```
logo
NTDB id 1358 HI 0366 AAC22024.1          ..MKTISKQLSAVIFPFIFSACVSQ...SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQ   80
NTDB id 29837 APJL RS06335 WP 012263163.1 MNLAKFLRNLTACMLLSVLFGCSSQLPTSQTTFNRSEAVKARINLALAYLDQHDLPKAKENIDKALSHDQNDYLPHSVLAYYYQQ   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 29837 APJL RS06335 WP 012263163.1 IGDQVHAEQAYQQALKLSN.......NRPDVLNNYGTFLCKQAKFDQAYQQFEQAVQSEQPYYHQADSLENIVLCAKQEPNLAKV  163
NTDB id 1386 A4U84 RS02735 WP 005710402.1 TGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQ  168
consensus                                 !**  !* *!**!**!*         **!!*!!*!!!!!*!**!**!*!!!! !**!   !!*!!!**!!!*!!!     ***

logo
NTDB id 1358 HI 0366 AAC22024.1          QQTLEKLRQIDGKRAEKFNSLK...  179
NTDB id 29837 APJL RS06335 WP 012263163.1 NEALSQLEKLDKARAAALR......  182
NTDB id 1386 A4U84 RS02735 WP 005710402.1 KTALSQLEKIDKSRAERLYNFMEIR  193
consensus                                * *!**!***!* !!***
```

X non conserved

X similar

X ≥ 50% conserved