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NTDB id 340979 ETT72 RS00685 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	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	EHDEE	68
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	LLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN.ED	72
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN.ED	72
NTDB id 231 SPD RS09885 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN.ED	72
NTDB id 197 SPR RS09420 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN.ED	72
NTDB id 162 SP RS10370 WP 000738627.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN.ED	72
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	LLLFVPNL	TKQ	KEAV	NDKG	KA	AAVVKV	VESQAELY	SLDKN.ED	72
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLKKL	R	VKAFTL	VEML	VLLI	ISVL	MLLFVPNL	TKQ	KEAV	SDTGNA	AAVVKV	VESQAELY	ELKNTG	DQ	69
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKK	IKRVS	VRAFTL	IEML	VLLI	ISIL	LLLFVPNL	TKQ	KDKV	KDTGG	S	AVVKV	VESQAELY	ELNKT.DS	68
NTDB id 387 SMU RS09025 WP 002263437.1		MKK	IKRVS	VRAFTL	IEML	VLLI	ISIL	LLLFVPNL	TKQ	KDKV	KDTGG	S	AVVKV	VESQAELY	ELNKT.DS	68
consensus	***	*	***	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	***

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NTDB id 340979 ETT72 RS00685 WP 002986552.1	PSLSQLKADGS	ITEKQEKAYQ	DYYD	KHKNEKAR	LSN	108
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELL	SAGMITQKQVTAYD	DYYDQ	NKNEQ	RNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQAKQYATAKK				93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQAKAYKE	YHAKQ	NTSQT	VAD.	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDKN	GGANR	KVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDKN	GGANR	KVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDKN	GGANR	KVND	108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYKE	YNDKN	GGANR	KVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYN	EYAKN	GGANR	KVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGN	ISQKQAD	SYKAYYG	KNNSET	QAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAKH	PSKTR	RHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAKH	PSKTR	RHVAN	104
consensus	*****	!!!!!!	!!!!!!	!!!!!!	****	***

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved