

logo

logo			MDEIDDLSLPMRPFIWGFAIFTPKGG	EVMHDEFEYLTHTTRAPRFTICRVVELEDMP	
Acr00259	type II-C	.	.MDEIDDLSLPMRPFIWGFAIFTPKGG.	.EVMHDEFEYLTHTTRAPRFTICRVVELEDMP.	56
Acr00260	type II-C	.	.MDEIDDLSLPMRPFIWGFAIFTPKGG.	.EVMHDEFEYLTHTTRAPRFTICRVVELEDMP.	56
Acr00261	type II-C	.	.MDEIDDLSLPMRPFIWGFAIFTPKGG.	.EVMHDEFEYLTHTTRAPRFTICRVVELEDMP.	56
Acr00267	type II-C	.	.MDEIDDLSLPMRPFIWGFAIFTPKGG.	.EVMHDEFEYLTHTTRSPRFTICRVVELEDMP.	56
Acr00248	type II-C	.	.MDEIDDLSLPMRPFIWGFAIFTPKGG.	.EVMHDEFEYLTHTTRSPRFTICRVVELEDMP.	56
Acr00100	type I-F	.	.	.MSYLISNG...QIVAKT.SATLA	19
Acr00099	type I-F	.	.	.MSYLISNG...QIIAKT.SATLA	19
Acr00097	type I-F	.	.MARLAPN.	.EDSTMSTAYLIFNS...SVAAVV.DTEIA	32
Acr00098	type I-F	.	.MVALATYNVFNCSCWLEPILL.	.RTMKMNIAVIIRNN...TVIATV.STGLP	46
Acr00312	type V-A, I-C	.	.MVGKSKIDWQSITDWTKTNAQIAECGRAYNTVCCK.	.MRGKLKGSHQGAKSPRKDKGISRP.QPHIN	63
Acr00083	type I-F	.	.MRFTSPSRPTIPSICLAIYGCGQLDRVADFINNSGEYVEDGVTFTPGMIAQQYAFAEQVKLDRKGYDTRRRLQKYLSFLADAGAFFSEEEALQRADDLI		96
Acr00311	type V-A	.	MHHTIARMNAFNKAFANAKDCYKKMQAWHLNLKPKHAFPPMQNTPLDLNGLAAIYELRGKGEDAHILSILSRILYLYGAWRNTLGIYQLDEETIKDCKEIP		100
consensus			* ** *	* ** *	

logo

[illegible]

logo

logo

Acr00259 type II-C
Acr00260 type II-C
Acr00261 type II-C
Acr00267 type II-C
Acr00248 type II-C
Acr00100 type I-F
Acr00099 type I-F
Acr00097 type I-F
Acr00098 type I-F
Acr00312 type V-A, I-C
Acr00083 type I-F
Acr00311 type V-A
consensus

128
128
128
128
128
81
81
92
107
168
239
300

logo

			SRKRLFRYFASITRSEYNTSSDYESSPMKLPIMAN	
Acr00259	type	II-C	128
Acr00260	type	II-C	128
Acr00261	type	II-C	128
Acr00267	type	II-C	128
Acr00248	type	II-C	128
Acr00100	type	I-F	81
Acr00099	type	I-F	81
Acr00097	type	I-F	92
Acr00098	type	I-F	107
Acr00312	type	V-A, I-C	168
Acr00083	type	I-F	RDAHKELPDVIRSEEITSMILQPMKLPIMAN	270
Acr00311	type	V-A	QAKEFRVRWQPAVFNNGRVSS.....	322
consensus				

- ☐ non conserved
- similar
- ≥ 50% conserved