

NTDB id 98	BSU 24710 NP 390351.1	MNEK G CFTLVEMLIVLFIIISILLITIPNVTKHNQTIQKGCEGLQNMVKAQMATAFELDHEGQT	63
NTDB id 32	MW RS07995 WP 000472256.1	MFKFLL..KK T QAFTLIEMLLVLLIISLLLILIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 16	SA RS07765 WP 000472256.1	MFKFLL..KK T QAFTLIEMLLVLLIISLLLILIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP	68
NTDB id 588	KW2 RS10845 WP 051303741.1	MSKALSLSVKIHGRKL C QKK K QAFTLIEMLIVLAIIISILLFVPNLIK E KAQVQKTGEAAVVKVVESQAQLYELDHDNDK	80
NTDB id 49768	SSUD12 RS00780 WP 002936550.1	MKKL V EMKV K GFTLVEMLVLVGIIISLLLLFVPNL S QKKDAV E KEGNAADVVKVVESQMELYELEHD.EE	68
NTDB id 460	GPW51 RS00790 WP 011921720.1	MKKL V EMKV K GFTLVEMLVLVGIIISLLLLFVPNL S QKKDA I TQKKGAADVVKVVESQMELYELEHD.EE	68
NTDB id 502	SM12261 RS00955 WP 000738641.1	MKKM M TF L KKAKVKAFTLVEMLVLLIISVLLLLFVPNLTKQKEAVNDKGKAADVVKVVESQAELYSLEKN.ED	72
NTDB id 162	SP RS10370 WP 000738627.1	MKKM M TF L KKAKVKAFTLVEMLVLLIISVFLFFVPNLTKQKEAVNDKGKAADVVKVVESQAELYSLEKN.ED	72
NTDB id 272	KZH43 RS09255 WP 000738626.1	MKKM M TF L KKAKVKAFTLVEMLVLLIISVFLFFVPNLTKQKEAVNDKGKAADVVKVVESQAELYSLEKN.ED	72
NTDB id 231	SPD RS09885 WP 000738626.1	MKKM M TF L KKAKVKAFTLVEMLVLLIISVFLFFVPNLTKQKEAVNDKGKAADVVKVVESQAELYSLEKN.ED	72
NTDB id 197	SPR RS09420 WP 000738626.1	MKKM M TF L KKAKVKAFTLVEMLVLLIISVFLFFVPNLTKQKEAVNDKGKAADVVKVVESQAELYSLEKN.ED	72
NTDB id 530	SMSK321 RS09230 WP 000738643.1	MKKM M TF L KKAKVKAFTLVEMLVLLIISVLLLLFVPNLTKQKEAVNDKGKAADVVKVVESQAELYSLEKN.ED	72
NTDB id 431	SGO RS09410 WP 012130940.1	MNKL K KL R VKAFTLVEMLVLLIISVLMFLFVPNLTKQKEAVSDTGNAADVVKVVESQAELYELKNTGDQ	69
NTDB id 396	FSA28 RS09395 WP 002263437.1	MKKI K RVSVRAFTLIEMLVLLIISILLFFVPNLTKQDKVKDTGGSAVVKVVESQAELYELNKTD.S	68
NTDB id 387	SMU RS09025 WP 002263437.1	MKKI K RVSVRAFTLIEMLVLLIISILLFFVPNLTKQDKVKDTGGSAVVKVVESQAELYELNKTD.S	68
consensus		* **K* *****! ! ! ! ! * ! ! ! ! ! * ! ! ! ! ! * ! ! ! ! ! * ! ! ! ! ! * ! ! ! ! ! * ! ! ! ! ! *	

NTDB	id	98	BSU	24710	NF	390351.1	PSLADLQSEGYVKKDAVCPNGK.RIIITGGEVKEH	98
NTDB	id	32	MW	RS07995	WP	000472256.1	SSIETDLIADGFIKEAQKTKCSGETITIS.NGEAVAN	103
NTDB	id	16	SA	RS07765	WP	000472256.1	SSIETDLIADGFIKEAQKTKCSGETITIS.NGEAVAN	103
NTDB	id	588	KW2	RS10845	WP	051303741.1	PNLSSELLSAGMITQKQVTAYDDYYDQNKNEQRNFDD	116
NTDB	id	49768	SSUD12	RS00780	WP	002936550.1	ATVEDLQQAGYITEKQAKQYATAKK.....	93
NTDB	id	460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGYITEKQAKQYATAKK.....	93
NTDB	id	502	SM12261	RS00955	WP	000738641.1	ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVD.	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	ATLSKLVAAGNISQKQADSYPKAYYGGKNNSETQAVAN	105
NTDB	id	396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDITQEQAANSYKDYAKHPSKTRHVAN	104
NTDB	id	387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDITQEQAANSYKDYAKHPSKTRHVAN	104
consensus							*** *!***! **** * * *	

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