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NTDB id 996288 LLNCD0184 RS11610 WP 010906318.1	MERTKKFILKKIQKIFL	VKICEKKRQKELKAFTL	IEMLIIVLA	IISILILLFVPNLIKESQVQKTGEAAVVKVVESQA	80
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	VKIHGRLCQKKQKAFTL	IEMLIIVLA	IISILILLFVPNLIKESQVQKTGEAAVVKVVESQA	69
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEMKVGFTLVEMLVVLG	IISLILLFVPNL	SQQDAIQKGDAAVVKVVESQM	58
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKMMTFLLKKA	KVKAFTLVEMLVLLIISVLF	LLFVPNLTKQKEAVNDKGKAAVVKVVESQA	62
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKMMTFLLKKA	KVKAFTLVEMLVLLIISVLF	LLFVPNLTKQKEAVNDKGKAAVVKVVESQA	62
NTDB id 231 SPD RS09885 WP 000738626.1		MKKMMTFLLKKA	KVKAFTLVEMLVLLIISVLF	LLFVPNLTKQKEAVNDKGKAAVVKVVESQA	62
NTDB id 197 SPR RS09420 WP 000738626.1		MKKMMTFLLKKA	KVKAFTLVEMLVLLIISVLF	LLFVPNLTKQKEAVNDKGKAAVVKVVESQA	62
NTDB id 162 SP RS10370 WP 000738627.1		MKKMMTFLLKKA	KVKAFTLVEMLVLLIISVLF	LLFVPNLTKQKEAVNDKGKAAVVKVVESQA	62
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKMMTFLLKKA	KVKAFTLVEMLVLLIISVLF	LLFVPNLTKQKEAVNDKGKAAVVKVVESQA	62
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLKCLR	VKAFTLVEMLVLLIISVLM	LLFVPNLTKQKEAVSDTGNAAVVKVVESQA	58
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKKIKRVS	VRAFTL	IEMLVLLIISILLFVPNLTKQKDKVKDTGGS	58
NTDB id 387 SMU RS09025 WP 002263437.1		MKKIKRVS	VRAFTL	IEMLVLLIISILLFVPNLTKQKDKVKDTGGS	58
consensus		**	*	*****	*****

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NTDB id	Accession	Protein	Sequence	Position
NTDB id 996288	LLNCDO184	RS11610	WP 010906318.1	127
NTDB id 588	KW2	RS10845	WP 051303741.1	116
NTDB id 460	GPW51	RS00790	WP 011921720.1	93
NTDB id 502	SM12261	RS00955	WP 000738641.1	107
NTDB id 272	KZH43	RS09255	WP 000738626.1	108
NTDB id 231	SPD	RS09885	WP 000738626.1	108
NTDB id 197	SPR	RS09420	WP 000738626.1	108
NTDB id 162	SP	RS10370	WP 000738627.1	108
NTDB id 530	SMK321	RS09230	WP 000738643.1	108
NTDB id 431	SGO	RS09410	WP 012130940.1	105
NTDB id 396	FSA28	RS09395	WP 002263437.1	104
NTDB id 387	SMU	RS09025	WP 002263437.1	104
consensus				

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