

NTDB id	32 MW RS07995 WP 000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	16 SA RS07765 WP 000472256.1	MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	588 KW2 RS10845 WP 051303741.1	MSKALSLVKIHGRKLCKKKQAFTLIEMLIVLAIIISILILLFVPNLIKEKAQVQKTGEAAVVKVVSQAQLYELDHDNDK	80
NTDB id	431 SGO RS09410 WP 012130940.1	MKKLKLLVKAFTLVEMLVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVSQAELYELKNTGDQ	69
NTDB id	396 FSA28 RS09395 WP 002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILLLLFVPNLTKQDKDKVDGTGSAVVVKVVSQAELYELNKTD.S	68
NTDB id	387 SMU RS09025 WP 002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILLLLFVPNLTKQDKDKVDGTGSAVVVKVVSQAELYELNKTD.S	68
NTDB id	460 GPW51 RS00790 WP 011921720.1	MKKLLEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVSQMELYELEHDEE	68
NTDB id	241556 CHF41 RS02725 WP 075105262.1	MKKLKKIKVKGFTLIEMLVLLIISILLLLFVPNLSKKKENAKQDTGNAVVKVVEGQAELYELQNE.GE	68
NTDB id	502 SM1261 RS00955 WP 000738641.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFFFVPNLTKQKEAVNDKGAAVVKVVSQAELYSLDKN.ED	72
NTDB id	272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFFFVPNLTKQKEAVNDKGAAVVKVVSQAELYSLDKN.ED	72
NTDB id	231 SPD RS09885 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFFFVPNLTKQKEAVNDKGAAVVKVVSQAELYSLDKN.ED	72
NTDB id	197 SPR RS09420 WP 000738626.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFFFVPNLTKQKEAVNDKGAAVVKVVSQAELYSLDKN.ED	72
NTDB id	162 SP RS10370 WP 000738627.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFFFVPNLTKQKEAVNDKGAAVVKVVSQAELYSLDKN.ED	72
NTDB id	530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKAQVKAFTLVEMLVLLIISVLFFFVPNLTKQKEAVNDKGAAVVKVVSQAELYSLDKN.ED	72
consensus		* * * * *	

NTDB id	32 MW RS07995 WP 000472256.1	SSTIEDLIADGFIKEAQKTKSGETITISNG.EAVAN	103
NTDB id	16 SA RS07765 WP 000472256.1	SSTIEDLIADGFIKEAQKTKSGETITISNG.EAVAN	103
NTDB id	588 KW2 RS10845 WP 051303741.1	PNLSELLSACMITQKQVTAAYDDYDQNKNEQRNFDD	116
NTDB id	431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYKYAYGKNNSETAQVAN	105
NTDB id	396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEASNYSKYDYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	ATLSKLVASGDITQEASNYSKYDYAKHPSKTRHVAN	104
NTDB id	460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id	241556 CHF41 RS02725 WP 075105262.1	ASLSRLVETGAITEAQAQAYNEYEYAKNTNQSRTVKP	104
NTDB id	502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKQNTNSTQVAD.	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHYDKNGGANRKVND	108
NTDB id	231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHYDKNGGANRKVND	108
NTDB id	197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHYDKNGGANRKVND	108
NTDB id	162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYHYDKNGGANRKVND	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYEYAKNGGANRKVND	108
consensus		*** * ! ** ! ! **** ! ***** ** *	

- X non conserved
- X similar
- X $\geq 50\%$ conserved