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logo

NTDB id 588 KW2 RS10845 WP 051303741.1        MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDK   80
NTDB id 460 GPW51 RS00790 WP 011921720.1      ...........MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHD.EE   68
NTDB id 502 SM12261 RS00955 WP 000738641.1    .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.ED   72
NTDB id 530 SMSK321 RS09230 WP 000738643.1    .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.ED   72
NTDB id 431 SGO RS09410 WP 012130940.1        ...........MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQ   69
NTDB id 396 FSA28 RS09395 WP 002263437.1      ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS   68
NTDB id 387 SMU RS09025 WP 002263437.1        ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS   68
NTDB id 98 BSU 24710 NP 390351.1              ..................MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQT   63
NTDB id 236156 CEP64 RS19660 WP 037588275.1   ..........MKRFKFLSKKEGFTLVEMLIVLLIISILLVIIIPNISKQSETVKSTGCQAQVKMVAGQVEAYTLKHDAKP   70
NTDB id 32 MW RS07995 WP 000472256.1          ............MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP   68
NTDB id 16 SA RS07765 WP 000472256.1          ............MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP   68
consensus                                                 *  *   ****!!!*!!!*!!*!!!*!*****!!******  **  *!  *****!**!*****!  *  *

logo

NTDB id 588 KW2 RS10845 WP 051303741.1        PNLSELLSAGMITQKQVTAYDDYYDQNKNEQRNFDD   116
NTDB id 460 GPW51 RS00790 WP 011921720.1      ATVADLQAKGYITEKQAKQYATAKK...........    93
NTDB id 502 SM12261 RS00955 WP 000738641.1    ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVAD.   107
NTDB id 530 SMSK321 RS09230 WP 000738643.1    ASLSKLQADGRITEEQAKAYNEYYAKNGGANRKVND   108
NTDB id 431 SGO RS09410 WP 012130940.1        ATLSKLVAAGNISQKQADSYKAYYGKNNSETQAVAN   105
NTDB id 396 FSA28 RS09395 WP 002263437.1      ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1        ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 98 BSU 24710 NP 390351.1              PSLADLQSEGYVKKD.AVCPNGKRIIITGGEVKVEH    98
NTDB id 236156 CEP64 RS19660 WP 037588275.1   NSINDLVSDGYINQKQLTCKSGETITLENGEVVAH.   105
NTDB id 32 MW RS07995 WP 000472256.1          SSIEDLIADGFIKEAQKTCKSGETITISNGEAVAN.   103
NTDB id 16 SA RS07765 WP 000472256.1          SSIEDLIADGFIKEAQKTCKSGETITISNGEAVAN.   103
consensus                                     *****!***!  **   **   *   *
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X non conserved

X similar

X ≥ 50% conserved