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NTDB id 133737 QI18 RS11685 WP 023188581.1	MLIVLAIISILILLFVPNL	IKESQVQKTGEAAVVKVVESQAQLYEL	DHDD	EKPSLSE	58									
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	VKIHGRKLCQKQKKAFTL	IEMLIVLAIISILILLFVPNL	IKESQVQKTGEAAVVKVVESQAQLYEL	DHDD	EKPNLSE	85							
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEMKVKGFTL	VEMLVLVGIISL	LLFVPNL	SQQKDAIQKKGDA	AVVKVVESQ	MELYEL	HD	EEATVAD	73				
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKMMTFLKKA	KVKAFTL	VEMLVLLIISV	LLFVPNL	TKQKEAV	NDKGKAA	VVKVVESQA	ELYSL	DKN	EDASLSK	77		
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKMMTFLKKA	KVKAFTL	VEMLVLLIISV	LLFVPNL	TKQKEAV	NDKGKAA	VVKVVESQA	ELYSL	EKN	EDASLRK	77		
NTDB id 231 SPD RS09885 WP 000738626.1		MKKMMTFLKKA	KVKAFTL	VEMLVLLIISV	LLFVPNL	TKQKEAV	NDKGKAA	VVKVVESQA	ELYSL	EKN	EDASLRK	77		
NTDB id 197 SPR RS09420 WP 000738626.1		MKKMMTFLKKA	KVKAFTL	VEMLVLLIISV	LLFVPNL	TKQKEAV	NDKGKAA	VVKVVESQA	ELYSL	EKN	EDASLRK	77		
NTDB id 162 SP RS10370 WP 000738627.1		MKKMMTFLKKA	KVKAFTL	VEMLVLLIISV	LLFVPNL	TKQKEAV	NDKGKAA	VVKVVESQA	ELYSL	EKN	EDASLRK	77		
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKMMTFLKKA	KVKAFTL	VEMLVLLIISV	LLFVPNL	TKQKEAV	NDKGKAA	VVKVVESQA	ELYSL	DKN	EDASLSK	77		
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLK	KL	RVKAFTL	VEMLVLLIISV	LMLFVPNL	TKQKEAV	SDTGNA	AVVKVVESQA	ELYEL	KNTG	DQATLSK	74	
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKKIK	RVSVRAFTL	IEMLVLLIIS	ILL	LLFVPNL	TKQKD	KVKDTGG	SAVVKVVESQA	ELYEL	NKT	DSATLSK	73	
NTDB id 387 SMU RS09025 WP 002263437.1		MKKIK	RVSVRAFTL	IEMLVLLIIS	ILL	LLFVPNL	TKQKD	KVKDTGG	SAVVKVVESQA	ELYEL	NKT	DSATLSK	73	
consensus		**	*	***	*****	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!

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NTDB id 133737 QI18 RS11685 WP 023188581.1	LLNAGMITQKQISAYDNYDQNKNEERNFND	89
NTDB id 588 KW2 RS10845 WP 051303741.1	LLSAGMITQKQVTAAYDDYDQNKNEQRNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	LQAKGYITEKQAKQYATAKK.....	93
NTDB id 502 SM12261 RS00955 WP 000738641.1	LQADGRITEEQAKAYKEEYHAKQNTSQTVAD.	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	LQADGRITEEQAKAYKEEYHDKNGGANRNVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	LQADGRITEEQAKAYKEEYHDKNGGANRNVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	LQADGRITEEQAKAYKEEYHDKNGGANRNVND	108
NTDB id 162 SP RS10370 WP 000738627.1	LQADGRITEEQAKAYKEEYNDKNGGANRNVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	LQADGRITEEQAKAYNEEYAKNGGANRNVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	LVAAGNISQKQADSYKAYYGKNNSETQAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	LVASGDITQEQANSYKDYYAKHPSKTRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	LVASGDITQEQANSYKDYYAKHPSKTRHVAN	104
consensus	!***! !***!***!*****	

- non conserved
- similar
- ≥ 50% conserved