

logo

NTDB id 404833	GE024	RS00850	WP 003046186.1		MTNQWNNL	RHKKL	KGFTL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	ETGHT	AVVKL	VENQAELYEL	SSQG	.SK	72		
NTDB id 588	KW2	RS10845	WP 051303741.1	MSKALSLV	KIHGRK	LCQKK	QKAFTL	IEML	IVLAI	ISIL	LLLFVPNL	LIKE	KAQV	QKTGE	AAVVKV	VESQAQ	LYEL	DHDND	80		
NTDB id 460	GPW51	RS00790	WP 011921720.1		MKKLVEM	KVKG	GFTL	VEML	VVLGI	ISL	LLLFVPNL	SQQK	DAIQ	KKGDA	AVVKV	VESQMELYEL	EH	.EE	68		
NTDB id 502	SM12261	RS00955	WP 000738641.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	LLLFVPNL	TKQKEA	VNDKG	KA	AVVKV	VESQAELYSL	DKN	.ED	72		
NTDB id 272	KZH43	RS09255	WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKV	VESQAELYSL	EKN	.ED	72		
NTDB id 231	SPD	RS09885	WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKV	VESQAELYSL	EKN	.ED	72		
NTDB id 197	SPR	RS09420	WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKV	VESQAELYSL	EKN	.ED	72		
NTDB id 162	SP	RS10370	WP 000738627.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKV	VESQAELYSL	EKN	.ED	72		
NTDB id 530	SMSK321	RS09230	WP 000738643.1		MKKKMTF	LKKA	KVKAFTL	VEML	VLLI	ISVL	LLLFVPNL	TKQKEA	VNDKG	KA	AVVKV	VESQAELYSL	DKN	.ED	72		
NTDB id 431	SGO	RS09410	WP 012130940.1		MNKKLKKL	RVKAFTL	VEML	VLLI	ISVL	MLLFVPNL	TKQKEA	VS	DTGNA	AAVVKV	VESQAELYEL	KNTG	DQ	69			
NTDB id 396	FSA28	RS09395	WP 002263437.1		MKKIKR	VS	VRAFTL	IEML	VLLI	ISIL	LLLFVPNL	TKQK	DV	KD	TGGS	AVVKV	VESQAELYEL	NKT	.DS	68	
NTDB id 387	SMU	RS09025	WP 002263437.1		MKKIKR	VS	VRAFTL	IEML	VLLI	ISIL	LLLFVPNL	TKQK	DV	KD	TGGS	AVVKV	VESQAELYEL	NKT	.DS	68	
consensus					**	*	***	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	***	***

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NTDB id 404833	GE024	RS00850	WP 003046186.1	PSLSQLKEDGS	ITDKQEKAYRDYYG	KHKDEKAS	LND	108												
NTDB id 588	KW2	RS10845	WP 051303741.1	PNLSELLSAGM	ITQKQVTAYDDYDQ	NKNEQ	RNFDD	116												
NTDB id 460	GPW51	RS00790	WP 011921720.1	ATVADLQAKGY	ITEKQAKQYATAKK			93												
NTDB id 502	SM12261	RS00955	WP 000738641.1	ASLSKLQADGR	ITEEQAKAYKEYHAK	QNTS	QTVAD	107												
NTDB id 272	KZH43	RS09255	WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKEYHDK	KNGGAN	RKVND	108												
NTDB id 231	SPD	RS09885	WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKEYHDK	KNGGAN	RKVND	108												
NTDB id 197	SPR	RS09420	WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKEYHDK	KNGGAN	RKVND	108												
NTDB id 162	SP	RS10370	WP 000738627.1	ASLRKLQADGR	ITEEQAKAYKEYND	KNGGAN	RKVND	108												
NTDB id 530	SMSK321	RS09230	WP 000738643.1	ASLSKLQADGR	ITEEQAKAYNEYYA	KNGGAN	RKVND	108												
NTDB id 431	SGO	RS09410	WP 012130940.1	ATLSKLVAAGN	ISQKQADSYKAYYG	KNNSETQ	AVAN	105												
NTDB id 396	FSA28	RS09395	WP 002263437.1	ATLSKLVASGD	ITQEQANSYKDYYA	KHPSKTR	RHVAN	104												
NTDB id 387	SMU	RS09025	WP 002263437.1	ATLSKLVASGD	ITQEQANSYKDYYA	KHPSKTR	RHVAN	104												
consensus				*****	!!!!!!	!!!!!!	!!!!!!	****	***	***	***	***	***	***	***	***	***	***	***	***

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved