```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1   MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNAIVERM   85
NTDB id 87607 TOL5 RS19745 WP 004825390.1   MAKINLLPWRDELREQKKKQFVAICVGVALIGLATVLLAWFYFDHKLEDQEQANQLVTSTNQNLDMQLKTLDGLQERRNAIIERM   85
NTDB id 1055 ABD1 RS15865 WP 000201227.1    MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDAIIERM   85
consensus                                   !**!!!!!!!**!!!**!!*!* ****!***!***!**!!!!*!!!*!!!!!!!!**!!!*!!*!!!!*!*!!!**!*!!*!!!

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1   KLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLAAEDNKDKTPS   170
NTDB id 87607 TOL5 RS19745 WP 004825390.1   KLIQGLQGQRPVIVRLVDELVRVVPAQMYITKFSRVGDKFTIEGKAESPNTVAELLRGLEASEWYRNAFMNSFLAAEENKNQAQT   170
NTDB id 1055 ABD1 RS15865 WP 000201227.1    KLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLANEEKKDKAAS   170
consensus                                   !!!!!!!*!!!!*!!!*!!!!!!*!**!!*!!!*!*!!!!!!!!!!!!!!!!!!!!!!!!*!!!! !!!!!!!!!!!!*!**!*** *

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1   SVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP   212
NTDB id 87607 TOL5 RS19745 WP 004825390.1   SVIPRVEESYGSFVVTVDLGEIASVADDSATKAEQVPAAGGAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1    SLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                   !**!!!!* !!!!*!*!!!!!****  **  ** * ***!***
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X non conserved

X similar

X ≥ 50% conserved