```
logo
NTDB id 1011 ACIAD RS15865 WP 000201227.1   MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNAIVERM   85
NTDB id 127653 IX88 RS00720 WP 000201224.1  MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDAIIERM   85
NTDB id 1055 ABD1 RS15865 WP 000201227.1    MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDAIIERM   85
consensus                                   !**!!!!!!!**!!!*!!!*!******!**!!***!**!!!!!*!!!*!!!!!!!!!!**!!!*!!!!!!*!*!!!*!!*!!*!!!

logo
NTDB id 1011 ACIAD RS15865 WP 000201227.1   KLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLAAEDNKDKTPS   170
NTDB id 127653 IX88 RS00720 WP 000201224.1  KLIQGLQSQRPVVVRLVDELVRVTPPAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLANEEKKDKAAS   170
NTDB id 1055 ABD1 RS15865 WP 000201227.1    KLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLANEEKKDKAAS   170
consensus                                   !!!!!!!*!!!!*!!!*!!!!!!!!**!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!*!**!!!**!

logo
NTDB id 1011 ACIAD RS15865 WP 000201227.1   SVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP   212
NTDB id 127653 IX88 RS00720 WP 000201224.1  SLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1    SLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
consensus                                   !**!!!!**!!!!*!*!!!!!*!*********!*!****!***
```

X non conserved

X similar

X ≥ 50% conserved