```
logo
                                                 MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDKPNLSE
NTDB id 64567 SPB RS09095 WP 003104947.1     .......MKKFIINFKNKSVQAFTLLEMLMVILIISILMLLFIPNLNKQKEKVLNTGGEALVKIVENQAELYALNNAGKVPSLND    78
NTDB id 588 KW2 RS10845 WP 051303741.1       MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDKPNLSE    85
NTDB id 502 SM12261 RS00955 WP 000738641.1   .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSK    77
NTDB id 272 KZH43 RS09255 WP 000738626.1     .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK    77
NTDB id 231 SPD RS09885 WP 000738626.1       .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK    77
NTDB id 197 SPR RS09420 WP 000738626.1       .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK    77
NTDB id 162 SP RS10370 WP 000738627.1        .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK    77
NTDB id 530 SMSK321 RS09230 WP 000738643.1   .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSK    77
NTDB id 431 SGO RS09410 WP 012130940.1       ...........MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSK    74
NTDB id 396 FSA28 RS09395 WP 002263437.1     ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK    73
NTDB id 387 SMU RS09025 WP 002263437.1       ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK    73
consensus                                               **************!!!!*!!!*!**!!!*!*!!!*!!!*!*!**!***!**!*!!*!!*!!*!!*!*!*** ****!**
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logo
                                                 LQADGRITEEQAKAYKEYYDKNGGANRKVND
NTDB id 64567 SPB RS09095 WP 003104947.1     LKSNVNLTEKQITAYNDYYKKHSDKIPNVK.    108
NTDB id 588 KW2 RS10845 WP 051303741.1       LLSAGMITQKQVTAYDDYYDQNKNEQRNFDD    116
NTDB id 502 SM12261 RS00955 WP 000738641.1   LQADGRITEEQAKAYKEYHAKQNTSQTVAD.    107
NTDB id 272 KZH43 RS09255 WP 000738626.1     LQADGRITEEQAKAYKEYHDKNGGANRKVND    108
NTDB id 231 SPD RS09885 WP 000738626.1       LQADGRITEEQAKAYKEYHDKNGGANRKVND    108
NTDB id 197 SPR RS09420 WP 000738626.1       LQADGRITEEQAKAYKEYHDKNGGANRKVND    108
NTDB id 162 SP RS10370 WP 000738627.1        LQADGRITEEQAKAYKEYNDKNGGANRKVND    108
NTDB id 530 SMSK321 RS09230 WP 000738643.1   LQADGRITEEQAKAYNEYYAKNGGANRKVND    108
NTDB id 431 SGO RS09410 WP 012130940.1       LVAAGNISQKQADSYKAYYGKNNSETQAVAN    105
NTDB id 396 FSA28 RS09395 WP 002263437.1     LVASGDITQEQANSYKDYYAKHPSKTRHVAN    104
NTDB id 387 SMU RS09025 WP 002263437.1       LVASGDITQEQANSYKDYYAKHPSKTRHVAN    104
consensus                                    !*********!***!**!* **   ** * *
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X non conserved

X similar

X ≥50% conserved