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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	KKRKA	FTLI	EMV	VVLAII	ALLILL	VAPNL	GAKK	QRAE	KKTD	QALV	TTLQT	QVEL	ASDE	70
NTDB id 32 MW RS07995 WP 000472256.1	MFKF	..L	KKTQA	FTLI	EMLV	LLIIS	LLLLI	ITPNI	AKQTA	HIQ	STGC	NAQV	KMVNS	QIEA	73
NTDB id 16 SA RS07765 WP 000472256.1	MFKF	..L	KKTQA	FTLI	EMLV	LLIIS	LLLLI	ITPNI	AKQTA	HIQ	STGC	NAQV	KMVNS	QIEA	73
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	LVKIH	GRKLC	QKKQ	KAF	TLI	EMLV	LAII	SILILL	FFV	PNLI	KEKA	QVQKT	GEAA	85
NTDB id 99690 QUD94 RS08050 WP 002356991.1	M	KKKQK	YAGF	TLLE	MLIV	LLIIS	VLILL	FFV	PNLA	AKHK	ETV	DKK	GNEA	IVK	71
NTDB id 460 GPW51 RS00790 WP 011921720.1	M	KKLV	EMK	VKG	FTL	VEM	LVVL	GIIS	LLLL	FFV	PNLS	SQK	DAIQ	KKG	73
NTDB id 502 SM12261 RS00955 WP 000738641.1	M	KKMT	FLK	KA	VKA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	TKQ	KEA	77
NTDB id 162 SP RS10370 WP 000738627.1	M	KKMT	FLK	KA	VKA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	TKQ	KEA	77
NTDB id 272 KZH43 RS09255 WP 000738626.1	M	KKMT	FLK	KA	VKA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	TKQ	KEA	77
NTDB id 231 SPD RS09885 WP 000738626.1	M	KKMT	FLK	KA	VKA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	TKQ	KEA	77
NTDB id 197 SPR RS09420 WP 000738626.1	M	KKMT	FLK	KA	VKA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	TKQ	KEA	77
NTDB id 530 SMSK321 RS09230 WP 000738643.1	M	KKMT	FLK	KA	VKA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	TKQ	KEA	77
NTDB id 431 SGO RS09410 WP 012130940.1	M	NK	LK	KL	R	V	KA	FTL	VEM	LVLL	IISV	LLFF	V	PNLT	74
NTDB id 396 FSA28 RS09395 WP 002263437.1	M	KK	I	K	R	V	S	V	R	A	F	T	L	I	73
NTDB id 387 SMU RS09025 WP 002263437.1	M	KK	I	K	R	V	S	V	R	A	F	T	L	I	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.TDPKYISKDQLKHAKKEKGITIEGGTVKQK.	99
NTDB id 32 MW RS07995 WP 000472256.1	LIADGFIKEAQKCTCKSGETITISNG.EAVAN	103
NTDB id 16 SA RS07765 WP 000472256.1	LIADGFIKEAQKCTCKSGETITISNG.EAVAN	103
NTDB id 588 KW2 RS10845 WP 051303741.1	LLSAGMITCKQVTAYPDLYDQNKNEQRNFDD	116
NTDB id 99690 QUD94 RS08050 WP 002356991.1	LVNEGYYITKEQLDKYTAEKQ.....	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	LQADGRITEEQAKAYKEYNDKNGGANRKYND	108
NTDB id 272 KZH43 RS09255 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKYND	108
NTDB id 231 SPD RS09885 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKYND	108
NTDB id 197 SPR RS09420 WP 000738626.1	LQADGRITEEQAKAYKEYHDKNGGANRKYND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	LQADGRITEEQAKAYNEYAKNGGANRKYND	108
NTDB id 431 SGO RS09410 WP 012130940.1	LVAAGNISQKQADSYKAYYGKNNSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	LVASGDITQEQANSYKDYAKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	LVASGDITQEQANSYKDYAKHPSKTRHVAN	104
consensus	! *** ***** ***** * * * * *	

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