```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1108000 ACN19N RS01355 WP 269229435.1 MAKINLLPWRDELREQKKKQFVAFCAGVAALGVVSVFSGWMYFDHKLNDQEQANQLIVSTNQNLDTQLKSLDGLQERRNA   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 1108000 ACN19N RS01355 WP 269229435.1 IIERMKLIQGLQGQRPITVRLVDELVRVTPPTMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1       IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                      !*!!!!!!!!!!*!!!**!!!*!!!!!!!!* !!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQ.PQTPPTAGVKP   212
NTDB id 1108000 ACN19N RS01355 WP 269229435.1 ADDKKDKTASSLVPRVEENYGSFVVTVDLGEIGTTAEADPTLEAVASGSGVVK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                      ****!!!**!!**!!!!* !!!!*!*!!!!!*!**    *   * ** * !* *
```

X non conserved

X similar

X $\geq$ 50% conserved