEDMDGTILALKVNRKWEN.....VNPDTNNS.....	71
consensus		** * * * * * * * * * * * * * * * * *	

-  non conserved
-  similar
-  ≥ 50% conserved