```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1      MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 1051002 O1Q82 RS08355 WP 386687064.1  ......MKFIPHFILILFLTACVSQDVEIDFDQKPAAKARVELALGYLSQQDWEQAKQNLDKALSYAPDYYLVHSAFAYF   74
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQSA.SSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                             *****  **  ****!!**!*  ****    !*!!!**!!!*!! !***  !! !*!!!! **  *!!*!!**!**

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1      YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 1051002 O1Q82 RS08355 WP 386687064.1  YQQQGNITQARESYLKAIKLDDK.......QGDVHNNYGTFLCSQGEFQAAYQQFDLAVASP.NYYRQADSYENKALCAL   146
NTDB id 1358 HI 0366 AAC22024.1                YQQQGQIENAFREYEIAVNLNHK.......QGDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVLCAY   149
consensus                                      !!!*!** *! * !  !**!  *        **!!*!!*!!!!!*!* !* !*!!!**!* !* *!! !!!**!!**!!!

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1      MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1051002 O1Q82 RS08355 WP 386687064.1  SENNQAVYQQSKIALEKADPERAKKLR......   173
NTDB id 1358 HI 0366 AAC22024.1                SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                      * * ********  !***!  !!***
```

X non conserved

X similar

X $\geq$ 50% conserved