```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1  MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
NTDB id 20807 PM RS10420 WP 010907447.1    ..MTRVFQVWIASIFSLFLSACVSQEGNKTFDSHEAAKARIELGLGYLSQHDFARAKQNFDKALSHAPNYYLSHAVLAYLYQKQG   83
NTDB id 1358 HI 0366 AAC22024.1            ..MKTISKQLSAVIFPFIFSACVSQSAS.SLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
consensus                                   * *******! ** ****!!**!*** ***  *!*!!!**!*!*!! ! ** *!! !*!!!! !* **!! !**!!**!!**!
```

```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1  DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT  170
NTDB id 20807 PM RS10420 WP 010907447.1    HLDLARHAYEKAIKLDKN.......QGDVLNNYGAFLCQQGEFEQAYKQFSQALNA.PHYYQHADTYENLVWCASFAKDTVRYQQ  160
NTDB id 1358 HI 0366 AAC22024.1            QIENAFREYEIAVNLNHK.......QGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDIYQQ  159
consensus                                   ** !  *!* !**! *        **!!*!!*!*!!! !* !**!**!!* !!** * !!**!!!*!!***!!  *********
```

```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1  ALSQLEKIDKSRAERLYNFMEIR  193
NTDB id 20807 PM RS10420 WP 010907447.1    NLQHLTQFDPNRAKKFPPHIE..  181
NTDB id 1358 HI 0366 AAC22024.1            TLEKLRQIDGKRAEKFNSLK...  179
consensus                                   !  ! **!  !!***   **
```

X non conserved

X similar

X ≥ 50% conserved