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logo
NTDB id 98 BSU 24710 NP 390351.1              ..........MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGYV   75
NTDB id 117756 BE24 RS04870 WP 029377192.1    ...MNNFKNKFNKKAFTLIEMLLVLLIISLLLILIIPNIAKQSSHIQTTSCEAQLKMVDSQIEAYSLKFNKKPTTMDELVNEGYI   82
NTDB id 32 MW RS07995 WP 000472256.1          .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
NTDB id 16 SA RS07765 WP 000472256.1          .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   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              KKDA.VCPNGKRIIITGGEVKVEH   98
NTDB id 117756 BE24 RS04870 WP 029377192.1    KENQKQCKSGALISI.NDGEAVAN  105
NTDB id 32 MW RS07995 WP 000472256.1          KEAQKTCKSGETITI.SNGEAVAN  103
NTDB id 16 SA RS07765 WP 000472256.1          KEAQKTCKSGETITI.SNGEAVAN  103
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