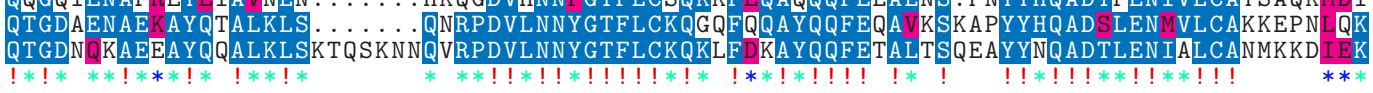
```
logo

NTDB id 1358 HI 0366 AAC22024.1             ..MKTISKQLSAVIFPFIFSACVSQSA....SSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQ   79
NTDB id 58288 MHH RS19190 WP 006251773.1    .MFAKFFPKMTACILIFLLAACSSLPPSSSEVEFNRSEAVKARINLALAYLEQNDFPKAKQNIDKALAHDEKDYLPYSVLAYYYQ   84
NTDB id 1386 A4U84 RS02735 WP 005710402.1   MTFAKFFRNLTACSVALWLVACANQ...TTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQ   82
consensus                                    **********!*** *** !! **    ** ***!***!*!!!**!!!*!!*!*****!! !*!!!! !! **!!**!*!!*!!!

logo                                        

NTDB id 1358 HI 0366 AAC22024.1             QQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDI  156
NTDB id 58288 MHH RS19190 WP 006251773.1    QTGDAENAEKAYQTALKLS.......QNRPDVLNNYGTFLCKQGQFQQAYQQFEQAVKSKAPYYHQADSLENMVLCAKKEPNLQK  162
NTDB id 1386 A4U84 RS02735 WP 005710402.1   QTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEK  167
consensus                                   !*!* **!***!* !**!*       * **!!*!!*!!!!!*!* !**!*!!!! !* !   !!*!!!**!!**!!!     ***

logo

NTDB id 1358 HI 0366 AAC22024.1             YQQTLEKLRQIDGKRAEKFNSLK...   179
NTDB id 58288 MHH RS19190 WP 006251773.1    IAETLTELEKLDKPRANLLKEN....   184
NTDB id 1386 A4U84 RS02735 WP 005710402.1   QKTALSQLEKIDKSRAERLYNFMEIR   193
consensus                                      *!* !***!* !!***
```

X non conserved

X similar

X ≥ 50% conserved