```
logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1006820 ABLC00 RS17050 WP 006581060.1  MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIKSTNQNLDQQLKALNGLQEQRDA   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 1006820 ABLC00 RS17050 WP 006581060.1  IIERMKLIQNLQSQRPVVVRLVDELVRVTPPAMYLTKFSRVGDKFTIEGKAESPNTVAELLRNLESSPWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1       IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                      !*!!!!!!!*!!*!!!!*!!!*!!!!!!!!**!!!!!!*!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!*!!!!!!!!!!!!

logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQPQT..PPTAGVKP  212
NTDB id 1006820 ABLC00 RS17050 WP 006581060.1  NEDKKDKTASSLLPRVEDNYGSFVVTVDLGEMGVTTTNNDADKPAVAKDTGAAK  214
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDS.AKPSTGESVGAAK  213
consensus                                      *!**!!!**!!**!!!!* !!!!*!*!!!!!*!****  *  **  * * !***
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X non conserved

X similar

X $\geq$ 50% conserved