```
logo

NTDB id 1358 HI 0366 AAC22024.1             ..MKTISKQLSAVIFPFIFSACVSQ...SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQ   80
NTDB id 122837 ACEE RS00335 WP 039195970.1  MNLAKFLRNLTACMLLSVLFGCSSQIPTSQTTFNRSEAVKARINLALAYLEQHDLPKAKENIDKALSHDQNDYLPHSVLAYYYQQ   85
NTDB id 1386 A4U84 RS02735 WP 005710402.1   MTFAKFFRNLTACSVALWLVACANQ..TTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQ   83
consensus                                   * **** **!*!***  ** *! *!  ******!***!*!!!**!!!*!!*! * **!!*!*!!!! !! **!!*!!*!!*!!!!

logo

NTDB id 1358 HI 0366 AAC22024.1             QGQIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDIY  157
NTDB id 122837 ACEE RS00335 WP 039195970.1  IGDQAHAERAYQQALKLSD.......NRPDVLNNYGTFLCKQAKFDQAYQQFEQAVQSEQSYYHQADSLENIVLCAKQEPNLAKV  163
NTDB id 1386 A4U84 RS02735 WP 005710402.1   TGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQ  168
consensus                                    !**  !***!**!**!*         **!!*!!*!!!!!*!**!**!*!!!! !**!   !!*!!!**!!!*!!!     ***

logo

NTDB id 1358 HI 0366 AAC22024.1             QQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 122837 ACEE RS00335 WP 039195970.1  NEALNQLEKLDKARAAALR......   182
NTDB id 1386 A4U84 RS02735 WP 005710402.1   KTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                   * *! *!***!* !!***
```

X non conserved

X similar

X ≥ 50% conserved