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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 1027118 AB3N25 RS07360 WP 058611611.1   ..........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          PNLSELLSACMITQKQVTAYDDYYDQNKNEQRNFDD   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        ATLSKLVASCDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1          ATLSKLVASCDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 98 BSU 24710 NP 390351.1                PSLADLQSEGYVKKD.AVCPNGKRIIITGGEVKVEH    98
NTDB id 1027118 AB3N25 RS07360 WP 058611611.1   NSINDLISDGYINQKQLTCKSGETITLENGEVVAH.   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