```
logo
NTDB id 1002959 ABEF88 RS01875 WP 347453499.1  MAKINLLPWRDELRVKRNNEFIAYGVGALLLGVTAAGGGWFYYDQKLQDQEQANQLISTNQNLDVQLKSLDGLQEQRDA   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 1002959 ABEF88 RS01875 WP 347453499.1  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 1002959 ABEF88 RS01875 WP 347453499.1  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