logo

```
NTDB id 1386 A4U84 RS02735 WP 005710402.1     MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 304444 DV369 RS08675 WP 005649140.1   ..MQTISKQLSAVIFPFIFSACVSQS.ASSLDHQAAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
NTDB id 1358 HI 0366 AAC22024.1               ..MKTISKQLSAVIFPFIFSACVSQS.ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                       * *****!*!********!!**!* ******* !*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!
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logo

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NTDB id 1386 A4U84 RS02735 WP 005710402.1     YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN  160
NTDB id 304444 DV369 RS08675 WP 005649140.1   YQQQGQIENAFREYEIAVKLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIALCAY  149
NTDB id 1358 HI 0366 AAC22024.1               YQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY  149
consensus                                     !!!*!****!***!**!**!*       ****!!*!!*!!!!!*!!*!**!*!!!!*!!*!  **!!*!!!!*!!!*!!!*
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logo

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NTDB id 1386 A4U84 RS02735 WP 005710402.1     MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 304444 DV369 RS08675 WP 005649140.1   SAKKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 1358 HI 0366 AAC22024.1               SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                     ***********!**!**!!**!!!******
```

X non conserved

X similar

X ≥ 50% conserved