

NTDB id 1136042	DQL11 RS00670 WP 002986552.1		MINQWNN	RHKK	KGFTL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	TETG	NAAVVK	LVEN	QAELY	EL	SQG	.SK	72
NTDB id 588	KW2 RS10845 WP 051303741.1	MSKALSLV	KIHGRK	LCKK	QKQFTL	TEML	IVLA	IIISIL	ILLFVPNL	TK	EKAQV	QKTG	EAAVVK	VVVSQA	QLY	EL	DHND	.K	80
NTDB id 460	GPW51 RS00790 WP 011921720.1		MKKL	VEVMK	KGFTL	VEML	VVLG	GIISL	LLFVPNL	SSQK	DAIQ	KKGD	AAVVKV	VVSQ	MELY	EL	EHD	.EE	68
NTDB id 502	SM12261 RS00955 WP 000738641.1		MKKMM	TF	LKKA	KVKAFTL	VEML	VLLIIISVL	LLFVPNL	TKQ	KEAV	NDKG	AAVVKV	VVSQA	EYLS	L	EKN	.ED	72
NTDB id 272	KZH43 RS09255 WP 000738626.1		MKKMM	TF	LKKA	KVKAFTL	VEML	VLLIIISVL	LLFVPNL	TKQ	KEAV	NDKG	AAVVKV	VVSQA	EYLS	L	EKN	.ED	72
NTDB id 231	SPD RS09885 WP 000738626.1		MKKMM	TF	LKKA	KVKAFTL	VEML	VLLIIISVL	LLFVPNL	TKQ	KEAV	NDKG	AAVVKV	VVSQA	EYLS	L	EKN	.ED	72
NTDB id 197	SPR RS09420 WP 000738626.1		MKKMM	TF	LKKA	KVKAFTL	VEML	VLLIIISVL	LLFVPNL	TKQ	KEAV	NDKG	AAVVKV	VVSQA	EYLS	L	EKN	.ED	72
NTDB id 162	SP RS10370 WP 000738627.1		MKKMM	TF	LKKA	KVKAFTL	VEML	VLLIIISVL	LLFVPNL	TKQ	KEAV	NDKG	AAVVKV	VVSQA	EYLS	L	EKN	.ED	72
NTDB id 530	SMSK321 RS09230 WP 000738643.1		MKKMM	TF	LKKA	KVKAFTL	VEML	VLLIIISVL	LLFVPNL	TKQ	KEAV	NDKG	AAVVKV	VVSQA	EYLS	L	EKN	.ED	72
NTDB id 431	SGO RS09410 WP 012130940.1		MNK	LKKL	RVKAFTL	VEML	VLLIIISVL	MLLFVPNL	TKQ	KEAV	SDTG	NAAVVK	VVVSQA	EYLS	EL	KNTG	.D	69	
NTDB id 396	FSA28 RS09395 WP 002263437.1		MKK	L	KRVSV	RAFTL	TEML	VLLIIISL	LLFVPNL	TKQ	DKVD	TGGS	AAVVKV	VVSQA	EYLS	EL	NKT	.DS	68
NTDB id 387	SMU RS09025 WP 002263437.1		MKK	L	KRVSV	RAFTL	TEML	VLLIIISL	LLFVPNL	TKQ	DKVD	TGGS	AAVVKV	VVSQA	EYLS	EL	NKT	.DS	68
consensus		***	*	***	*****	***	*****	*	***	*****	***	*****	***	*****	*	***	*****		

NTDB id	1136042 DQL11 RS00670 WP 002986552.1	P S L S Q L K A D G S I T E K Q E K A Y Q D Y Y D K H K N E K A R L S N	108
NTDB id	588 KW2 RS10845 WP 051303741.1	P N L S E L L S A G M I T C K Q V T A Y D D Y Y D Q N K N E Q R N F D D	116
NTDB id	460 GPW51 RS00790 WP 011921720.1	A T V A D L Q A K G Y I T E K Q A K Q Y A T A K K	93
NTDB id	502 SM12261 RS00955 WP 000738641.1	A S L S K L Q A D G R I T E E Q A K A Y K E Y H A K N T S C T V A D .	107
NTDB id	272 KZH43 RS09255 WP 000738626.1	A S L R K 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	A S L R K 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	A S L R K 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	A S L R K 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	A S L S K 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	A T L S K 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	A T L S K 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	A T L S K 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