```
logo

NTDB id 1143472 AC2117 RS01575 WP 133971489.1  MATINLLPWRDGLREQRKKQFIILCFGVVVLGIATVFSGWFYLNQKLNDQEQANQLITSTNQTLDQQLKTLNGLQEQRDA   80
NTDB id 1055 ABD1 RS15865 WP 000201227.1       MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA   80
NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTNQGLDQQLKSLDGLQQQRNA   80
consensus                                      !**!!!!!!!* !!!*!!!*!*****!**!!***!* !!!!* *!!*!!!!!!!!!*!!! !!!!!!*!*!!!*!!*!

logo

NTDB id 1143472 AC2117 RS01575 WP 133971489.1  IIERMKLIQGLQSQRPVVVRLVDELVRVTPPAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASSWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1       IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
NTDB id 1011 ACIAD RS15190 WP 004923716.1      IVERMKLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLA  160
consensus                                      !*!!!!!!!!!!*!!!!*!!!*!!!!!!!!**!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!

logo

NTDB id 1143472 AC2117 RS01575 WP 133971489.1  TEEKKDKTASSLLPRVEDSYGSFVVTVDLGEAGVTTTDDTANTSVSTDKGAVK  213
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK  213
NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQP.QTPPTAGVKP  212
consensus                                       !**!!!**!!**!!!!* !!!!*!*!!!!!*!******** **  * !* *
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X non conserved

X similar

X $\geq$ 50% conserved