```
logo
NTDB id 403817 GK020 RS14140 WP 156189589.1  MAKINLLPWRDELREKRKKEFIAVCIGAALIGMLAVTLAWFYYNHKLQDQEQANQLVISTNQNLDVQLKSLEGLQEQRNA  80
NTDB id 1055 ABD1 RS15865 WP 000201227.1     MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA  80
NTDB id 1011 ACIAD RS15190 WP 004923716.1    MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA  80
consensus                                    !**!!!!!!!**!!!*!!!*!* ********!* *! **!!!* !!!*!!!!!!!!*****!!!*!!*!!!*!*!!!*!!*!

logo
NTDB id 403817 GK020 RS14140 WP 156189589.1  IVERMKLIQGLQSQRPVAVRLVDELVRVTPSSMYLTKVARIGNKFTFEGKAESPNTVAEFLRNLEASPWYRNAFMNSFLA  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 403817 GK020 RS14140 WP 156189589.1  VEENQNKAPSSVVPRVEESYGSFVVTADLDEIALPGTAPASG..TQATAGAVQ  211
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK  213
NTDB id 1011 ACIAD RS15190 WP 004923716.1    AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP  212
consensus                                     !****!**!!**!!!!* !!!!*!**!!*!**** *     * !  **!*
```

X non conserved

X similar

X $\geq$ 50% conserved