```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1   MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
NTDB id 123478 H733 RS01345 WP 038439288.1  ..MQTISKQLSAVIFPFIFSACVSQS.ASSLDHQAAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
NTDB id 1358 HI 0366 AAC22024.1             ..MKTISKQLSAVIFPFIFSACVSQS.ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
consensus                                     * *****!*!********!!**!* ******* !*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!!!!*!
```

```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1   DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT   170
NTDB id 123478 H733 RS01345 WP 038439288.1  QIKNAFREYEIAVKLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQTDTFENIALCAYSAKKMNIYQQ   159
NTDB id 1358 HI 0366 AAC22024.1             QIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDIYQQ   159
consensus                                   ** *!***!**!**!**       ***!!*!!*!!!!!*!!*!**!*!!!!*!!*! **!!*!*!!*!!!*!!!******!****
```

```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1   ALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 123478 H733 RS01345 WP 038439288.1  TLEKLRQIDGKRAEKFNSLK...   179
NTDB id 1358 HI 0366 AAC22024.1             TLEKLRQIDGKRAEKFNSLK...   179
consensus                                   *!**!**!!**!!!******
```

X non conserved

X similar

X $\geq$ 50% conserved