```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 368234 E8E07 RS01600 WP 086399992.1  MATINLLPWRDGLREQRKKQFIILCFGVVVLGITTVFAGWFYLNQKLNDQEQANQLIISTNQNLDQQLKTLNGLQEQRDA   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 368234 E8E07 RS01600 WP 086399992.1  IIERMKLIQGLQSQRPVVVRLVDELVRVTPAAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMSSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1     IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                    !*!!!!!!!!!!*!!!!*!!!*!!!!!!!!**!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!*!!!!

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQP.QTPPTAGVKP   212
NTDB id 368234 E8EKKDKAISSLLPRVEDSYGSFVVTVDVGEMGVTTTDDTANTSVDMNKEVAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                     !**!!!* !!**!!!!* !!!!*!*!!*!!*!********  **  * ****
```

X non conserved

X similar

X $\geq$ 50% conserved