

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	MSKALSLVKIHGRKLCQKKQKAFTLIEMLLVLAIIISILILLFVPNLIKKEKAQVKGTGEAAVVKVVESQAQLYELDHDNDK	80
NTDB id 460	GPW51 RS00790 WP 011921720.1	MKKLVEMKVKCFTLVEMLVVLGIISLLLLFVPNLSQKKDAIKKKGDAAVVKVVESQMELYELEHDEE	68
NTDB id 502	SM12261 RS00955 WP 000738641.1	MKKMTFLKKAKVKAFTLVEMLVLLIISVLLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED	72
NTDB id 272	KZH43 RS09255 WP 000738626.1	MKKMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED	72
NTDB id 231	SPD RS09885 WP 000738626.1	MKKMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED	72
NTDB id 197	SPR RS09420 WP 000738626.1	MKKMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED	72
NTDB id 162	SP RS10370 WP 000738627.1	MKKMTFLKKAKVKAFTLVEMLVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED	72
NTDB id 530	SMSK321 RS09230 WP 000738643.1	MKKMTFLKKAKVKAFTLVEMLVLLIISVLLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNED	72
NTDB id 431	SGO RS09410 WP 012130940.1	MNKLLKLRVKAFTLVEMLVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQ	69
NTDB id 206813	BSR14 RS09545 WP 037611125.1	MKKLNAVKLRAFTLIEMLVLLIISILLFLLFVPNLSKQKDSVKETGNAAVVKVVDQAELYEMKNNTKT	68
NTDB id 396	FSA28 RS09395 WP 002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILLFLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKTDS	68
NTDB id 387	SMU RS09025 WP 002263437.1	MKKIKRVSVRAFTLIEMLVLLIISILLFLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKTDS	68
consensus		*** ** ** ** **	68

NTDB id 32	MW	RS07995	WP	000472256.1	SSIEDLIADGF	IK	EAQK	TKCS	GETITI	.SNGEAV	AN	103
NTDB id 16	SA	RS07765	WP	000472256.1	SSIEDLIADGF	IK	EAQK	TKCS	GETITI	.SNGEAV	AN	103
NTDB id 588	KW2	RS10845	WP	051303741.1	PNLSELLSAGMIT	QK	QVTA	YDDY	YDQ	NKNEQ	RNFDD	116
NTDB id 460	GPW51	RS00790	WP	011921720.1	ATVADLQAKGYIT	EK	QAKQ	YATA	HK			93
NTDB id 502	SM12261	RS00955	WP	000738641.1	ASLSKLQADGRIT	E	EQA	KAYKE	YHAK	QNTS	QTVD	107
NTDB id 272	KZH43	RS09255	WP	000738626.1	ASLRKLQADGRIT	E	EQA	KAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 231	SPD	RS09885	WP	000738626.1	ASLRKLQADGRIT	E	EQA	KAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 197	SPR	RS09420	WP	000738626.1	ASLRKLQADGRIT	E	EQA	KAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 162	SP	RS10370	WP	000738627.1	ASLRKLQADGRIT	E	EQA	KAYKE	YHDK	KNGGAN	RKVND	108
NTDB id 530	SMSK321	RS09230	WP	000738643.1	ASLSKLQADGRIT	E	EQA	KAYNE	YAYK	KNGGAN	RKVND	108
NTDB id 431	SGO	RS09410	WP	012130940.1	ATLSKLVAAGNIS	QK	QADS	YKAY	YGGK	NNSET	QAVAN	105
NTDB id 206813	BSR14	RS09545	WP	037611125.1	ASLAALVSEGGIT	QK	QADS	YNDY	YAKH	GGES	RSVAN	104
NTDB id 396	FSA28	RS09395	WP	002263437.1	ATLSKLVASGDIT	Q	EQANS	YKDY	YAKH	PSKT	RHVAN	104
NTDB id 387	SMU	RS09025	WP	002263437.1	ATLSKLVASGDIT	Q	EQANS	YKDY	YAKH	PSKT	RHVAN	104
consensus					***	*	***	!	***	!	***	*

- X non conserved
- X similar
- X > 50% conserved