```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 362234 FDQ49 RS11075 WP 004711656.1   MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIKSTNQNLDQQLKALNGLQEQRDA   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 362234 FDQ49 RS11075 WP 004711656.1   IIERMKLIQNLQSQRPVVVRLVDELVRVTPPAMYLTKFSRVGDKFTIEGKAESPNTVAELLRNLESSPWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1      IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                     !*!!!!!!!*!!*!!!!*!!!*!!!!!!!!**!!!!!!*!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!*!!!!!!!!!!!!

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQPQTPPT..AGVKP   212
NTDB id 362234 FDQ49 RS11075 WP 004711656.1   NEDKKDKTASSLLPRVEDNYGSFVVTVDLGEMGVTTTNNDAVKPAVAKDTGAAK   214
NTDB id 1055 ABD1 RS15865 WP 000201227.1      NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDS.AKPSTGESVGAAK   213
consensus                                     *!**!!!**!!**!!!!* !!!!*!*!!!!!*!****  *  **  *   !***
```

X non conserved

X similar

X $\geq$ 50% conserved