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logo

NTDB id 98 BSU 24710 NP 390351.1            ..........MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQTPSLADLQSEGYV   75
NTDB id 63331 STP1 RS00600 WP 023372658.1   ...MIKPNHSFKTKAFTLIEMLLVLLIISLLLILIIPNVAKQTANIQSTGCDAQVKMINSQIEAYSLKHNRNPNTIQDLITEGYI   82
NTDB id 32 MW RS07995 WP 000472256.1        .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
NTDB id 16 SA RS07765 WP 000472256.1        .....MFKFLKKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIEDLIADGFI   80
NTDB id 460 GPW51 RS00790 WP 011921720.1    ....MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHDE.EATVADLQAKGYI   80
NTDB id 431 SGO RS09410 WP 012130940.1      ....MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSKLVAAGNI   81
NTDB id 502 SM12261 RS00955 WP 000738641.1  MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNE.DASLSKLQADGRI   84
NTDB id 162 SP RS10370 WP 000738627.1       MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNE.DASLRKLQADGRI   84
NTDB id 530 SMSK321 RS09230 WP 000738643.1  MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNE.DASLSKLQADGRI   84
consensus                                       * **   ****!!!*!!!*!!*!!!*!*****!!*********  *!  *********!****  !***    ***  *!***!**
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logo

NTDB id 98 BSU 24710 NP 390351.1            KKDA.VCPNGKRIIITGGEVKVEH   98
NTDB id 63331 STP1 RS00600 WP 023372658.1   KENQKSCKTGETISI.SNGEAIAN  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 460 GPW51 RS00790 WP 011921720.1    TEKQAKQYATAKK...........   93
NTDB id 431 SGO RS09410 WP 012130940.1      SQKQADSYKAYYGKNNSETQAVAN  105
NTDB id 502 SM12261 RS00955 WP 000738641.1  TEEQAKAYKEYHAKQNTSQTVAD.  107
NTDB id 162 SP RS10370 WP 000738627.1       TEEQAKAYKEYNDKNGGANRKVND  108
NTDB id 530 SMSK321 RS09230 WP 000738643.1  TEEQAKAYNEYYAKNGGANRKVND  108
consensus                                   ** **  * *       *    *
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X non conserved

X similar

X $\geq$ 50% conserved