

logo

NTDB id 308557 DZS39 RS00670 WP 002986552.1		MINQWNNL	RHKKL	KGFTLL	LEMLLV	ILVISVL	MLLFVPNL	SKQK	DRVT	ETGNA	AAVVKLV	ENQAELYEL	SSQG	.SK	72	
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALSLV	KIHGRK	LCQKK	QKAFTL	IEMLIV	LAIIS	TLILLFVPNL	LIKEA	QVQKT	GEEAA	VVKVVSQA	QLYEL	DHDND	K	80	
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEM	KVK	GFTL	VEMLV	VLGIIS	LLLLFVPNL	SQQK	DAIQK	KGDA	AVVKVVSQ	MELYEL	EH	.EE	68	
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKKMTF	LKKA	KVKAFTL	VEMLV	VLLIISVL	LLLFVPNL	TKQKEA	VNDKG	KA	AVVKVVSQA	ELYSL	DKN	.ED	72	
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEMLV	VLLIISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKVVSQA	ELYSL	EKN	.ED	72	
NTDB id 231 SPD RS09885 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEMLV	VLLIISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKVVSQA	ELYSL	EKN	.ED	72	
NTDB id 197 SPR RS09420 WP 000738626.1		MKKKMTF	LKKA	KVKAFTL	VEMLV	VLLIISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKVVSQA	ELYSL	EKN	.ED	72	
NTDB id 162 SP RS10370 WP 000738627.1		MKKKMTF	LKKA	KVKAFTL	VEMLV	VLLIISVL	FLLFVPNL	TKQKEA	VNDKG	KA	AVVKVVSQA	ELYSL	EKN	.ED	72	
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKKMTF	LKKA	KVKAFTL	VEMLV	VLLIISVL	LLLFVPNL	TKQKEA	VNDKG	KA	AVVKVVSQA	ELYSL	DKN	.ED	72	
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLKKL	R	VKAFTL	VEMLV	VLLIISVL	MLLFVPNL	TKQKEA	VSDTG	NA	AAVVKVVSQA	ELYEL	KNTG	DQ	69	
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKK	IKRVS	VRAFTL	IEMLV	VLLIIS	TLILLFVPNL	TKQK	DVK	DTGG	S	AVVKVVSQA	ELYEL	NKT	.DS	68
NTDB id 387 SMU RS09025 WP 002263437.1		MKK	IKRVS	VRAFTL	IEMLV	VLLIIS	TLILLFVPNL	TKQK	DVK	DTGG	S	AVVKVVSQA	ELYEL	NKT	.DS	68
consensus		**	*	**	***	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	***	**

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NTDB id 308557 DZS39 RS00670 WP 002986552.1	PSLSQLKADGS	ITEKQEKAYQ	DYYD	KHKNEKAR	L	SN	108					
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELLSAGM	ITCKQVTA	YD	DYYDQ	NKNEQ	RNFDD	116					
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQAKQY	ATAKK				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	EYHD	KNGGAN	RK	VND	108					
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHD	KNGGAN	RK	VND	108					
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYK	EYHD	KNGGAN	RK	VND	108					
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYK	EYND	KNGGAN	RK	VND	108					
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYN	EY	YAK	KNGGAN	RK	VND	108				
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGN	ISQKQAD	SYKAYY	GKNN	SETQ	AVAN	105					
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAK	HPSKTR	RH	VAN	104					
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAK	HPSKTR	RH	VAN	104					
consensus	*****	!***!	!***!	***!	***!	***	***	*	***	*		

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved