```
logo
NTDB id 1358 HI 0366 AAC22024.1              ..MKTISKQLSAVIFPFIFSACVSQ...SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALA   75
NTDB id 301286 DRF63 RS06905 WP 005598298.1  MNLAKFLRNLTACMLLSVLFGCSSQLPTSQITFNRSEAVKARINLALAYLDQHDLPKAKENIDKALSHDQNDYLPHSVLA   80
NTDB id 1386 A4U84 RS02735 WP 005710402.1    MTFAKFFRNLTACSVALWLVACANQ..TTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLA   78
consensus                                    * ***** **!*!***  ** *! *!  ******!***!*!!!**!!!*!!*! * **!!*!*!!!! !! **!!*!!*!!

logo
NTDB id 1358 HI 0366 AAC22024.1              HYYQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLC  147
NTDB id 301286 DRF63 RS06905 WP 005598298.1  YYYQQIGDQVHAEQAYQQALKSS.......DNRPDVLNNYGTFLCKQAKFDQAYQQFEQAVKSEQPYYHQADSLENIVLC  153
NTDB id 1386 A4U84 RS02735 WP 005710402.1    YYYQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALC  158
consensus                                    *!!!! !**  !* *!**!****         **!!*!!*!!!!!*!**!**!*!!!! !* !   !!*!!!**!!!*!!

logo
NTDB id 1358 HI 0366 AAC22024.1              AYSAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...  179
NTDB id 301286 DRF63 RS06905 WP 005598298.1  AKQEPNLAKVNEALSQLEKLDKARAAALR......  182
NTDB id 1386 A4U84 RS02735 WP 005710402.1    ANMKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR  193
consensus                                    !     *** * *!**!***!* !!***
```

X non conserved

X similar

X ≥ 50% conserved