```
logo
NTDB id 1358 HI 0366 AAC22024.1          ..MKTISKQLSAVIFPFIFSACVSQS....ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQ   79
NTDB id 21065 HD RS01300 WP 010944348.1  ..MAKILKNLTACFAIVMLLGCSSSYTGSSVSTFNRNKIVQARINLALAYLEQHHFAKAKENIDKALQHDAQDYLPYSALAYYYQ   83
NTDB id 1386 A4U84 RS02735 WP 005710402.1 MTFAKFFRNLTACSVALWLVACANQTT...QVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQ   82
consensus                                 **** **!*!*   *****! ****    ***!** ***!!**!!!*!!*!  * *!!*!*!!!!*!! **!!**!*!!*!!!

logo
NTDB id 1358 HI 0366 AAC22024.1          QQGQIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDI  156
NTDB id 21065 HD RS01300 WP 010944348.1  QIGDLTQAANAYQTALTLGN.......NTPDVLNNYGTFLCHQGKFDLAYQQFEQAIKSQQSYYHQADSLENIILCAKQEKNTAK  161
NTDB id 1386 A4U84 RS02735 WP 005710402.1 QTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEK  167
consensus                                ! !** *!  *!* !* !          *!!*!!*!!!!! !**!* !*!!!! !* !*  !!*!!!**!!!*!!!   * ***

logo
NTDB id 1358 HI 0366 AAC22024.1          YQQTLEKLRQIDGKRAEKFNSLK...  179
NTDB id 21065 HD RS01300 WP 010944348.1  IAIALQQLTALDKTRADALR......  181
NTDB id 1386 A4U84 RS02735 WP 005710402.1 QKTALSQLEKIDKSRAERLYNFMEIR 193
consensus                                 *! *!  *!**!!***
```

X non conserved

X similar

X ≥ 50% conserved