```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1   MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
NTDB id 39457 HICON RS04495 WP 006996133.1  ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQAAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
NTDB id 1358 HI 0366 AAC22024.1             ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
consensus                                     *******!*!********!!**! *****!** !*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!!!!*!
```

```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1   DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT   170
NTDB id 39457 HICON RS04495 WP 006996133.1  EIENAHREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIALCAYSAQKMDIYQQ   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 39457 HICON RS04495 WP 006996133.1  TLEKLRQIDGKRAEKFNSLK...   179
NTDB id 1358 HI 0366 AAC22024.1             TLEKLRQIDGKRAEKFNSLK...   179
consensus                                   *!**!**!!**!!!******
```

X non conserved

X similar

X $\geq$ 50% conserved