```
logo
NTDB id 1002876 ABEF84 RS01695 WP 347473746.1  MAKINLLPWRDELRVKRNNEFIAYGVGALLLGVAAAGGGWFYYDQKLQDQEQANQLILSTNQNLDVQLKSLDGLQEQRDA   80
NTDB id 1055 ABD1 RS15865 WP 000201227.1       MATINLLPWREELREQRKKQFITLCFGVVVLGITTVFGGWFYFDHKLDDQEQANQLIMSTNQNLDQQLKTLNGLQEQRDA   80
NTDB id 1011 ACIAD RS15190 WP 004923716.1      MTKINLLPWRDKLREKRKKEFLVFSILVGLLGCAAVLMGWFYFGHKLNDQEQANQLILTNQGLDQQLKSLDGLQQQRNA   80
consensus                                      !**!!!!!!**!!**!***!* ******!!**** *!!!!***!!*!!!!!!!!!*!!!*!!*!!!*!*!!!*!!*!

logo
NTDB id 1002876 ABEF84 RS01695 WP 347473746.1  IVERMKLIQGLQTQRPIAVHLIDEIVRVTPSDMYVTKFVRTGDKFTIEGKAASPNTVAELLRNLEASPWYRNAFMNSFLA  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 1002876 ABEF84 RS01695 WP 347473746.1  AEEKKDKAPSSLIPRIEESYGTFTVTVDLDDQIAQPVLSEQ.QPQAAETTTQGATS  214
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESV..GAAK  213
NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTA..GVKP  212
consensus                                      *!**!!!**!!**!!*!* !!*!*!*!!!**** ** **  *!*****   !*
```

X non conserved

X similar

X $\geq$ 50% conserved