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logo

NTDB id 1383 A4U84 RS08530 WP 005711781.1      MKK.IRLIRPLAKAFTLIELMIVIAIIAILATIAIPSYTSYTQKAALSELLSASASYKSDVEICIYNVGDKK.......I   72
NTDB id 210443 BTV17 RS01355 WP 075293673.1    MKI.TYIFSSAKKGFTLIELMIVIAIVAILATIAIPSYQNYTKKAAITELLQASAPYRAEVELCVYNNDA.........P   70
NTDB id 1365 NTHI RS01955 WP 011271973.1       MKL.T.TQQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNE.........T   69
NTDB id 1354 HI 0299 AAC21963.1                MKL.T.TLQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNE.........T   69
NTDB id 1400 DSB67 RS12660 WP 010643261.1      ...MKTNKQKKQQGFTLIELMIVVAIIGILAALAIPAYQDYTKKATMAEFPKIASSTKLAVELCAHEHAADA....GAFK   73
NTDB id 1424 RS RS02795 WP 011000517.1         MKSMRHLNKRVQKGFTLIELMIVVAIVGILAAIAIPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSVGSSVFT   80
consensus                                      **        * **!!!!!!!!!*!!**!!!**!!!*!* !!**!***!** *** ****!*****

logo

NTDB id 1383 A4U84 RS08530 WP 005711781.1      SECSSGKNGVRADKSDVPDAKYLKSISVASGVITV.....AGKGNIDGY.....GYTMTPKF.ANNNITWATSCQGADIS   141
NTDB id 210443 BTV17 RS01355 WP 075293673.1    TNCNAGTNGIQAAINT..PKGKVKSVSVQGGAITV.....TGNGNLKDI.....TYTLTPTGDASTGVTWSANCNN.KAD   137
NTDB id 1365 NTHI RS01955 WP 011271973.1       TNCTGGKNGIAADITT..AKGYVKSVTTSNGAITV.....KGDGTLANM.....EYILQATGNAATGVTWTTTCKGTDAS   137
NTDB id 1354 HI 0299 AAC21963.1                TSCTGGKNGIAADIKT..AKGYVASVITQSGGITV.....KGNGTLANM.....EYILQAKGNAAAGVTWTTTCKGTDAS   137
NTDB id 1400 DSB67 RS12660 WP 010643261.1      TKCHQTANGIPAAFTLNN.LTVTPAAGAASGAISITVNAANDKCPIKQNE....KYILT.ATYASNGVTWSSECKDA...   144
NTDB id 1424 RS RS02795 WP 011000517.1         PT...........KNVAN.LTI.AGTGTVPGQITVTYTTAAGGGTLALVPTAAGTALPV.SSAPSGPIMWICYAQNKAQA   146
consensus                                      * *  * *** *          * **    !*!**      * ! *  *      ***     *  ***!** *

logo

NTDB id 1383 A4U84 RS08530 WP 005711781.1      L...FP.......ANFCGQKATP..   154
NTDB id 210443 BTV17 RS01355 WP 075293673.1    L...FP.......AGFCSSTSTNSP   152
NTDB id 1365 NTHI RS01955 WP 011271973.1       L...FP.......ANFCGSVTQ...   149
NTDB id 1354 HI 0299 AAC21963.1                L...FP.......ANFCGSVTK...   149
NTDB id 1400 DSB67 RS12660 WP 010643261.1      ........TNAVQTSYCP.......   154
NTDB id 1424 RS RS02795 WP 011000517.1         ASSVAPSGTMSLAAKYAPAECR...   168
consensus                                      *   **       * **
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X non conserved

X similar

X $\geq$ 50% conserved