```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 276623 C6I29 RS13330 WP 005018044.1  MTTINLLPWRDELREQRKKQFIIICIGVALLGLAGVVLTWMYYDRKLDDQEQANQLIISTNQNLDIQLKSLDGLQEQRDA   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 276623 C6I29 RS13330 WP 005018044.1  IIERMNLIQGLQGKRPVTVRLVDELARVIPANMYLTKFARTGDKFTLEGKAESPNTVAELLRGLEASPWYRNAFMNSFLA  160
NTDB id 1055 ABD1 RS15865 WP 000201227.1     IIERMKLIQGLQSQRPVVVRLVDELVRVTPSAMYLTKFSRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  160
consensus                                    !*!!!*!!!!!!**!!!*!!!*!!!*!!*!**!!!!!!*!!!!!!!*!!!!!!!!!!!!!!!!*!!!!*!!!!!!!!!!!!

logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1     AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSDQPQTPPTAGVKP.......   212
NTDB id 276623 C6I29 RS13330 WP 005018044.1  AEEKKEQTPTSLIPRPEENYGSFVVTVDAGEIAADSPPSADPNAPTNTAATGPAAGAAQ   219
NTDB id 1055 ABD1 RS15865 WP 000201227.1     NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSA..KPS....TGESVGAAK   213
consensus                                    *!**!*****!**!!*!* !!!!*!*!!*!!** **  *  ** !*  * **   ***
```

X non conserved

X similar

X ≥ 50% conserved