```
logo

NTDB id 232172 B4U42 RS09510 WP 005710402.1   MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 1386 A4U84 RS02735 WP 005710402.1     MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 1358 HI 0366 AAC22024.1               ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                     *********!*!********!!**!******!***!*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!

logo

NTDB id 232172 B4U42 RS09510 WP 005710402.1   YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 1386 A4U84 RS02735 WP 005710402.1     YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 1358 HI 0366 AAC22024.1               YQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY   149
consensus                                     !!!*!****!***!**!**!************!!*!!*!!!!!*!!*!**!*!!!!*!!*!***!!*!!!!*!!!*!!!*

logo

NTDB id 232172 B4U42 RS09510 WP 005710402.1   MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1386 A4U84 RS02735 WP 005710402.1     MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1358 HI 0366 AAC22024.1               SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                     ***********!**!**!!**!!!*********
```

X non conserved

X similar

X $\geq$ 50% conserved