```
logo
NTDB id 1358 HI 0366 AAC22024.1              ..MKTISKQLSAVIFPFIFSACVSQS....ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSAL   74
NTDB id 298822 CT138 RS04625 WP 112110586.1  .MFAKYLKNITACITLFLLAACSSYTETASETEFNRSEAVKARINLALAYLEQNDFPKAKQNIDKALTHDPNDYLPHSVL   79
NTDB id 1386 A4U84 RS02735 WP 005710402.1    MTFAKFFRNLTACSVALWLVACANQ...TTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVL   77
consensus                                      **** ****!**  *** !! ***   * ***!***!*!!!**!!!*!!*!*****!! !*!!!! !! **!!*!!*!

logo
NTDB id 1358 HI 0366 AAC22024.1              AHYYQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVL  146
NTDB id 298822 CT138 RS04625 WP 112110586.1  AYYYQQTADAQNAEKAYQQALKLS.......QNRPDVRNNYGTFLCKQGQFQQAYQQFEQAVASPKPYYHQADSLENIIL  152
NTDB id 1386 A4U84 RS02735 WP 005710402.1    AYYYQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIAL  157
consensus                                    !*!!!!*** **!***!**!**!*       * **!! !!*!!!!!*!* !**!*!!!! !* !*  !!*!!!**!!!*!

logo
NTDB id 1358 HI 0366 AAC22024.1              CAYSAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...  179
NTDB id 298822 CT138 RS04625 WP 112110586.1  CAKKEPNPQKVAETFIELEKLDKPRATLLKGN....  184
NTDB id 1386 A4U84 RS02735 WP 005710402.1    CANMKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR  193
consensus                                    !!      ***   **  !***!* !!***
```

X non conserved

X similar

X $\geq$ 50% conserved