```
logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1      .MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAY   79
NTDB id 1142579 DQL22 RS00230 WP 109842957.1  MVNIQFIFYRFVPFIFPFLFSACVVHSDAPDFDKQAAAKARVELALGYLQQQDGSQAKLNLDKALSYAPKYYLVHAALAY   80
NTDB id 1358 HI 0366 AAC22024.1                ...MKTISKQLSAVIFPFIFSACVSQS.ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAH   76
consensus                                       * *  ******* *******!!* ** * ***** !*!!!**!!!*!!*!**   !!*!*!!!! ** **!!*!**!!*

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1      YYQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCA   159
NTDB id 1142579 DQL22 RS00230 WP 109842957.1  FYQQQGDVEHAKQSYLTAIKLDDK.......QGDVFNNFGAFLCSQGEYQAAYEQFKQALKSP.NYYHQADTYENLALCA   152
NTDB id 1358 HI 0366 AAC22024.1                YYQQQGQIENAFREYEIAVNLNHK.......QGDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVLCA   148
consensus                                      *!!!*!*** !   !  !**!  *       **!! !!*!*!!!*!* ** !**!!* !! !* *!!*!!!!*!!**!!!

logo
NTDB id 1386 A4U84 RS02735 WP 005710402.1      NMKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1142579 DQL22 RS00230 WP 109842957.1  LSAKEQKIYQENLAALEKIEPARAKKLTQFVK..   184
NTDB id 1358 HI 0366 AAC22024.1                YSAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                       *********  !  !**!*  !!*** ***
```

X non conserved

X similar

X $\geq$ 50% conserved