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NTDB id 342220 ETT49 RS00670 WP 002986552.1		MINQWNNL	RHKKL	KGFTL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	ETGNA	AAVVK	LVEN	QAELY	EL	SSQG	.SK	72				
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALSLV	KIHGRK	L	CQKK	QKAFTL	IEML	IVLAI	IS	IL	LLFVPNL	LIKE	KAQV	QKTG	EAAVVK	VV	ESQA	QLYEL	DHDNDK	80			
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEM	KVK	GFTL	VEML	VVL	GIIS	LLLLFVPNL	SQK	DAIQ	KKGD	AAVVK	VV	ESQM	ELYEL	EH	.EE	68				
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKMTF	L	KKA	KVKAFTL	VEML	VLLI	ISVL	LLFVPNL	TKQ	KEAV	NDKG	KA	AAVVK	VV	ESQA	ELYS	LDKN	.ED	72		
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKMTF	L	KKA	KVKAFTL	VEML	VLLI	ISVL	FLFVPNL	TKQ	KEAV	NDKG	KA	AAVVK	VV	ESQA	ELYS	LEKN	.ED	72		
NTDB id 231 SPD RS09885 WP 000738626.1		MKKMTF	L	KKA	KVKAFTL	VEML	VLLI	ISVL	FLFVPNL	TKQ	KEAV	NDKG	KA	AAVVK	VV	ESQA	ELYS	LEKN	.ED	72		
NTDB id 197 SPR RS09420 WP 000738626.1		MKKMTF	L	KKA	KVKAFTL	VEML	VLLI	ISVL	FLFVPNL	TKQ	KEAV	NDKG	KA	AAVVK	VV	ESQA	ELYS	LEKN	.ED	72		
NTDB id 162 SP RS10370 WP 000738627.1		MKKMTF	L	KKA	KVKAFTL	VEML	VLLI	ISVL	FLFVPNL	TKQ	KEAV	NDKG	KA	AAVVK	VV	ESQA	ELYS	LEKN	.ED	72		
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKMTF	L	KKA	KVKAFTL	VEML	VLLI	ISVL	LLFVPNL	TKQ	KEAV	NDKG	KA	AAVVK	VV	ESQA	ELYS	LDKN	.ED	72		
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLK	KL	R	VKAFTL	VEML	VLLI	ISVL	MLLFVPNL	TKQ	KEAV	SDTG	NA	AAVVK	VV	ESQA	EYEL	LKNTG	DQ	69		
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKK	I	KRV	SVRAFTL	IEML	VLLI	IS	IL	LLFVPNL	TKQ	KD	TKV	KDTG	G	SAVVK	VV	ESQA	EYEL	LNKT	.DS	68
NTDB id 387 SMU RS09025 WP 002263437.1		MKK	I	KRV	SVRAFTL	IEML	VLLI	IS	IL	LLFVPNL	TKQ	KD	TKV	KDTG	G	SAVVK	VV	ESQA	EYEL	LNKT	.DS	68
consensus		**	*	**	***	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	**	**

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NTDB id 342220 ETT49 RS00670 WP 002986552.1	PSLSQLKADGS	ITEKQ	EKAYQ	DYYD	KHKNE	KAR	L	SN	108
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELL	SAGMIT	QKQVT	AYDD	DYYDQ	NKNE	Q	RNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQ	AKQY	ATAK					93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQ	AKAYK	EYHAK	QNTS	Q	T	VAD.	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQ	AKAYK	EYHD	KNGG	AN	RK	VND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQ	AKAYK	EYHD	KNGG	AN	RK	VND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQ	AKAYK	EYHD	KNGG	AN	RK	VND	108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQ	AKAYK	EYND	KNGG	AN	RK	VND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQ	AKAYN	EY	YAKNGG	AN	RK	VND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGN	ISQKQ	ADS	SYKAY	YGKNN	SET	Q	AVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQ	ANSYK	DY	YAKHPSK	TRH	V	AN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQ	ANSYK	DY	YAKHPSK	TRH	V	AN	104
consensus	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	****	****	*

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