```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1       MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 1065197 ACHWS2 RS08770 WP 014391406.1  ..MNRTFRFGIASIFSLLLCACVSQESNKTFDSHEAAKARIELGLGYLSQHDFARAKQNFDKALSHAPNYYLSHAVLAYL   78
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQSA.SSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                        * * *****! ** *** !!**!*** ***  *!*!!!**!*!*!! ! ** *!! !*!!!! !* **!! !**!!**

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1       YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN  160
NTDB id 1065197 ACHWS2 RS08770 WP 014391406.1  YQKQGHLDLARHAYEKAIKLDKN.......QGDVLNNYGTFLCQQGEFEQAYKQFSQALNA.PHYYQHADTYENLVWCAS  150
NTDB id 1358 HI 0366 AAC22024.1                YQQQGQIENAFREYEIAVNLNHK.......QGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY  149
consensus                                      !!**! ** !  *!* !**! *        **!!*!!*!!!!! !* !**!**!!* !!** * !!**!!!*!!***!!

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1       MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR  193
NTDB id 1065197 ACHWS2 RS08770 WP 014391406.1  FAKDAVRYQQNLQHLTQFDPSRAKKFPPYIE..  181
NTDB id 1358 HI 0366 AAC22024.1                SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...  179
consensus                                       ******** !  ! **! *!!***  ***
```

X non conserved

X similar

X $\geq$ 50% conserved