```
logo
NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTTNQGLDQQLKSLDGLQQQRNA   80
NTDB id 313071 AhaeAN4 RS01380 WP 004641325.1  MAKINLLPWRDELREQRKKQFIAFCVGVAALGIASVFTAWMYFDHKLNDQEQANQFIVSTNQNLDVQLKSLEGLQERRNA   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 313071 AhaeAN4 RS01380 WP 004641325.1  IIERMKLIQGLQGQRPVIVRLVDELVRVTPPTMYITKFTRTGDKFTIEGKAESPNTVAELLRNLEASPWYRNAFMNSFLA  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 313071 AhaeAN4 RS01380 WP 004641325.1  SEEKKDKAVSSLVPRVEENYGSFVVTVDLGEIGTTVTTDPASESAASVTVGGAQ  214
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTG.ESVGAAK  213
consensus                                       !**!!!* !!**!!!!* !!!!*!*!!!!!*!****** *  * ** *!***
```

X non conserved

X similar

X ≥ 50% conserved