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NTDB id 341200 ETT68 RS00670 WP 002986552.1		MINQWNN	LRHKKL	KGFTLL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	ETGNA	AAVVK	LVEN	QAELYEL	LSQG	.SK	72	
NTDB id 588 KW2 RS10845 WP 051303741.1		MSKALSLV	KIHGRK	LCQKK	QKAFTL	IEML	IVLAI	ISIL	LLLFVPNL	LIKE	KAQV	QKTGE	AAVVKV	VESQA	QLYEL	DHDNDK	80	
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEM	KVK	GFTL	VEML	VVLGI	ISL	LLLFVPNL	SQK	DAIQ	KKGDA	AAVVKV	VESQ	MELYEL	EHDE	.EE	68	
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	LLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	.ED	72	
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	.ED	72	
NTDB id 231 SPD RS09885 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	.ED	72	
NTDB id 197 SPR RS09420 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	.ED	72	
NTDB id 162 SP RS10370 WP 000738627.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	.ED	72	
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLII	ISVL	LLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN	.ED	72	
NTDB id 431 SGO RS09410 WP 012130940.1		MNKKL	KKL	R	VKAFTL	VEML	VLLII	ISVL	MLLFVPNL	TKQ	KEAV	SDTG	NA	AAVVKV	VESQAELYEL	KNTGDQ	69	
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKK	IKR	VSV	RAFTL	IEML	VLLII	ISIL	LLLFVPNL	TKQ	KD	TKV	KGDT	GG	SAVVKV	VESQAELYEL	NKT.DS	68
NTDB id 387 SMU RS09025 WP 002263437.1		MKK	IKR	VSV	RAFTL	IEML	VLLII	ISIL	LLLFVPNL	TKQ	KD	TKV	KGDT	GG	SAVVKV	VESQAELYEL	NKT.DS	68
consensus	***	*	***	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	***	**	

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NTDB id 341200 ETT68 RS00670 WP 002986552.1	PSLSQLKADGS	ITEKQEKAYQ	DYYD	KHKNEKARLSN	108		
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELLSAGM	ITCKQVTA	YDD	DYYDQNKNEQRNFDD	116		
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQAKQYATA	KK.....	93			
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQAKAYKE	YHAK	QNTS	QTVAD. 107		
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDK	KN	GGANR	KVND 108	
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDK	KN	GGANR	KVND 108	
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDK	KN	GGANR	KVND 108	
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYKE	YND	KN	GGANR	KVND 108	
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYNE	Y	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	HPSKT	RH	VAN 104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQAN	SYKDY	YAK	HPSKT	RH	VAN 104
consensus	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	****	*** *

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