```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 171833 AYJ52 RS03260 WP 017387055.1  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 171833 AYJ52 RS03260 WP 017387055.1  IIERMKLIQGLQSQRPVVVRLVDELVRVTPAAMYLTKFSRSGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMSSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1     IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                    !*!!!!!!!!!!*!!!!*!!!*!!!!!!!!**!!!!!!*!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!*!!!!

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQPQTPPTAGVKP....   212
NTDB id 171833 AYJ52 RS03260 WP 017387055.1  TEEKKDKAVSSLLPRVEDSYGSFVVTVDVGEMGVTTTDDTA..NT.SVGTDKEAAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSA..KP.STGESVGAAK   213
consensus                                     !**!!!* !!**!!!!* !!!!*!*!!*!!*!********   * * !     ***
```

X non conserved

X similar

X $\geq$ 50% conserved