```
logo
NTDB id 1358 HI 0366 AAC22024.1          ..MKTISKQLSAVIFPFIFSACVSQS.ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQQG   82
NTDB id 56700 WQG RS09125 WP 015433244.1 MFFAKLPQYLSACLLIFSLSGCVNSQTSADFDRDEALIARASLALAYLEQQDFSKAKENIEKALSHNSQHYLAHLVAAYYYQQLS   85
NTDB id 1386 A4U84 RS02735 WP 005710402.1 MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQTG   85
consensus                                * **** **!*!*** * ***!*****  **** *!**!!* !!!*!!*!*** *!!*!**!!! !* * !! !***!*!!!! *

logo
NTDB id 1358 HI 0366 AAC22024.1          QIENAFREYEIAVNLNHKQ.......GDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVLCAYSAQKMDIYQQ  159
NTDB id 56700 WQG RS09125 WP 015433244.1 DVQHAEQSYQAAIRLSHQI.KQNSPLPEVLNNYGTFLCQQGKFVQAYTQFEQALTGEKAYLRQGDTLENMVLCAKQANQLEKMAQ  169
NTDB id 1386 A4U84 RS02735 WP 005710402.1 DNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQKT  170
consensus                                *** !*  !* !**!** * ***   **!*!!*!!!!! !**!**!**!!! !!**  *!* !*!!*!!**!!!  ** ***  *

logo
NTDB id 1358 HI 0366 AAC22024.1          TLEKLRQIDGKRAEKFNSLK...  179
NTDB id 56700 WQG RS09125 WP 015433244.1 ALARLALLDAERATKLAKIVK..  190
NTDB id 1386 A4U84 RS02735 WP 005710402.1 ALSQLEKIDKSRAERLYNFMEIR  193
consensus                                *! *!  *!* !!***  **
```

X non conserved

X similar

X ≥ 50% conserved