```
logo
NTDB id 127125 JL26 RS06680 WP 021110392.1  MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
NTDB id 1386 A4U84 RS02735 WP 005710402.1   MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
NTDB id 1358 HI 0366 AAC22024.1             ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
consensus                                   *********!*!********!!**!******!***!*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!!!!*!
```

```
logo
NTDB id 127125 JL26 RS06680 WP 021110392.1  DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKQFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT   170
NTDB id 1386 A4U84 RS02735 WP 005710402.1   DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT   170
NTDB id 1358 HI 0366 AAC22024.1             QIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDIYQQ   159
consensus                                   ****!***!**!**!************!!*!!*!!!!!*!! !**!*!!!!*!!*!***!!*!!!!*!!!*!!!***********
```

```
logo
NTDB id 127125 JL26 RS06680 WP 021110392.1  ALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1386 A4U84 RS02735 WP 005710402.1   ALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1358 HI 0366 AAC22024.1             TLEKLRQIDGKRAEKFNSLK...   179
consensus                                   *!**!**!!**!!!*********
```

X non conserved

X similar

X $\geq$ 50% conserved