```
logo
NTDB id 155765 AMQ28 RS09720 WP 053579418.1  MARINLLPWRDELREKRKKEFIAICIGAALIGAFIVALTWFFYNHKLEDQEQANQLISSTNQNLDVQLKSLEGLEAQRNA   80
NTDB id 1011 ACIAD RS15190 WP 004923716.1    MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 1055 ABD1 RS15865 WP 000201227.1     MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA   80
consensus                                    !**!!!!!!!**!!!*!!!*!* ********!    ! **!!** !!!*!!!!!!!!!*!!!*!!*!!!*!*!!* !!*!

logo
NTDB id 155765 AMQ28 RS09720 WP 053579418.1  IIERMKLIQGLQSQRPIAVRLIDEMARVTPSNLYITKFVRVGDKFTIEGKAESPNTVAEFLRNLEASAWYRNAFMNSFLA  160
NTDB id 1011 ACIAD RS15190 WP 004923716.1    IVERMKLIQGLQAQRPVTVRLIDELVRVTPSNMYLTKFTRTGDKFTIEGKAESPNTVAELLRNLEASAWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1     IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                    !*!!!!!!!!!!*!!!* !!!*!!**!!!!!**!*!!!*!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!*!!!!!!!!!!!!

logo
NTDB id 155765 AMQ28 RS09720 WP 053579418.1  VEEKKDKAPSSVVPRVEESYGSFVVTADLDEIASPSEPEAPQ..TTVEAAQ..  209
NTDB id 1011 ACIAD RS15190 WP 004923716.1    AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP  212
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK  213
consensus                                     !**!!!**!!**!!!!* !!!!*!**!!*!*!*****  *  ** !  ***
```

X non conserved

X similar

X $\geq$ 50% conserved