```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 207777 BSR55 RS01640 WP 004825390.1  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 207777 BSR55 RS01640 WP 004825390.1  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 207777 BSR55 RS01640 WP 004825390.1  AEENKNQAQTSVIPRVEESYGSFVVTVDLGEIASVADDSATKAEQVPAAGGAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                    *!**!*** *!**!!!!* !!!!*!*!!!!!****  **  ** * ***!***
```

X non conserved

X similar

X $\geq$ 50% conserved