

NTDB id 1136646	DQL59 RS07405 WP 111681331.1		MINQWNN	RHKK	KGFTL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	TETG	NAAVVK	LVEN	QAELYE	LSQG	.SK	72	
NTDB id 588	KW2 RS10845 WP 051303741.1	MSKALSLV	KIHGRK	LCKK	QKQAF	TLIEM	LIVL	AIIS	ILILLFVPNL	TK	EKAQV	QKTG	EAAVVK	VVVSQA	QLYE	LHDN	.DK	80	
NTDB id 460	GPW51 RS00790 WP 011921720.1		MKKL	VEMKV	KGFTL	VEML	VVLG	GIIS	LILLFVPNL	SQLK	DAIQ	KKGD	AAVVKV	VVSQ	MELEYE	LHD	.EE	68	
NTDB id 502	SM12261 RS00955 WP 000738641.1		MKKMM	TFLLKKA	KVKAFTL	VEML	VLLIIS	SVLF	LFLFVPNL	TKQKEA	VNDK	GKAA	AVVKV	VVSQA	EYSL	EKN	.ED	72	
NTDB id 272	KZH43 RS09255 WP 000738626.1		MKKMM	TFLLKKA	KVKAFTL	VEML	VLLIIS	SVLF	LFLFVPNL	TKQKEA	VNDK	GKAA	AVVKV	VVSQA	EYSL	EKN	.ED	72	
NTDB id 231	SPD RS09885 WP 000738626.1		MKKMM	TFLLKKA	KVKAFTL	VEML	VLLIIS	SVLF	LFLFVPNL	TKQKEA	VNDK	GKAA	AVVKV	VVSQA	EYSL	EKN	.ED	72	
NTDB id 197	SPR RS09420 WP 000738626.1		MKKMM	TFLLKKA	KVKAFTL	VEML	VLLIIS	SVLF	LFLFVPNL	TKQKEA	VNDK	GKAA	AVVKV	VVSQA	EYSL	EKN	.ED	72	
NTDB id 162	SP RS10370 WP 000738627.1		MKKMM	TFLLKKA	KVKAFTL	VEML	VLLIIS	SVLF	LFLFVPNL	TKQKEA	VNDK	GKAA	AVVKV	VVSQA	EYSL	EKN	.ED	72	
NTDB id 530	SMSK321 RS09230 WP 000738643.1		MKKMM	TFLLKKA	KVKAFTL	VEML	VLLIIS	SVLF	LFLFVPNL	TKQKEA	VNDK	GKAA	AVVKV	VVSQA	EYSL	EKN	.ED	72	
NTDB id 431	SGO RS09410 WP 012130940.1		MNK	LKKL	RVKAFTL	VEML	VLLIIS	SVLM	LLFVPNL	TKQKEA	VS	DTG	NAAVVK	VVVSQA	EYEL	LKNTG	.DQ	69	
NTDB id 396	FSA28 RS09395 WP 002263437.1		MKK	LKRV	SVRAFTL	TEML	VLLIIS	TL	LLFVPNL	TKQK	DK	VD	TG	SAVVKV	VVSQA	EYEL	LKNT	.DS	68
NTDB id 387	SMU RS09025 WP 002263437.1		MKK	LKRV	SVRAFTL	TEML	VLLIIS	TL	LLFVPNL	TKQK	DK	VD	TG	SAVVKV	VVSQA	EYEL	LKNT	.DS	68
consensus			**	*	**	**	**	**	*	*	*	*	**	**	**	**	*		

NTDB id	1136646 DQL59 RS07405 WP 111681331.1	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 H 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 Q 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		* * * * * ! * * * ! * * * ! * * * ! * * * ! * * * ! * * *	

- ☐ non conserved
- ☒ similar
- ☒ >50% conserved