```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 386457 F2A31 RS01315 WP 150024851.1  MAKINLLPWRDELREQRKKQFIAFCVGVAALGIASVFTAWMYFDHKLNDQEQANQFIVSTNQNLDVQLKSLEGLQERRNA   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 386457 F2A31 RS01315 WP 150024851.1  IIERMKLIQGLQGQRPVIVRLVDELVRVTPPTMYITKFTRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  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 386457 F2A31 RS01315 WP 150024851.1  SEEKKDKAVSSLVPRVEENYGSFVVTVDLGEIGTTVANDSASDTAASATVGGAQ   214
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTG.ESVGAAK   213
consensus                                     !**!!!* !!**!!!!* !!!!*!*!!!!!*!**** ***   * ****!***
```

X non conserved

X similar

X $\geq$ 50% conserved