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NTDB id 237529 CFA72 RS00665 WP 002986552.1	MINQWNNL	RHKKL	KGFTLL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	ETGNA	AAVVK	LVEN	QAELYEL	LSQG	.SK	72
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALSLV	KIHGRK	LCCKK	QKAFTL	IEML	IVLAI	ISIL	LLLFVPNL	LIKE	KAQV	QKTGE	AAVVKV	VESQA	QLYEL	DHDND	80
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEM	KVK	GFTL	VEML	VVLGI	ISL	LLLFVPNL	SQK	DAIQ	KKGDA	AAVVKV	VESQ	MELYEL	EH	68
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	LLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	72
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SL	72
NTDB id 231 SPD RS09885 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SL	72
NTDB id 197 SPR RS09420 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SL	72
NTDB id 162 SP RS10370 WP 000738627.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SL	72
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	LLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SL	72
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLK	KKL	R	VKAFTL	VEML	VLLII	ISVL	MLLFVPNL	TKQ	KEAV	SDTG	NA	AAVVKV	VESQAELYEL	69
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKK	IKR	VSV	RAFTL	IEML	VLLII	ISIL	LLLFVPNL	TKQ	KD	TGG	S	AVVKV	VESQAELYEL	68
NTDB id 387 SMU RS09025 WP 002263437.1		MKK	IKR	VSV	RAFTL	IEML	VLLII	ISIL	LLLFVPNL	TKQ	KD	TGG	S	AVVKV	VESQAELYEL	68
consensus	***	*	***	*****	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	***

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NTDB id 237529 CFA72 RS00665 WP 002986552.1	PSLSQLKADGS	ITEKQ	EKAYQ	DYYD	KHK	NEKAR	LSN	108
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELL	SAGMIT	QKQVT	AYDD	DYYD	QNK	NEQRNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQAK	QYATA	KK				93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQAKAYK	EYHAK	QNTS	QTVAD			107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KN	GGANR	KVND		108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KN	GGANR	KVND		108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHDK	KN	GGANR	KVND		108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYK	EYND	KN	GGANR	KVND		108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYN	EY	YAK	KN	GGANR	KVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGN	ISQKQAD	SYKAY	YG	KNN	SETQ	AVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQAN	SYKDY	YAK	HPSK	TRH	VAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQAN	SYKDY	YAK	HPSK	TRH	VAN	104
consensus	*****	!!!!	!!!!	!!!!	!!!!	****	***	

- X non conserved
- X similar
- X ≥ 50% conserved