```
logo

NTDB id 588 KW2 RS10845 WP 051303741.1        MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDK   80
NTDB id 502 SM12261 RS00955 WP 000738641.1    .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.ED   72
NTDB id 530 SMSK321 RS09230 WP 000738643.1    .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.ED   72
NTDB id 431 SGO RS09410 WP 012130940.1        ...........MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQ   69
NTDB id 396 FSA28 RS09395 WP 002263437.1      ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS   68
NTDB id 387 SMU RS09025 WP 002263437.1        ...........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DS   68
NTDB id 204921 BI355 RS03285 WP 057866609.1   ................MKKIHKAFTLIEMVIVLFIISLLLLIMIPNLVAQKNHAEDRSQEAFITTIKTQKELYSDNE.GKI   64
NTDB id 619 LCA RS06470 WP 011374999.1        ................MKKKRKAFTLIEMVVVLAIIALLILLVAPNLGAQKQRAEKKTDQALVTTLQTQVELASDEI.GHQ   64
consensus                                                *  *** ***!!!!*!!**!!*!!**!*!***!!!***!* * * *  !*******!**!* *    *

logo

NTDB id 588 KW2 RS10845 WP 051303741.1        PNLSELLSAGMITQKQVTAYDDYYDQNKNEQRNFDD   116
NTDB id 502 SM12261 RS00955 WP 000738641.1    ASLSKLQADCRITEEQAKAYKEYHAKQNTSQTVAD.   107
NTDB id 530 SMSK321 RS09230 WP 000738643.1    ASLSKLQADCRITEEQAKAYNEYYAKNGGANRKVND   108
NTDB id 431 SGO RS09410 WP 012130940.1        ATLSKLVAACNISQKQADSYKAYYGKNNSETQAVAN   105
NTDB id 396 FSA28 RS09395 WP 002263437.1      ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 387 SMU RS09025 WP 002263437.1        ATLSKLVASGDITQEQANSYKDYYAKHPSKTRHVAN   104
NTDB id 204921 BI355 RS03285 WP 057866609.1   PSLGELEKEKYITQKQYEKAQKIDELK.........    91
NTDB id 619 LCA RS06470 WP 011374999.1        IKSLDELTDKYISKDQLKHAKEKGITIEGGTVKQK.    99
consensus                                     ******** * !***!*  *******
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X non conserved

X similar

X ≥ 50% conserved