```
logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1      MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 1058285 O1Q83 RS06855 WP 132302241.1   ......MKFITHFILILFLTACVSQNAEIDFDQKPAAKARVELALGYLSQQDWEPAKQNLDKALSYAPDYYLVHSAFAYF   74
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQSAS.SLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                             ***** ** ****!!**!** ****   !*!!!**!!!*!! !***  !! !*!!!! **  *!!*!!**!**

logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1      YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 1058285 O1Q83 RS06855 WP 132302241.1   YQQQGNITQAREAYLKAIKLDDK.......QGDVHNNYGTFLCSQGEFQAAYQQFDLAVVSP.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 1058285 O1Q83 RS06855 WP 132302241.1   SENNQAVYQQSKIALEKADPERAKKLR......   173
NTDB id 1358 HI 0366 AAC22024.1                SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                      * * ********  !***!  !!***
```

X non conserved

X similar

X $\geq$ 50% conserved