

NTDB id	32 MW RS07995 WP 000472256.1	MFKFL..KKTKQAFTLIEMLVLLIIISLILLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	16 SA RS07765 WP 000472256.1	MFKFL..KKTKQAFTLIEMLVLLIIISLILLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id	588 KW2 RS10845 WP 051303741.1	MSKALSLVKIHGRLCKKKQKAFTLIEMLVLAIIISLILLFVPNLIKBAQVQKTGEAAVVKVVSQAQLYELDHNDK	80
NTDB id	1127548 FGL06 RS00635 WP 021141433.1	MKKLWKKLRKKSVEAFTLVLEMLIVLLIICVLMLFVPNLISKQKEATHEKGDAAVVKVVSQKELYEVKTG.KE	72
NTDB id	460 GPW51 RS00790 WP 011921720.1	MKKLVMKVKCFITLVLEMLVVLGIISLILLFVPNLSQQKDAIQKGDAAAVVKVVSQKMELEYEHD.EE	68
NTDB id	502 SM12261 RS00955 WP 000738641.1	MKKMMTFLKKA KVKAFTLVLEMLVLLIIISVLILLFVPNLTKQKEAVNDKGKAAAVVKVVSQAELYSLDKN.ED	72
NTDB id	162 SP RS10370 WP 000738627.1	MKKMMTFLKKA KVKAFTLVLEMLVLLIIISVLILLFVPNLTKQKEAVNDKGKAAAVVKVVSQAELYSLEKN.ED	72
NTDB id	272 KZH43 RS09255 WP 000738626.1	MKKMMTFLKKA KVKAFTLVLEMLVLLIIISVLILLFVPNLTKQKEAVNDKGKAAAVVKVVSQAELYSLEKN.ED	72
NTDB id	231 SPD RS09885 WP 000738626.1	MKKMMTFLKKA KVKAFTLVLEMLVLLIIISVLILLFVPNLTKQKEAVNDKGKAAAVVKVVSQAELYSLEKN.ED	72
NTDB id	197 SPR RS09420 WP 000738626.1	MKKMMTFLKKA KVKAFTLVLEMLVLLIIISVLILLFVPNLTKQKEAVNDKGKAAAVVKVVSQAELYSLEKN.ED	72
NTDB id	530 SMSK321 RS09230 WP 000738643.1	MKKMMTFLKKA KVKAFTLVLEMLVLLIIISVLILLFVPNLTKQKEAVNDKGKAAAVVKVVSQAELYSLDKN.ED	72
NTDB id	431 SGO RS09410 WP 012130940.1	MNKLKKLRVKAFTLVLEMLVLLIIISVLMILLFVPNLTKQKEAVSDTGNAAAVVKVVSQAELYELKNTGDQ	69
NTDB id	396 FSA28 RS09395 WP 002263437.1	MKKTKRVSVRRAFTLIEMLVLLIIISLILLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNKT.DS	68
NTDB id	387 SMU RS09025 WP 002263437.1	MKKTKRVSVRRAFTLIEMLVLLIIISLILLFVPNLTKQKDKVKDTGGSAVVKVVSQAELYELNKT.DS	68
consensus		*** ** * **** * *** ** * ** * ** * ** * ** * ** * ** * ** * ** *	

NTDB id	32 MW RS07995 WP 000472256.1	SSIEDLIADGFIKEAQKTCKSGETITISN.GEAVAN	103
NTDB id	16 SA RS07765 WP 000472256.1	SSIEDLIADGFIKEAQKTCKSGETITISN.GEAVAN	103
NTDB id	588 KW2 RS10845 WP 051303741.1	PNLSELLSAGMITQKQVTAIDDYDQNKNENRNFDD	116
NTDB id	1127548 FGL06 RS00635 WP 021141433.1	ATIKDLVKEGYISQQQAQTYNEKAAPKE.....	100
NTDB id	460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVAD.	107
NTDB id	162 SP RS10370 WP 000738627.1	ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVND	108
NTDB id	272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id	231 SPD RS09885 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id	197 SPR RS09420 WP 000738626.1	ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND	108
NTDB id	530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGRITEEQAKAYNEYAKNGGANRKVND	108
NTDB id	431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNISQKQADSYPKAYYGKINSETQAVAN	105
NTDB id	396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
NTDB id	387 SMU RS09025 WP 002263437.1	ATLSKLVASGDITQEANSYKDYYAKHPSKTRHVAN	104
consensus		*** *! *** !****! * **** *	

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