```
logo
NTDB id 1358 HI 0366 AAC22024.1          ..MKTISKQLSAVIFPFIFSACVSQS....ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQ   79
NTDB id 115043 X808 RS02300 WP 025216722.1 .MFAKYLKNITACITLFLLAACSSTPETVSETEFNRSEAVKARINLALAYLEQNDFPKAKQNIDKALAHDSNDYLPHSVLAYYYQ   84
NTDB id 1386 A4U84 RS02735 WP 005710402.1 MTFAKFFRNLTACSVALWLVACANQT...TQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQ   82
consensus                                  ****  ****!**  ***  !!  ***   * ***!***!*!!!**!!!*!!*!*****!!  !*!!!!  !!  **!!*!!*!!*!!!

logo
NTDB id 1358 HI 0366 AAC22024.1          QQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVLCAYSAQKMDI  156
NTDB id 115043 X808 RS02300 WP 025216722.1 QTADAQNAEKAYQQALKLS.......QNRPDVRNNYGTFLCKQGQFQQAYQQFEQAVASPKPYYHQADSLENIILCAKKEPNPQK  162
NTDB id 1386 A4U84 RS02735 WP 005710402.1 QTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEK  167
consensus                                  !***  **!***!**!**!*       * **!! !!*!!!!!*!* !**!*!!!! !* !*  !!*!!!**!!!*!!!     ***

logo
NTDB id 1358 HI 0366 AAC22024.1          YQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 115043 X808 RS02300 WP 025216722.1 VAETLIELEKLDKPRATLLKGN....   184
NTDB id 1386 A4U84 RS02735 WP 005710402.1 QKTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                     *!  !***!* !!***
```

X non conserved

X similar

X ≥ 50% conserved