

logo




Accession	Type	Sequence	Length
Acr00072	I-E, I-F	MRFTSPSRTIPSICLAIIYGCQQLDRVADFINNSGEYVEDGVTFPTPGMIAGQYAFEQVKLDRKGYDTRRLQKYLSFLADAGAFFSEEEALQRADDLINNFT	0
Acr00074	I-F	0
Acr00073	I-F	0
Acr00194	II-A	0
Acr00193	II-A	0
Acr00198	II-A	0
Acr00197	II-A	0
Acr00195	II-A	0
Acr00192	II-A	0
Acr00199	II-A	0
Acr00196	II-A	0
Acr00119	I-F	0
Acr00083	I-F	MRFTSPSRTIPSICLAIIYGCQQLDRVADFINNSGEYVEDGVTFPTPGMIAGQYAFEQVKLDRKGYDTRRLQKYLSFLADAGAFFSEEEALQRADDLINNFT	100
Acr00316	VI-AMIYYIKDLKVKGKIFENLMN.....KEAVEGLITFLKKAEFEIYSRENYSKYNKWFEMWKSPTSSSLVFWKNYSFRCHLLFVIEKD	80
Acr00208	II-AMADMTLRQFCERYRKGDFLAKDRETQIEAGWYDWFCDKALAGRL	45
Acr00313	V-AMYEIKLNDTLIHQTDDRVNAFVAYRYLLRRGDLPKCENIARMYYDGKVIKTDVIDHDSVHSDEQAKVSNNDIKMAISELGVNNF	85
consensus			

logo

Accession	Type	Sequence	Length
Acr00072	type I-E, I-F	MKVPAFFAANI...LTIEQIIIEA...INNDGSAMTSAPETIAGYYAWDAATDALES...ENDLEQLTDEDDF...VA	63
Acr00074	type I-F	MKVPAFFAANI...LTIEQIIIEA...INNDGSAMTSAPETIAGYYAWDAATDALES...ENDLEQLTDEDDF...VA	63
Acr00073	type I-F	MKVPAFFAANI...LTIEQIIIEA...INNDGSAMTSAPETIAGYYAWDAATDALES...ENDLEQLTDEDDF...VA	63
Acr00194	type II-A	MNLKELVRE...IKNKDYTAKLS...GTDSNSITQLIIHVNNDGNEYGIS...ESNFESIVEK...F...VS	57
Acr00193	type II-A	MNLKELVRE...IKNKDYTAKLS...GTDSNSITQLIIHVNNDGNEYGIS...ESNFESIVEK...F...VS	57
Acr00198	type II-A	MNINELIRE...IKNKDYTAKLS...GTDSNSITQLIIRVNNDGNEYVIS...ESENESIVEK...F...IS	57
Acr00197	type II-A	MNINELIRE...IKNKDYTAKLS...GTDSNSIAQLIIRVNNDGNEYVIS...ESENESIVEK...F...IS	57
Acr00195	type II-A	MNINELIRE...IKNKDYTAKLS...GTDSNSITQLIIHVNNDGNEYVIS...ESENESIVEK...F...IS	57
Acr00192	type II-A	MNINELIRE...IKNKDYTVKLS...GTDSNSITQLIIRVNNDGNEYVIS...ESENESIVEK...F...IS	57
Acr00199	type II-A	GSMNINELIRE...IKNKDYTVKLS...GTDSNSITQLIIRVNNDGNEYVIS...ESENESIVEK...F...IS	59
Acr00196	type II-A	MNISELIRE...IKNKDYAVRLE...GTDDNSITQLIIDVDNDGNEYVIS...ESKNESIAEK...F...AS	57
Acr00119	type I-F	MTTITIAEY...VSNQKVTETIK...TMVESQIHNVNFNGEFTIE...RGDFTSIDKDE...51	51
Acr00083	type I-F	ADFGHDEGDGATFRIKAQSYRLKKYDCHALWSM...NQASLTIES...VKPETFCDESDDEDISSD...SLDLVLTSRLFNKQNEAFATLTQYYTVSEGLMTVEDLL	199
Acr00316	type VI-A	GECLGIPASVFESVLQIYLADPPAPDTKELFVEVCNLYECLADVTVVEHFEAEESAWHKLT...HNETEVSKRVYSKDDDEL...LKYIPEFLDTIATNK...KS	176
Acr00208	type II-A	AKIWGILKGITSDYILDNYRVWFKNNCPMVGPLYD...DVRFEPLDEEQRDELYFGVAIDDKRREKKYVIFTARNDYENECGFNNVREVRQFINGWEDELKNE	145
Acr00313	type V-A	KSLIKKQGYPSFN...GHINSWFTDDPVKSKTMMHND...MYVQALIRACIIEID...LYTEQLYNIKSLPYDKRPNV...VYSDQPLDPNNLDLS...EPE	176

logo

[illegible]

-  non conserved
-  similar
-  $\geq 50\%$ conserved