logo

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NTDB id 1354 HI 0299 AAC21963.1                .MKLTTLQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNET.....TSCTG   74
NTDB id 44253 CHBNTHi RS04070 WP 005666683.1   .MKLTTQQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNET.....TNCTG   74
NTDB id 1365 NTHI RS01955 WP 011271973.1       .MKLTTQQTLKKGFTLIELMIVIAIIAILATIAIPSYQNYTKKAAVSELLQASAPYKADVELCVYSTNET.....TNCTG   74
NTDB id 1383 A4U84 RS08530 WP 005711781.1      MKKIRLIRPLAKAFTLIELMIVIAIIAILATIAIPSYTSYTQKAALSELLSASASYKSDVEICIYNVGDKK...ISECSS   77
NTDB id 1467 ASU2 RS05095 WP 014991735.1       ..MKHIIKPIQQAFTLIELMIVIAIIAILATVAIPSYNNYTKKAALSELLAASSAYKTDVEICIYNTGAT.....NSCFS   73
NTDB id 1400 DSB67 RS12660 WP 010643261.1      ..MKTNKQKKQQGFTLIELMIVVAIIGILAALAIPAYQDYTKKATMAEFPKIASSTKLAVELCAHEHAADAGAFKTKCHQ   78
NTDB id 1274 LPP RS09530 WP 011214168.1        ........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNAL....PA..TQ   66
NTDB id 1258 GCO85 RS09745 WP 014844212.1      ........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNAL....PA..TQ   66
consensus                                                * **!!!!!!!!!*!!!*!!!**!!!*!* !!**!***!** **  *! *!****** *       * **
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NTDB id 1354 HI 0299 AAC21963.1                GKNGIAADIKT..AKGYVASVITQSGGIT.....VKGNGTLANMEYILQAKGNAAAGVTWTTTCKGTDASLFPANFCGSV   147
NTDB id 44253 CHBNTHi RS04070 WP 005666683.1   GKNGIAADITT..AKGYVASVKTQSGGIT.....VKGDGTLANMEYILQATGNAATGVTWTTTCKGTEASLFPANFCGSV   147
NTDB id 1365 NTHI RS01955 WP 011271973.1       GKNGIAADITT..AKGYVKSVTTSNGAIT.....VKGDGTLANMEYILQATGNAATGVTWTTTCKGTDASLFPANFCGSV   147
NTDB id 1383 A4U84 RS08530 WP 005711781.1      GKNGVRADKSDVPDAKYLKSISVASGVIT.....VAGKGNIDGYGYTMTPK.FANNNITWATSCQGADISLFPANFCGQK   151
NTDB id 1467 ASU2 RS05095 WP 014991735.1       GANGIKN.GDDLKNSKYLDSVSVTAGTIT.....VNGKGSVAGYGYTLTPD.YSSKTLTWKTTCSGEDKSLFPAAFCSE.   145
NTDB id 1400 DSB67 RS12660 WP 010643261.1      TANGIPAAFT.LNNLTVTPAAGAASGAISITVNAANDKGPIKQNEKYILTATYASNGVTWSSECKDATNAVQT.SYCP..   154
NTDB id 1274 LPP RS09530 WP 011214168.1        AATGYVSPAA.TP..NVASIAIGANGVITITYTAAAGGGTIVMT.....PTLQANGDVTWTCTGGTLLAKYRP.ASCRP.   136
NTDB id 1258 GCO85 RS09745 WP 014844212.1      AATGYVSPAA.TP..NVQSIAIGANGVITITYTAAAGGGTIIMT.....PTLQANGDVTWTCTGGTLLAKYRP.ASCRP.   136
consensus                                      * *!* *          ** **    ! !*     * * !**    * *    **   *!!**** * *******  *!
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NTDB id 1354 HI 0299 AAC21963.1                TK.   149
NTDB id 44253 CHBNTHi RS04070 WP 005666683.1   TQ.   149
NTDB id 1365 NTHI RS01955 WP 011271973.1       TQ.   149
NTDB id 1383 A4U84 RS08530 WP 005711781.1      ATP   154
NTDB id 1467 ASU2 RS05095 WP 014991735.1       ...   145
NTDB id 1400 DSB67 RS12660 WP 010643261.1      ...   154
NTDB id 1274 LPP RS09530 WP 011214168.1        ...   136
NTDB id 1258 GCO85 RS09745 WP 014844212.1      ...   136
consensus
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X non conserved

X similar

X ≥ 50% conserved