```
logo
NTDB id 32 MW RS07995 WP 000472256.1              ..........MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALK    63
NTDB id 16 SA RS07765 WP 000472256.1              ..........MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALK    63
NTDB id 588 KW2 RS10845 WP 051303741.1            MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELD    75
NTDB id 1111661 HMPREF9319 RS00395 WP 003062921.1 .......MKKIWKKLRKKSVAAFTLVEMLIVLLIISVLMLLFVPNLSKQKDVVREKGDAAVVKVVESQMDLYELK    68
NTDB id 460 GPW51 RS00790 WP 011921720.1          ...........MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELE    64
NTDB id 502 SM12261 RS00955 WP 000738641.1        .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLD    68
NTDB id 272 KZH43 RS09255 WP 000738626.1          .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLE    68
NTDB id 231 SPD RS09885 WP 000738626.1            .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLE    68
NTDB id 197 SPR RS09420 WP 000738626.1            .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLE    68
NTDB id 162 SP RS10370 WP 000738627.1             .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLE    68
NTDB id 530 SMSK321 RS09230 WP 000738643.1        .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLD    68
NTDB id 431 SGO RS09410 WP 012130940.1            ...........MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELK    64
NTDB id 396 FSA28 RS09395 WP 002263437.1          ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELN    64
NTDB id 387 SMU RS09025 WP 002263437.1            ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELN    64
consensus                                                ** ** **** ****!!!*!!!*!!*!!!*!**!**!!*********** *!*!!*!*!!***! !*
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logo
NTDB id 32 MW RS07995 WP 000472256.1              HNRNPSSIEDLIADGFIKEAQKTCKSGETITIS.NGEAVAN   103
NTDB id 16 SA RS07765 WP 000472256.1              HNRNPSSIEDLIADGFIKEAQKTCKSGETITIS.NGEAVAN   103
NTDB id 588 KW2 RS10845 WP 051303741.1            HDNDKPNLSELLSAGMITQKQVTAYDDYYDQNKNEQRNFDD   116
NTDB id 1111661 HMPREF9319 RS00395 WP 003062921.1 TG.DKPTVDDLVEVGYITSEQAKTYNEAKK...........    97
NTDB id 460 GPW51 RS00790 WP 011921720.1          HD.EEATVADLQAKGYITEKQAKQYATAKK...........    93
NTDB id 502 SM12261 RS00955 WP 000738641.1        KN.EDASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVAD.   107
NTDB id 272 KZH43 RS09255 WP 000738626.1          KN.EDASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 231 SPD RS09885 WP 000738626.1            KN.EDASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 197 SPR RS09420 WP 000738626.1            KN.EDASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND   108
NTDB id 162 SP RS10370 WP 000738627.1             KN.EDASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVND   108
NTDB id 530 SMSK321 RS09230 WP 000738643.1        KN.EDASLSKLQADGRITEEQAKAYNEYYAKNGGANRKVND   108
NTDB id 431 SGO RS09410 WP 012130940.1            NTGDQATLSKLVAAGNISQKQADSYKAYYGKNNSETQAVAN   105
NTDB id 396 FSA28 RS09395 WP 002263437.1          KT.DSATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1            KT.DSATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
consensus                                         ** * *** *! **! !***!** ****  **     * *
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X non conserved

X similar

X ≥50% conserved