```
logo
NTDB id 391516 GBN67 RS00595 WP 234963356.1  MAKINLLPWRDALREKRKKEFIAYGVGAALVGVLAVALAWFFYSQKLEDQEQANQLVVSTNQNLDVQLKSLEGLQEQRNA   80
NTDB id 1011 ACIAD RS15190 WP 004923716.1    MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1055 ABD1 RS15865 WP 000201227.1     MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA   80
consensus                                    !**!!!!!!!* !!!*!!!*!* ********!* *! **!!** *!!*!!!!!!!!*!!!*!!*!!!*!*!!!*!!*!

logo
NTDB id 391516 GBN67 RS00595 WP 234963356.1  IVERMKLIQGLQGQRPVAVRLVDEMVRVTPSNMYITKFVRTGDKFTIEGKAESPNTVAEFLRNLEASAWYRNAFMNSFLV  160
NTDB id 1011 ACIAD RS15190 WP 004923716.1    IVERMKLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1     IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                    !*!!!!!!!!!!*!!!! !!!*!!*!!!!!!*!!*!!!*!!!!!!!!!!!!!!!!!!!!!*!!!!!!!*!!!!!!!!!!!*

logo
NTDB id 391516 GBN67 RS00595 WP 234963356.1  AEEKKDKAPSSIVPRVEETYGTFVVTADLDEIAQPVVSDATTPEAANTVNNSTAGAAQ  218
NTDB id 1011 ACIAD RS15190 WP 004923716.1    AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP.....  212
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK.....  213
consensus                                    *!**!!!**!!**!!!!* !!*!*!**!!*!** ** **   ! ** ***
```

X non conserved

X similar

X $\geq$ 50% conserved