```
logo
NTDB id 86295 KI216 RS01525 WP 005013206.1  MAKINLLPWRDALREKRKKEFIAICIGAFFIGVLGVGSTWLYFDNRLQDQEQANQLIVSTNQNLDIQLKSLEGLQEQRNAIVERM   85
NTDB id 1055 ABD1 RS15865 WP 000201227.1    MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDAIIERM   85
NTDB id 1011 ACIAD RS15190 WP 004923716.1   MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNAIVERM   85
consensus                                   !**!!!!!!!* !!!*!!*!* ***** **!* *!  *!*!!***!*!!!!!!!!!**!!!*!!*!!!*!*!!!*!!*!!*!!!

logo
NTDB id 86295 KI216 RS01525 WP 005013206.1  KLIQGLQGQRPIAVHLVDELVRVTPSNVYITKFSRLGDKFTLEGKAESPNTVAELLRNLEASSWYRNAFMSSFLATEDKKDQAAS  170
NTDB id 1055 ABD1 RS15865 WP 000201227.1    KLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLANEEKKDKAAS  170
NTDB id 1011 ACIAD RS15190 WP 004923716.1   KLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLAAEDNKDKTPS  170
consensus                                   !!!!!!!*!!!* !*!*!!!!!!!!!**!*!!!*!*!!!!!*!!!!!!!!!!!!!!!!!!!! !!!!!!!*!!!! !**!!***!

logo
NTDB id 86295 KI216 RS01525 WP 005013206.1  SIVPRIEDSYGTFTVTADLDAIAQPELSGVDSVAAAQPGNAVSQGTTAASTPEAAK  226
NTDB id 1055 ABD1 RS15865 WP 000201227.1    SLLPRVEDHYGSFVVTVDLGEMGVTTTDD.....SAKPSTGESVGAAK........  213
NTDB id 1011 ACIAD RS15190 WP 004923716.1   SVVPRVEAGYGSFLVSVDLGELGTSLATS.....D.QPQTPPTAGVKP........  212
consensus                                   !**!!*!* !!*!*!**!!**** *  *        **! ** * !
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X non conserved

X similar

X ≥ 50% conserved