```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1       MTFAKFFRNLTACSVALWLVACANQT.TQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAY   79
NTDB id 1071073 ACIRXM RS05290 WP 046097864.1  ...MKIYQCILSILFSLFLSACVSSPLNKQDFDKQQAAKARVELGIGYLEQNELGLAKQNLDKALAYAPNYYLVHSALAY   77
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQSA..SSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAH   76
consensus                                          ************ ****!!****    ***** !*!!!**!***!!*!**  *!! !*!!!! ** **!!*!!*!!*

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1       YYQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCA   159
NTDB id 1071073 ACIRXM RS05290 WP 046097864.1  FYQLQGENTRAENAYLKALKLDDK.......QGDVFNNYAVFLCKQGQFEQAYLRFEQALAT.PNYYQQADTYENIVLCA   149
NTDB id 1358 HI 0366 AAC22024.1                YYQQQGQIENAFREYEIAVNLNHK.......QGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCA   148
consensus                                      *!!**!** *!* *!  !**!  *       **!! !!***!!!*!* !**!***!! !! * **!!*!!!!*!!!*!!!

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1       NMKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1071073 ACIRXM RS05290 WP 046097864.1  LAEKNAIRYEKYKGLLEKLDPKRNQKTNVIH...   180
NTDB id 1358 HI 0366 AAC22024.1                YSAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                         * ****   *  !***! *!*** * *
```

X non conserved

X similar

X ≥ 50% conserved