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NTDB id 98 BSU 24710 NP 390351.1	MNEKGF	TLVEM	LIVLFI	IISILL	LLITIP	NVTKHN	QTIQ	KKGC	EGLQ	NMVKA	QMTAF	ELDH	EGQT	PSIAD	68
NTDB id 619 LCA RS06470 WP 011374999.1	M	KKKR	KAFTL	TEMV	VVLAII	ALLILL	VAPNL	GAKK	QRAEK	KT	DQAL	VTTL	QTQ	VELAS	70
NTDB id 32 MW RS07995 WP 000472256.1	M	FKF..	LKKT	QAF	TLIEM	LVLLII	ISLLLI	LIIP	NTAK	QTAHI	QSTG	CNAQ	VKMV	NSQIE	73
NTDB id 16 SA RS07765 WP 000472256.1	M	FKF..	LKKT	QAF	TLIEM	LVLLII	ISLLLI	LIIP	NTAK	QTAHI	QSTG	CNAQ	VKMV	NSQIE	73
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	LVKIHGR	KLCQ	KQK	KAFTL	IEML	IVLAI	IISIL	ILLF	VPNLI	KEKA	QVQKT	GEAA	VVKV	85
NTDB id 46134 OG1RF RS08465 WP 002356991.1	M	KKKQ	KYAG	FTLLE	MLIV	LLIIS	VLIIL	LFVP	NLAK	HKHET	VDK	KGNEA	IVK	I	71
NTDB id 460 GPW51 RS00790 WP 011921720.1	M	KKLV	EMKVK	GFTL	VEMLV	VLGII	ISL	LLLL	LFVP	NLSQ	QKDAI	QKKG	DAAV	VVKV	73
NTDB id 502 SM12261 RS00955 WP 000738641.1	M	KKMT	FLKKA	KVKAFTL	VEMLV	VLLIIS	VLLLF	VFPNL	TKQKE	AVND	DKGKA	AVVKV	VESQA	EYSL	77
NTDB id 162 SP RS10370 WP 000738627.1	M	KKMT	FLKKA	KVKAFTL	VEMLV	VLLIIS	VLFLL	VFPNL	TKQKE	AVND	DKGKA	AVVKV	VESQA	EYSL	77
NTDB id 272 KZH43 RS09255 WP 000738626.1	M	KKMT	FLKKA	KVKAFTL	VEMLV	VLLIIS	VLFLL	VFPNL	TKQKE	AVND	DKGKA	AVVKV	VESQA	EYSL	77
NTDB id 231 SPD RS09885 WP 000738626.1	M	KKMT	FLKKA	KVKAFTL	VEMLV	VLLIIS	VLFLL	VFPNL	TKQKE	AVND	DKGKA	AVVKV	VESQA	EYSL	77
NTDB id 197 SPR RS09420 WP 000738626.1	M	KKMT	FLKKA	KVKAFTL	VEMLV	VLLIIS	VLFLL	VFPNL	TKQKE	AVND	DKGKA	AVVKV	VESQA	EYSL	77
NTDB id 530 SMSK321 RS09230 WP 000738643.1	M	KKMT	FLKKA	KVKAFTL	VEMLV	VLLIIS	VLLLF	VFPNL	TKQKE	AVND	DKGKA	AVVKV	VESQA	EYSL	77
NTDB id 431 SGO RS09410 WP 012130940.1	M	NKLK	KL	R	VKAFTL	VEMLV	VLLIIS	VLM	LLFVP	NLTK	QKEA	VSDT	GNA	AVVKV	74
NTDB id 396 FSA28 RS09395 WP 002263437.1	M	KKIK	RVSV	RAFTL	IEML	VLLIIS	ILL	LLFVP	NLTK	QKDK	VKDT	TGG	S	AVVKV	73
NTDB id 387 SMU RS09025 WP 002263437.1	M	KKIK	RVSV	RAFTL	IEML	VLLIIS	ILL	LLFVP	NLTK	QKDK	VKDT	TGG	S	AVVKV	73
consensus		*	***	*****	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	!!!!	*

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NTDB id 98 BSU 24710 NP 390351.1	LQSEGYVKKDAV.CPNGKRIIITGGEVKKVEH	98
NTDB id 619 LCA RS06470 WP 011374999.1	L.TDPKYISKDQLKHAKEKGITIEGGTVKKQK.	99
NTDB id 32 MW RS07995 WP 000472256.1	LIADGFIKEAQKTCKSGETITISNG.EAVAN	103
NTDB id 16 SA RS07765 WP 000472256.1	LIADGFIKEAQKTCKSGETITISNG.EAVAN	103
NTDB id 588 KW2 RS10845 WP 051303741.1	LLSAGMITCKQVTA YDDYDQNKNEQRNFDD	116
NTDB id 46134 OG1RF RS08465 WP 002356991.1	LVN EGYITKEQLDKYTAEKQ.....	91
NTDB id 460 GPW51 RS00790 WP 011921720.1	LQAKGYITEKQAKQYATAKK.....	93
NTDB id 502 SM12261 RS00955 WP 000738641.1	LQADGRITEEQAKAYKEYHAKQNTSQTVAD.	107
NTDB id 162 SP RS10370 WP 000738627.1	LQADGRITEEQAKAYKEYNDKNGGANRKYVND	108
NTDB id 272 KZH43 RS09255 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKYVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKYVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKYVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	LQADGRITEEQAKAYNEYAKNGGANRKYVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	LVAAGNISQKQADSYYKYYGKNNSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	LVASGDITQEANSYKDYAKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	LVASGDITQEANSYKDYAKHPSKTRHVAN	104
consensus	! * * * * * * * * * * * * * * *	

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