```
logo

NTDB id 1358 HI 0366 AAC22024.1          ..MKTISKQLSAVIFPFIFSACVS...QSASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALA   75
NTDB id 1147532 CKV75 RS00480 WP 014990914.1 MNLAKFLRNLTACMLLPVLFGCSSLLPPPQTTFNRSEAVKARINLALAYLDQHDLPKAKENIDKALSHDQNDYLPHSVLA   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 1147532 CKV75 RS00480 WP 014990914.1 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 1147532 CKV75 RS00480 WP 014990914.1 AKQEPNLAKVNEALSQLEKLDKARAAALR......   182
NTDB id 1386 A4U84 RS02735 WP 005710402.1 ANMKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                !     *** * *!**!***!* !!***
```

X non conserved

X similar

X $\geq$ 50% conserved