```
logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1043288 ACEN3N RS19705 WP 151806341.1 MAKINLLPWRDELREQKKKQFVAICVGVALIGLATVLLAWFYFDHKLEDQEQANQLVTSTNQNLDMQLKTLDGLQERRNA   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 1043288 ACEN3N RS19705 WP 151806341.1 IIERMKLIQGLQGQRPVIVRLVDELVRVVPAQMYITKFSRVGDKFTIEGKAESPNTVAELLRGLEASEWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1      IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                      !*!!!!!!!!!!*!!!!*!!!*!!!!!!*!**!!*!!!*!*!!!!!!!!!!!!!!!!!!!!!!!!*!!!! !!!!!!!!!!!!

logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP   212
NTDB id 1043288 ACEN3N RS19705 WP 151806341.1 AEENKNESQTSVIPRVEESYGSFVVTVDLGEIASVADDSATKAEQVPAAGGAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                      *!**!*** *!**!!!!* !!!!*!*!!!!!**** **  ** * ***!***
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X non conserved

X similar

X ≥ 50% conserved