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logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1      MTFAKFFRNLTACSVALWLVACANQTTQVDFNRSEAVKARINLALAYLEQSDFPKAKENIDKALEHDVKDYLPHSVLAYY   80
NTDB id 1151736 LS075 RS03310 WP 011272006.1   ..MQTISKQLSAVIFPFIFSACVSQS.ASSLDHQAAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
NTDB id 1358 HI 0366 AAC22024.1                ..MKTISKQLSAVIFPFIFSACVSQS.ASSLNHQTAAKARVELALSYLQQNNPQLAKINLDKALQHDKNYYLVHSALAHY   77
consensus                                        * *****!*!********!**!* *******  !*!!!**!!!*!!*!*****!!*!*!!!!*!!***!!*!!*!!*!

logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1      YQQTGDNQKAEEAYQQALKLSKTQSKNNQVRPDVLNNYGTFLCKQKLFDKAYQQFETALTSQEAYYNQADTLENIALCAN   160
NTDB id 1151736 LS075 RS03310 WP 011272006.1   YQQQGQIENAFREYEIAVKLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIALCAY   149
NTDB id 1358 HI 0366 AAC22024.1                YQQQGQIENAFREYEIAVNLN.......HKQGDVHNNFGTFLCSQKKFEQAQQQFELALNS.PNYYHQADTFENIVLCAY   149
consensus                                      !!!*!****!***!**!**!*        ****!!*!!*!!!!!*!!*!**!*!!!!*!!*!  **!!*!!!!*!!!*!!!*

logo

NTDB id 1386 A4U84 RS02735 WP 005710402.1      MKKDIEKQKTALSQLEKIDKSRAERLYNFMEIR   193
NTDB id 1151736 LS075 RS03310 WP 011272006.1   SAKKMDIYQQTLEKLRQINGKRAEKFNSLK...   179
NTDB id 1358 HI 0366 AAC22024.1                SAQKMDIYQQTLEKLRQIDGKRAEKFNSLK...   179
consensus                                      ***********!**!**!***!!!******
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X non conserved

X similar

X ≥ 50% conserved