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logo
NTDB id 98 BSU 24710 NP 390351.1            ..........MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGYV   75
NTDB id 73316 SAP2 RS06590 WP 103387866.1   .....MY.KRTVTKAFTLIEMLLVLLIISLLLILIIPNIAKQSKHIQNTGCKAQLKMVDSQIEAYTLKNNQAPATIDDLVREGYI   79
NTDB id 32 MW RS07995 WP 000472256.1        .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
NTDB id 16 SA RS07765 WP 000472256.1        .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
NTDB id 460 GPW51 RS00790 WP 011921720.1    ....MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHD.EEATVADLQAKGYI   80
NTDB id 502 SM12261 RS00955 WP 000738641.1  MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSKLQADGRI   84
NTDB id 162 SP RS10370 WP 000738627.1       MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQADGRI   84
NTDB id 272 KZH43 RS09255 WP 000738626.1    MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQADGRI   84
NTDB id 231 SPD RS09885 WP 000738626.1      MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQADGRI   84
NTDB id 197 SPR RS09420 WP 000738626.1      MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQADGRI   84
NTDB id 530 SMSK321 RS09230 WP 000738643.1  MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSKLQADGRI   84
NTDB id 431 SGO RS09410 WP 012130940.1      ....MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSKLVAAGNI   81
NTDB id 396 FSA28 RS09395 WP 002263437.1    ....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSKLVASGDI   80
NTDB id 387 SMU RS09025 WP 002263437.1      ....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSKLVASGDI   80
consensus                                     * **** ****!!!*!!!*!!*!!!*!*****!!************! ******!**!**** !*** * *** *!***! *
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logo
NTDB id 98 BSU 24710 NP 390351.1            KKDAV.CPNGKRIIITGGEVKVEH    98
NTDB id 73316 SAP2 RS06590 WP 103387866.1   KENQKQCKSGANIT.ITNGEAVAS   102
NTDB id 32 MW RS07995 WP 000472256.1        KEAQKTCKSGETIT.ISNGEAVAN   103
NTDB id 16 SA RS07765 WP 000472256.1        KEAQKTCKSGETIT.ISNGEAVAN   103
NTDB id 460 GPW51 RS00790 WP 011921720.1    TEKQAKQYATAKK...........    93
NTDB id 502 SM12261 RS00955 WP 000738641.1  TEEQAKAYKEYHAKQNTSQTVAD.   107
NTDB id 162 SP RS10370 WP 000738627.1       TEEQAKAYKEYNDKNGGANRKVND   108
NTDB id 272 KZH43 RS09255 WP 000738626.1    TEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 231 SPD RS09885 WP 000738626.1      TEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 197 SPR RS09420 WP 000738626.1      TEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 530 SMSK321 RS09230 WP 000738643.1  TEEQAKAYNEYYAKNGGANRKVND   108
NTDB id 431 SGO RS09410 WP 012130940.1      SQKQADSYKAYYGKNNSETQAVAN   105
NTDB id 396 FSA28 RS09395 WP 002263437.1    TQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1      TQEQANSYKDYYAKHPSKTRHVAN   104
consensus                                   *****   ****   *        *
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X non conserved

X similar

X ≥ 50% conserved