```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 254575 CS557 RS01550 WP 004909633.1  MAKINLLPWRDELREQRKKQFIAYCVGVAALGVASVLAGWGYLDYKLNDQEQANQLIVSTNQNLDAQLKSLDGLQERRNA   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 254575 CS557 RS01550 WP 004909633.1  IIERMKLIQGLQGQRPVAVRLIDELVRVTPPTMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  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 254575 CS557 RS01550 WP 004909633.1  AEEKKDKTVSSLIPRIEENYGSFVVTVDLGEIGTTVEADPLAETVASGTVGAAQ   214
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPS.TGESVGAAK   213
consensus                                    *!**!!!* !!**!!*!* !!!!*!*!!!!!*!***  *    *  ***!***
```

X non conserved

X similar

X $\geq$ 50% conserved