


[illegible]



Sequence	Position 12
LKFAE ST AGIN DA WE EE SEYVRLP VE EQKD FL AAA.NGKE I FKI	116
LKFAE ST AGIN DA WE EE SEYVRLP VE EQKD FL AAAANGKE I FKI	117
LKFAE ST AGIN DA WE EE SEYVRLP VE EQKD FL AAA.NGKE I FKI	138
LKFAE ST AGIN DA WE EE SEYVRLP VE EQKD FL AAA.NGKE I FKI	116
LKFAE ST CTGIS DA WE EE SEYVRLP VE RRD FL AAA.NGKE I FKI	116
LKFAE ST CTGIS DA WE EE SEYVRLP VE RRD FL AAA.NGKE I FKI	116
LKFAE ST AGIN DA WE EE SEYVRLP VE RRD FL AAA.NGKE I FKI	116
LKFAE ST AGIN DA WE EE SEYVRLP VE RRD FL AAA.NGKE I FKI	116
LEVLE ER GAKIDR DA AA I AV AL Q FF .AAAVN..DLHSGD.... E	100
LEVLE ER GAKIDR DA AA I AV AL Q FF .AAAVN..DLHSGD.... E	100
LEFLINAGLVFD V QEA I NKARSLH.IANQS..PC.....	86
L K FL R ELGIKF DA DQALRYAVYLSG I LDYQSSKLKWL.....	99
YFDIDN TE AQ LE ALREHGAK FD FGK AL EL.....	89
V ITK V ILK FK YENGEL K LG IK FEKS I NKIS.....	116
D K I LE YNKVSVGG ET TVFKN VE VEL N TEEFMN L LF.....	129

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