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logo
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NTDB id 460 GPW51 RS00790 WP 011921720.1        ....MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHDEE.ATVADLQ    75
NTDB id 502 SM12261 RS00955 WP 000738641.1      MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED.ASLSKLQ    79
NTDB id 162 SP RS10370 WP 000738627.1           MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNED.ASLRKLQ    79
NTDB id 272 KZH43 RS09255 WP 000738626.1        MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNED.ASLRKLQ    79
NTDB id 231 SPD RS09885 WP 000738626.1          MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNED.ASLRKLQ    79
NTDB id 197 SPR RS09420 WP 000738626.1          MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNED.ASLRKLQ    79
NTDB id 530 SMSK321 RS09230 WP 000738643.1      MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED.ASLSKLQ    79
NTDB id 98 BSU 24710 NP 390351.1                ..........MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQ    70
NTDB id 1055584 ACFDHX RS05020 WP 107632995.1   MLKIINIL..KNKKGFTLIEMLLVLLIISVLLILIIPNIAKQSDHIQKTGCAAQLKLVDSQIEAYTLKYNRKPTSIDELI    78
NTDB id 32 MW RS07995 WP 000472256.1            ...MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLI    75
NTDB id 16 SA RS07765 WP 000472256.1            ...MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLI    75
consensus                                       ****************!!!*!!!*!!*!!!*!*****!!***********!*******!**!*****!*****  ***  *!*
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logo
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NTDB id 460 GPW51 RS00790 WP 011921720.1        AKGYITEKQAKQYATAKK...........    93
NTDB id 502 SM12261 RS00955 WP 000738641.1      ADGRITEEQAKAYKEYHAKQNTSQTVAD.   107
NTDB id 162 SP RS10370 WP 000738627.1           ADGRITEEQAKAYKEYNDKNGGANRKVND   108
NTDB id 272 KZH43 RS09255 WP 000738626.1        ADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 231 SPD RS09885 WP 000738626.1          ADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 197 SPR RS09420 WP 000738626.1          ADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 530 SMSK321 RS09230 WP 000738643.1      ADGRITEEQAKAYNEYYAKNGGANRKVND   108
NTDB id 98 BSU 24710 NP 390351.1                SEGYVKKD.AVCPNGKRIIITGGEVKVEH    98
NTDB id 1055584 ACFDHX RS05020 WP 107632995.1   QEGYIKDNQKNCKSGETITIRGGEAVVS.   106
NTDB id 32 MW RS07995 WP 000472256.1            ADGFIKEAQKTCKSGETITISNGEAVAN.   103
NTDB id 16 SA RS07765 WP 000472256.1            ADGFIKEAQKTCKSGETITISNGEAVAN.   103
consensus                                       **!**********  **   **  ***  ***
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X non conserved

X similar

X $\geq$ 50% conserved