```
logo
NTDB id 172120 AZE33 RS18395 WP 020753500.1   MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQREA   80
NTDB id 1055 ABD1 RS15865 WP 000201227.1     MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA   80
NTDB id 1011 ACIAD RS15190 WP 004923716.1    MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
consensus                                    !**!!!!!!!**!!!*!!!*!******!**!!***!**!!!!!*!!!*!!!!!!!!!**!!!*!!!!!!*!*!!!*!!*!

logo
NTDB id 172120 AZE33 RS18395 WP 020753500.1   IIERMKLIQGLQSQRPVVVRLIDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA   160
NTDB id 1055 ABD1 RS15865 WP 000201227.1     IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA   160
NTDB id 1011 ACIAD RS15190 WP 004923716.1    IVERMKLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLA   160
consensus                                    !*!!!!!!!!!!*!!!!*!!!*!!!!!!!!!*!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!

logo
NTDB id 172120 AZE33 RS18395 WP 020753500.1   NEEKKDKAASSLLPMVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK   213
NTDB id 1011 ACIAD RS15190 WP 004923716.1    AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP   212
consensus                                    *!**!!!**!!**!*!!**!!!!*!*!!!!!*!*********!*!****!***
```

X non conserved

X similar

X $\geq$ 50% conserved