```
logo
NTDB id 1358 HI 0366 AAC22024.1          ..MKTISKQLSAVIFPFIFSACVSQ...SASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHYYQQ   80
NTDB id 27622 APL RS06640 WP 011848557.1 MNLAKFLRNLTACMLLSVLFGCSSQLPTSQTTFNRSEAVKARINLALAYLDQHDLPKAKENIDKALSHDQNDYLPHSVLAYYYQQ   85
NTDB id 1386 A4U84 RS02735 WP 005710402.1 MTFAKFFRNLTACSVALWLVACANQ..TTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYYYQQ   83
consensus                                * *****  **!*!***  ** *! *!  ******!***!*!!!**!!!*!!*! * **!!*!*!!!! !! **!!*!!*!!*!!!!

logo
NTDB id 1358 HI 0366 AAC22024.1          QGQIENAFREYEIAVNLNH.......KQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAYSAQKMDIY  157
NTDB id 27622 APL RS06640 WP 011848557.1 IGDQVHAEQAYQQALKLSD.......NRPDVLNNYGTFLCKQAKFDQAYQQFEQAVKSEQPYYHQADSLENIVLCAKQEPNLAKV  163
NTDB id 1386 A4U84 RS02735 WP 005710402.1 TGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCANMKKDIEKQ  168
consensus                                 !**  !* *!**!**!*         **!!*!!*!!!!!*!**!**!*!!!! !* !   !!*!!!**!!!*!!!     ***

logo
NTDB id 1358 HI 0366 AAC22024.1          QQTLEKLRQIDGKRAEKFNSLK...  179
NTDB id 27622 APL RS06640 WP 011848557.1 NEALSQLEKLDKSRAAALR......  182
NTDB id 1386 A4U84 RS02735 WP 005710402.1 KTALSQLEKIDKSRAERLYNFMEIR  193
consensus                                * *!**!***!**!!***
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X non conserved

X similar

X ≥ 50% conserved