```
logo
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQ...SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALA   75
NTDB id 312054 D1106 RS06660 WP 237593742.1    MNLAKFLRNLSACMLLSVLFGCSSQLPTSQTTFNRSEAVKARINLALAYLDQHDLPKAKENIDKALSHDQNDYLPHSVLA   80
NTDB id 1386 A4U84 RS02735 WP 005710402.1      MTFAKFFRNLTACSVALWLVACANQ..TTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLA   78
consensus                                      * ***** **!*!***  ** *! *!  ******!***!*!!!**!!!*!!*! * **!!*!*!!!! !! **!!*!!*!!

logo
NTDB id 1358 HI 0366 AAC22024.1                HYYQQQGQIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLC  147
NTDB id 312054 D1106 RS06660 WP 237593742.1    YYYQQIGDQVHAEQAYQQALKLSN.......NRPDVLNNYGTFLCKQAKFDQAYQQFEQAVQSEQPYYHQADSLENIVLC  153
NTDB id 1386 A4U84 RS02735 WP 005710402.1      YYYQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALC  158
consensus                                      *!!!! !**  !* *!**!**!*         **!!*!!*!!!!!*!**!**!*!!!! !**!   !!*!!!**!!!*!!

logo
NTDB id 1358 HI 0366 AAC22024.1                AYSAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 312054 D1106 RS06660 WP 237593742.1    AKQEPNLAKANEALSQLEKLDKSRAAALR......   182
NTDB id 1386 A4U84 RS02735 WP 005710402.1      ANMKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                      !      *** * *!**!***!**!!***
```

X non conserved

X similar

X $\geq$ 50% conserved