```
logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1     MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY    80
NTDB id 210474 BTV22 RS06550 WP 075294142.1   ...MRYRTIFTFIIFSLFLSACVSQPITVRFNKQQAAKARIELGLGYLKLRNFSQAKLNFDKALFYQPDYYLVHSAFAYL    77
NTDB id 1358 HI 0366 AAC22024.1               ..MKTISKQLSAVIFPFIFSACVSQSA.SSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY    77
consensus                                         ** ******* ****!!**!*  * *!** !*!!!**!*!*!! * **  !!*!*!!!! **  *!!*!!**!**

logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1     YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 210474 BTV22 RS06550 WP 075294142.1   YQLQGDISQAKSSYEKAISLDKSQ.......GDVKNNYGAFLCSQGEFARAYELFEQALTTP.NYYQQADTYANIICAE   149
NTDB id 1358 HI 0366 AAC22024.1               YQQQGQIENAFREYEIAVNLNHKQ.......GDVHNNFGTFLCSQKKFEQAQQQFELALNSP.NYYHQADTFENIVLCAY   149
consensus                                     !!**!** *!   !* !* ! **!       *!! !!*!*!!!*!* !**!***!! !!*** *!!*!!!!**!!**!!

logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1     MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 210474 BTV22 RS06550 WP 075294142.1   AEHNIPRYQKFLQLLKNIDPKRAEEIKKVS...   179
NTDB id 1358 HI 0366 AAC22024.1               SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                         ***** !  !**!! *!!!**  *
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X non conserved

X similar

X ≥ 50% conserved