```
logo
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQS.ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY    77
NTDB id 351770 EXH44 RS07670 WP 162857554.1    MTFAKFFRNLTACSVALWLVACTNQTAKVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDAKDYLPHSVLAYY    80
NTDB id 1386 A4U84 RS02735 WP 005710402.1      MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY    80
consensus                                      *********!*!********!! *!*  ***!***!*!!!**!!!*!!*!*****!!*!*!!!!*!! **!!*!!*!!*!

logo
NTDB id 1358 HI 0366 AAC22024.1                YQQQGQIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY   149
NTDB id 351770 EXH44 RS07670 WP 162857554.1    YQQTGDSPKAEEAYQQALKLSKTQSKSNQFRPDVLNNYGTFLCKQKQFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 1386 A4U84 RS02735 WP 005710402.1      YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
consensus                                      !!!*!*  *!***!**!**!****** ** **!!*!!*!!!!!*!! !**!*!!!!*!!*!***!!*!!!!*!!!*!!!*

logo
NTDB id 1358 HI 0366 AAC22024.1                SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 351770 EXH44 RS07670 WP 162857554.1    MEKNVAKQNEALEALAKLDQTRADKLRPLFK..   191
NTDB id 1386 A4U84 RS02735 WP 005710402.1      MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                      * * ***** *!* ! **! *!!***  *
```

X non conserved

X similar

X ≥ 50% conserved