```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1     MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 324187 EEX22 RS06200 WP 156732369.1  ..MTRVFQVWIASIFSLFLSACVSQEGNKTFDSHEAAKARIELGLGYLSQHDFARAKQNFDKALSHAPNYYLSHAVLAYL   78
NTDB id 1358 HI 0366 AAC22024.1              ..MKTISKQLSAVIFPFIFSACVSQSAS.SLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                      * *******! ** ****!!**!*** ***  *!*!!!**!*!*!! ! ** *!! !*!!!! !* **!! !**!!**

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1     YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN  160
NTDB id 324187 EEX22 RS06200 WP 156732369.1  YQKQGHLDLARHAYEKAIKLDKN.......QGDVLNNYGTFLCQQGEFEQAYKQFSQALKA.PYYYQHADTYENLVWCAS  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 324187 EEX22 RS06200 WP 156732369.1  FAKDAVRYQQNLQHLTQFDPNRAKKFPPHIE..  181
NTDB id 1358 HI 0366 AAC22024.1              SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...  179
consensus                                     ********* !  ! **!  !!***   **
```

X non conserved

X similar

X $\geq$ 50% conserved