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NTDB id 98 BSU 24710 NP 390351.1            ..................MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLAD   68
NTDB id 619 LCA RS06470 WP 011374999.1      ................MKKKRKAFTLIEMVVVLAIIALLILLVAPNLGAQKQRAEKKTDQALVTTLQTQVELASDEIGHQIKSLDE   70
NTDB id 32 MW RS07995 WP 000472256.1        ..........MFKF..LKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIED   73
NTDB id 16 SA RS07765 WP 000472256.1        ..........MFKF..LKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIED   73
NTDB id 588 KW2 RS10845 WP 051303741.1      MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDKPNLSE   85
NTDB id 69429 SR187 RS00800 WP 024532254.1  ..........MKKIIEMKNKAFTLVEMLVVLLIISVLLLLFVPNLSKHSETVQKKGNEAVVKIVESQQELYQLEHG.HQPSAEE   73
NTDB id 460 GPW51 RS00790 WP 011921720.1    ..........MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHD.EEATVAD   73
NTDB id 502 SM12261 RS00955 WP 000738641.1  .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSK   77
NTDB id 162 SP RS10370 WP 000738627.1       .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK   77
NTDB id 272 KZH43 RS09255 WP 000738626.1    .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK   77
NTDB id 231 SPD RS09885 WP 000738626.1      .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK   77
NTDB id 197 SPR RS09420 WP 000738626.1      .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK   77
NTDB id 530 SMSK321 RS09230 WP 000738643.1  .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSK   77
NTDB id 431 SGO RS09410 WP 012130940.1      ..........MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSK   74
NTDB id 396 FSA28 RS09395 WP 002263437.1    ..........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK   73
NTDB id 387 SMU RS09025 WP 002263437.1      ..........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK   73
consensus                                             *  *** ****!!!*!!**!!*!!**!******!!******* ***** **********!**** **    * *** *
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NTDB id 98 BSU 24710 NP 390351.1            LQSEGYVKKDAV.CPNGKRIIITGGEVKVEH    98
NTDB id 619 LCA RS06470 WP 011374999.1      L.TDKYISKDQLKHAKEKGITIEGGTVKQK.    99
NTDB id 32 MW RS07995 WP 000472256.1        LIADGFIKEAQKTCKSGETITIS.NGEAVAN   103
NTDB id 16 SA RS07765 WP 000472256.1        LIADGFIKEAQKTCKSGETITIS.NGEAVAN   103
NTDB id 588 KW2 RS10845 WP 051303741.1      LLSAGMITQKQVTAYDDYYDQNKNEQRNFDD   116
NTDB id 69429 SR187 RS00800 WP 024532254.1  LKNKGYITEKQLEQYQNAKK.............    93
NTDB id 460 GPW51 RS00790 WP 011921720.1    LQAKGYITEKQAKQYATAKK.............    93
NTDB id 502 SM12261 RS00955 WP 000738641.1  LQADGRITEEQAKAYKEYHAKQNTSQTVAD.   107
NTDB id 162 SP RS10370 WP 000738627.1       LQADGRITEEQAKAYKEYNDKNGGANRKVND   108
NTDB id 272 KZH43 RS09255 WP 000738626.1    LQADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 231 SPD RS09885 WP 000738626.1      LQADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 197 SPR RS09420 WP 000738626.1      LQADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 530 SMSK321 RS09230 WP 000738643.1  LQADGRITEEQAKAYNEYYAKNGGANRKVND   108
NTDB id 431 SGO RS09410 WP 012130940.1      LVAAGNISQKQADSYKAYYGKNNSETQAVAN   105
NTDB id 396 FSA28 RS09395 WP 002263437.1    LVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1      LVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
consensus                                   ! *** ******  ****  *       *
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X non conserved
X similar
X ≥ 50% conserved