```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 276064 C6N17 RS02270 WP 002116944.1  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 276064 C6N17 RS02270 WP 002116944.1  IIERMKLIQGLQSQRPVVVRLVDELVRVTPAAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMSSFLA  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 276064 C6N17 RS02270 WP 002116944.1  TEEKKDKAISSLLPRVEDSYGSFVVTVDVGEMGVTTTDDTAN..T.SVGTDKEAAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAK..P.STGESVGAAK   213
consensus                                     !**!!!* !!**!!!!* !!!!*!*!!*!!*!********   * * !    ***
```

X non conserved

X similar

X $\geq$ 50% conserved