

logo

Acr00221	type	II-CMNKTYKIGKN..AGYD..CCGLCLAAISENEATKVKYLRD.....	36
Acr00230	type	II-CMANKTYKIGKN..AGYD..CCGLCLAAISENEATKVKYLRD.....	37
Acr00223	type	II-CMTLTDWIVDRVRHNTMNVFKVPQSLADKYH..GAGYALAATVAGQLVDIVYLAD.....M	53
Acr00225	type	II-C	MIERSAKRARPIRRIPRQGRRLHPPACARCAQGMVGARWPQPGVTLTDWIVDRVRHNTMNVFKVPQSLADKYH..GAGYALAATVAGQLVDIVYLAD.....M	97
Acr00222	type	II-CMNVFPIPDSLAEAYH..GAGWALGAVLGGQLVALRYVAD.....	37
Acr00220	type	II-CMNVYPVPDSFAGNY..GVGYALAAIAGGQVLSIKYIAEH..VDEKT	42
Acr00224	type	II-C	MTDSQGREALSVIHIN.....VPRAMKARWVKASQSREQKLTDWIEKLESRTMNVYPVPETLAAKYH..GAGHALAAVTGGQIVDLVYIED.....	85
Acr00227	type	II-CMNVYPVPDALQHHYK..GRCIALGATRNCELVALVYVRD.....	37
Acr00229	type	II-CMASVTTTPDPNLRERIE..ETGWALAAISGNQIAAVREFRR...LSPQV	45
Acr00219	type	II-CMKEVFKLKEPEL.VTYK..CCGWALACIKDGETIDLTYYVRDL..G..IE	41
Acr00226	type	II-CMNFLVCPKLGINYEKIQNTPAAEYYRGCGKAWIATANGIPQAIKYMD.....	48
Acr00068	type	I-FMSSTISDRIISRSVIEAARFIQSWEDADPDNTESQVLAASSFAARLHEGLQATVLQRLVDESNRDEYREFQAW	75
consensus			*****	

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Acr00221	type	II-C	.ICPDYDGGDDKAEDWLRW.....GTDSRVKAAALEME.QYAYTSVG.....MASCWEFVEL.....	85
Acr00230	type	II-C	.ICPDYDGGDDKAEDWLRW.....GTDSRVKAAALEME.QYAYTSVG.....MASCWEFVEL.....	86
Acr00223	type	II-C	LPDFGGQDGPTRADAQTA.....IDEPVLAPTVRHLQ.ALGSVHMG.....MLSGWAFVEL.....	103
Acr00225	type	II-C	LPDFGGQDGPTRADAQTA.....IDEPVLAPTVRHLQ.ALGSVHMG.....MLSGWAFVEL.....	147
Acr00222	type	II-C	LGVSVEPCAAGVQAVRAW.....IESPGAAPVVRELQ.ALGRVTIG.....MCSGWQFVEQ.....	87
Acr00220	type	II-C	QDELAEGCAPARNAAFKW.....IGSQSAGPVVRELQ.ALGRVCAG.....MCSGWEFVEL.....	92
Acr00224	type	II-C	..ALLDYDADAPGALAAA.....IDDQRLAPTVRRLQ.ALGQVSAG.....MCGAWEFVEL.....	133
Acr00227	type	II-C	...LIPDFADTVEAAPSI.....GDNARVAQVARALEKDFCFVRVC.....IVSSWEFCEL.....	85
Acr00229	type	II-C	AQHLPIGAEEQAAEVIRRW.....IDTLPASSVVDELQ.QLGRITVG.....MCHEGQFLEC.....	95
Acr00219	type	II-C	EYDENFDGLEPEIIYYDV.....VASQACKEVAYRYE.EMGEFTFG.....LCSCWEFNVN.....	91
Acr00226	type	II-CFAYDPDLEASQR.....EQFGVYRQKCRDYLSKCRTEG.....HCSCWEFTTNAYYCEAESLLAAIESEREYME	117
Acr00068	type	I-F	EALLNADGRVTSNPFADWGWYR..IANVMLATASQNVGVAWCSHVHCRLMAIFQDRFQHHYEDEEC.....	140
consensus			* * * * *	

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved