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logo

NTDB id 338989 EQ029 RS06420 WP 057504819.1   MKNHFKNLKK..LKAFTLIEMLLVLLIISVLLILIIPNIAKQTAHIQSTGCEAQVKMINSQIEAYTLKHNRTPSNINDLI   78
NTDB id 32 MW RS07995 WP 000472256.1          ...MFKFLKK..TQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLI   75
NTDB id 16 SA RS07765 WP 000472256.1          ...MFKFLKK..TQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLI   75
NTDB id 502 SM12261 RS00955 WP 000738641.1    MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSKLQ   79
NTDB id 162 SP RS10370 WP 000738627.1         MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQ   79
NTDB id 272 KZH43 RS09255 WP 000738626.1      MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQ   79
NTDB id 231 SPD RS09885 WP 000738626.1        MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQ   79
NTDB id 197 SPR RS09420 WP 000738626.1        MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRKLQ   79
NTDB id 530 SMSK321 RS09230 WP 000738643.1    MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSKLQ   79
NTDB id 431 SGO RS09410 WP 012130940.1        ....MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSKLV   76
NTDB id 396 FSA28 RS09395 WP 002263437.1      ....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSKLV   75
NTDB id 387 SMU RS09025 WP 002263437.1        ....MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSKLV   75
consensus                                     ** ** **!* ***!!!!*!!!*!!!!!!*!**!**!!**!!****** ! *!*!!***!!*!*!*! ** * *** *!

logo

NTDB id 338989 EQ029 RS06420 WP 057504819.1   NDGLIKEEQKQCRSGETIT.ISNGEAIAN   106
NTDB id 32 MW RS07995 WP 000472256.1          ADGFIKEAQKTCKSGETIT.ISNGEAVAN   103
NTDB id 16 SA RS07765 WP 000472256.1          ADGFIKEAQKTCKSGETIT.ISNGEAVAN   103
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 431 SGO RS09410 WP 012130940.1        AAGNISQKQADSYKAYYGKNNSETQAVAN   105
NTDB id 396 FSA28 RS09395 WP 002263437.1      ASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1        ASGDITQEQANSYKDYYAKHPSKTRHVAN   104
consensus                                     **! !***!*  ****  ** *  * *
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X non conserved

X similar

X $\geq$ 50% conserved