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NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	LV	IKIHGRK	LCQKQKQ	KAFTLI	EMLV	LVLA	IISIL	ILLFVPNL	IK	EKAQV	QKT	G	EAAVVKV	VESQAQ	LYEL	DH	DN	KPNL	SE	85								
NTDB id 124404 GT95 RS01065 WP 000793380.1		MKN	L	LLKCK	DKK	VKAFTL	LEMLV	VLI	IISVLL	LLFVPNL	SK	QKES	V	TRTG	NAAVVKV	VESQAEL	FEL	QETGR	KASL	ST	78								
NTDB id 460 GPW51 RS00790 WP 011921720.1			MKK	L	VEMKVK	GFTL	VEMLV	VLG	IISLL	LLFVPNL	SQ	QKDA	I	QKK	G	DAVVKV	VESQM	EYEL	EH	EEAT	VAD	73							
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKK	M	TFLK	KA	KVKAFTL	VEMLV	VLLI	IISVLL	LLFVPNL	TK	QKEAV	N	DKG	K	AAVVKV	VESQAEL	YSL	DKN	EDAS	LSK	77							
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKK	M	TFLK	KA	KVKAFTL	VEMLV	VLLI	IISVLF	LLFVPNL	TK	QKEAV	N	DKG	K	AAVVKV	VESQAEL	YSL	EKN	EDAS	LRK	77							
NTDB id 231 SPD RS09885 WP 000738626.1		MKK	M	TFLK	KA	KVKAFTL	VEMLV	VLLI	IISVLF	LLFVPNL	TK	QKEAV	N	DKG	K	AAVVKV	VESQAEL	YSL	EKN	EDAS	LRK	77							
NTDB id 197 SPR RS09420 WP 000738626.1		MKK	M	TFLK	KA	KVKAFTL	VEMLV	VLLI	IISVLF	LLFVPNL	TK	QKEAV	N	DKG	K	AAVVKV	VESQAEL	YSL	EKN	EDAS	LRK	77							
NTDB id 162 SP RS10370 WP 000738627.1		MKK	M	TFLK	KA	KVKAFTL	VEMLV	VLLI	IISVLF	LLFVPNL	TK	QKEAV	N	DKG	K	AAVVKV	VESQAEL	YSL	EKN	EDAS	LRK	77							
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKK	M	TFLK	KA	KVKAFTL	VEMLV	VLLI	IISVLF	LLFVPNL	TK	QKEAV	N	DKG	K	AAVVKV	VESQAEL	YSL	DKN	EDAS	LSK	77							
NTDB id 431 SGO RS09410 WP 012130940.1			MNK	L	KKL	R	VKAFTL	VEMLV	VLLI	IISVLM	LLFVPNL	TK	QKEAV	S	D	TG	NAAVVKV	VESQAEL	YEL	KNTG	D	QATLSK	74						
NTDB id 396 FSA28 RS09395 WP 002263437.1			MKK	I	K	R	VSVRAFTL	I	EMLV	VLLI	IIS	I	LLFVPNL	TK	QK	D	K	V	K	D	T	G	S	AVVKV	VESQAEL	YEL	NKT	DSATLSK	73
NTDB id 387 SMU RS09025 WP 002263437.1			MKK	I	K	R	VSVRAFTL	I	EMLV	VLLI	IIS	I	LLFVPNL	TK	QK	D	K	V	K	D	T	G	S	AVVKV	VESQAEL	YEL	NKT	DSATLSK	73
consensus			**	*	*		***	*****	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	

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NTDB id 588 KW2 RS10845 WP 051303741.1	L	L	S	A	G	M	I	T	Q	K	Q	V	T	A	Y	D	D	Y	D	Q	N	K	N	E	Q	R	N	F	D	D	116	
NTDB id 124404 GT95 RS01065 WP 000793380.1	L	K	S	G	G	Y	I	T	E	K	Q	E	K	A	Y	L	D	Y	K	D	S	S	N	G	S	Q	K	I	S	S	109	
NTDB id 460 GPW51 RS00790 WP 011921720.1	L	Q	A	K	G	Y	I	T	E	K	Q	A	K	Q	Y	A	T	A	K											93		
NTDB id 502 SM12261 RS00955 WP 000738641.1	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	A	K	Q	N	T	S	Q	T	V	A	D	107	
NTDB id 272 KZH43 RS09255 WP 000738626.1	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	D	K	N	G	G	A	N	R	K	V	N	D	108
NTDB id 231 SPD RS09885 WP 000738626.1	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	D	K	N	G	G	A	N	R	K	V	N	D	108
NTDB id 197 SPR RS09420 WP 000738626.1	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	D	K	N	G	G	A	N	R	K	V	N	D	108
NTDB id 162 SP RS10370 WP 000738627.1	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	N	D	K	N	G	G	A	N	R	K	V	N	D	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	N	E	Y	Y	A	K	N	G	G	A	N	R	K	V	N	D	108
NTDB id 431 SGO RS09410 WP 012130940.1	L	V	A	A	G	N	I	S	Q	K	Q	A	D	S	Y	K	A	Y	Y	G	K	N	N	S	E	T	Q	A	V	A	N	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	L	V	A	S	G	D	I	T	Q	E	Q	A	N	S	Y	K	D	Y	Y	A	K	H	P	S	K	T	R	H	V	A	N	104
NTDB id 387 SMU RS09025 WP 002263437.1	L	V	A	S	G	D	I	T	Q	E	Q	A	N	S	Y	K	D	Y	Y	A	K	H	P	S	K	T	R	H	V	A	N	104
consensus	!	*	*	!	!	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****		

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