```
logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1040150 ACDW34 RS01405 WP 130074426.1  MAKINLLPWREELREQRKKQFVAISIGIAVLGFIIVLLAWMYFNQRLDDQEQANQLVTSTNQNLDTQLKTLDGLQEQRNA   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 1040150 ACDW34 RS01405 WP 130074426.1  IIERMKLIQGLQGQRPVTVRLVDELVRVVPPAMYITKFSRTGDKFTIEGKAESPNTVADFLRGLEASEWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1       IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                      !*!!!!!!!!!!*!!!!*!!!*!!!!!!*!**!!*!!!*!!!!!!!!!!!!!!!!!!!!!**!!*!!!! !!!!!!!!!!!!

logo

NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATS.DQPQTPPTAGVKP   212
NTDB id 1040150 ACDW34 RS01405 WP 130074426.1  AEENKNKPATSVVPRAEESYGSFVVSVDIGEIAQAAKPDAEQAQA...AGGAK   210
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                      *!**!*! **!**!!*!* !!!!*!*!!*!!** *   * *****  **!***
```

X non conserved

X similar

X $\geq$ 50% conserved