```
logo
```

```
NTDB id 1386 A4U84 RS02735 WP 005710402.1    MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
NTDB id 44222 K6J73 RS03310 WP 221253412.1   ..MKTISKQLSAVIFPFIFSACVSQ.SASNLNHQAAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
NTDB id 1358 HI 0366 AAC22024.1              ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
consensus                                      *******!*!********!!**!  ***  *!**  !*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!!!!*!
```

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logo
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```
NTDB id 1386 A4U84 RS02735 WP 005710402.1    DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT   170
NTDB id 44222 K6J73 RS03310 WP 221253412.1   EIENAHREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSSQKIDIYQQ   159
NTDB id 1358 HI 0366 AAC22024.1              QIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDIYQQ   159
consensus                                    *****!  **!**!**!**       ***!!*!!*!!!!!*!!*!**!*!!!!*!!*!  **!!*!!!!*!!!*!!!** ********
```

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logo
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```
NTDB id 1386 A4U84 RS02735 WP 005710402.1    ALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 44222 K6J73 RS03310 WP 221253412.1   TLEKLRQIDGKRAEKFNSLK...   179
NTDB id 1358 HI 0366 AAC22024.1              TLEKLRQIDGKRAEKFNSLK...   179
consensus                                    *!**!**!!**!!!******
```

X non conserved

X similar

X ≥ 50% conserved