```
logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1006862 ABLC01 RS01475 WP 002116944.1 MATINLLPWRDGLREQRKKQFIILCFGVVVLGITTVFAGWFYLNQKLNDQEQANQLIISTNQNLDQQLKTLNGLQEQRDA   80
NTDB id 1055 ABD1 RS15865 WP 000201227.1      MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA   80
consensus                                      !**!!!!!!!* !!!*!!!*!*****!**!!***!**!!!!* *!!*!!!!!!!!!*!!!*!!!!!!*!*!!!*!!*!

logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      IVERMKLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLA  160
NTDB id 1006862 ABLC01 RS01475 WP 002116944.1 IIERMKLIQGLQSQRPVVVRLVDELVRVTPAAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMSSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1      IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                      !*!!!!!!!!!!*!!!!*!!!*!!!!!!!!**!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!*!!!!

logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQPQTPPTAGVKP....   212
NTDB id 1006862 ABLC01 RS01475 WP 002116944.1 TEEKKDKAISSLLPRVEDSYGSFVVTVDVGEMGVTTTDDTAN..T.SVGTDKEAAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1      NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAK..P.STGESVGAAK   213
consensus                                       !**!!!* !!**!!!!* !!!!*!*!!*!!*!********   * * !     ***
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X non conserved

X similar

X $\geq$ 50% conserved