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LESKNNLRKLLNEYEEIDINEMLKNFRSMFKRAITMDSEELRNLLYIEYPESDINEIIVLD...TVPAAAAETVVRILLVNYTEHGINYNDTVDDIVDD67
Acr00005 type I-DMIKAITMDSEELRNLLYIEYPESDINEIIVLD.....TVPAAAAETVVRILLVNYTEHGINYNDTVDDIVDD67
Acr00007 type I-DMIKAITMDSEELRNLLYIEYPESNINEIIVLD.....TVPAAAAETVVRILLVNYTEHGINYNDTVDDIVDD67
Acr00006 type I-DMIKAITMDSEQLRNLLYIEYPESNIDEIIVLD.....TVPAAAAETVVRILLVNYTEHGINYNDTVDDIVDD67
Acr00294 type II-CMFKRAIIFTS.....FNGLETVSQTEKRRLA.....KIINSRVSTIDEYLRAKDTNASLDGQYRAFLFNDE61
Acr00292 type II-CMFKRAIIFTS.....FNGFEKVSRTEKRRLA.....KIINSRVSTIDEYLRAKDTNASLDGQYRAFLFNDE61
Acr00301 type II-CMFKRAIIFTS.....FNGFEKVSRTEKRRLA.....KIINSRVSTIDEYLRAKDTNASLDGQYRAFLFNDE18
Acr00298 type II-CMFKRAIIFTS.....FNGFEKVSRTEKRRLA.....KIINSEVSIINEHLEAKATNASLDGQYRAFLFNDE61
Acr00193 type II-AMNKLKELVR.....EIKN.....KDYTAKLSGTDSNSITGLIHVN36
Acr00196 type II-AMNISELIR.....EIKN.....KDYAVRLEGTDDBNSTITLIIDVDN36
Acr00044 type I-DMEKQDLEKIESDIIN.....DWTEADDLDDALDFLFMEKVSEFKIKFDPLKVT50
Acr00001 type I-B LESKNNLRKLLNEYEEIDINEMLKNFRSINKSGTKNDIEIFLHEKAIKFEKSSISSTVVFSIEDNEILGYFTIANRSVIPKENFGILSKTQQKKLGNSAAALKNG105
Acr00263 type II-CMDEIEDLSDLPMPRFIWG.....FAVWAGKGGEVMHDEFEYLTHTRSPRTCFVVELEDV55
consensus * * * *

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Acr00005 type I-D EPYYNDIPMKILFDYMKDYGKTCFS...IILKLT E E Y S S D F L P L D T A Q S...ANDITKTLIK 123

Acr00007 type I-D EPYYNDIPMKVLFYDMKDYGKTCFS...IILKLT E E Y S S D F L P L D T A R S...ANDITKTMIK 123

Acr00006 type I-D EPYYNDIPMKVLFYDMKDYGKTCFS...IILKLT E E Y S S D F L P L D T A Q S...ANDITKTLIK 123

Acr00294 type II-C SP.....AMTEFLAKLKAFAESTAG...INIDAW E I E E S E Y N S L P L K Q K D F L A A A N G K E I F K I 116

Acr00292 type II-C SP.....AMTEFLAKLKAFAESTAG...INIDAW E I E E S E Y N S L P L K Q K D F L A A A N G K E I F K I 116

Acr00301 type II-C SP.....AMTEFLAKLKAFAESTAG...INIDAW E I E E S E Y N S L P L K Q K D F L A A A N G K E I F K I 73

Acr00298 type II-C SP.....AMTEFLAKLKAFAESTAG...INIDAW E I E E S E Y N S L P L K Q K D F L A A A N G K E I F K I 116

Acr00193 type II-A DGN...EYGISESNFESIVEKFVS...NFENGWDGAYEDEEEFYNDMQ...DIVNR.HFK 86




Acr00196 type II-A DGN...EYVISESKNESIAEKFAS...TFKNGWNKEYEDEEEFYNDMQ...SIILKSELN 87

Acr00044 type I-D EEYR...ELLGNYDSSNSVSSNGIT...IDQYTYDEDDDIMYKLEFTYRKEDNKIYIYEVQGWREKKK 112

Acr00001 type I-B DLMTSSFLLGQLGKNYSDDIENLITGRELLTFAYDLFKIKELINVKYIWLECNPKLISFYQNFGFKMLESLTSEEGLKVMIMELK 193

Acr00263 type II-C PADNGEDDIDGRIVHHDPRLLFYIT.....DAGMALVNVEFFDKLPKEAKLKKVKCDEAIANWMLRREFLDEEEED 126

consensus *** * **** * * * * * * * * * * * * * * *

-  non conserved
-  similar
-  > 50% conserved