```
logo
NTDB id 1107059 ACLKM2 RS11890 WP 005094324.1  MAKINLLPWRDELRIKRNNEFVAICVGALFLGLAAAGSTWFYYDHKLQDQEQANQLIQSTNQNLDVQLKALEGLQEQRNA   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 1107059 ACLKM2 RS11890 WP 005094324.1  IVERMKLIQGLQTQRPIAVHLIDEIVRVTPANMYITKFQRTGDKFSIEGRAESPNTVAELLRNLEASSWYRNAFMNAFLV  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 1107059 ACLKM2 RS11890 WP 005094324.1  AEESKEKAPSSIIPRVEESYGSFTVTVDLDDQIAEPVLNEEQQAAN.TTPGASS  212
NTDB id 1055 ABD1 RS15865 WP 000201227.1       NEEKKDKAASSLLPRVEDHYGSFVVTVDLGEMGVTTTDDSAKPSTGESVGAAK  213
NTDB id 1011 ACIAD RS15190 WP 004923716.1      AEDNKDKTPSSVVPRVEAGYGSFLVSVDLGELGTSLATSD.QPQTPPTAGVKP  212
consensus                                      *!* !*!**!!**!!!!* !!!!*!*!!!**** **  ** ** *  * !*
```

X non conserved

X similar

X $\geq$ 50% conserved