```
logo
NTDB id 1142626 DQL18 RS03700 WP 042611895.1  ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQAAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSTLAHY   77
NTDB id 1358 HI 0366 AAC22024.1               ..MKTISKQLSAVIFPFIFSACVSQ.SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
NTDB id 1386 A4U84 RS02735 WP 005710402.1     MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
consensus                                       *******!*!********!!**! *****!** !*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!! !!*!

logo
NTDB id 1142626 DQL18 RS03700 WP 042611895.1  YQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY   149
NTDB id 1358 HI 0366 AAC22024.1               YQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY   149
NTDB id 1386 A4U84 RS02735 WP 005710402.1     YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
consensus                                     !!!*!****!***!**!**!*       ****!!*!!*!!!!!*!!*!**!*!!!!*!!*! **!!*!!!!*!!!*!!!*

logo
NTDB id 1142626 DQL18 RS03700 WP 042611895.1  SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 1358 HI 0366 AAC22024.1               SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 1386 A4U84 RS02735 WP 005710402.1     MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                     ***********!**!**!!**!!!******
```

X non conserved

X similar

X $\geq$ 50% conserved