

logo

NTDB id 183097 LLUC06 RS12070 WP 012898622.1	MERTKKFILKKIQKIFLFV	KICEKKCKQKELRAFTL	IEMLVLA	IISL	LLFVPNL	IKESQV	QKTGE	AAVVKV	VESQA	80
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	LVIHGRKLCQKQKQ	KAFTL	IEMLVLA	IISL	LLFVPNL	IKESQV	QKTGE	AAVVKV	69
NTDB id 460 GPW51 RS00790 WP 011921720.1		MKKLVEMKVKGFTL	VEMLVVL	GIISL	LLFVPNL	SQKQ	DAIQK	KGD	AAVVKV	58
NTDB id 502 SM12261 RS00955 WP 000738641.1		MKKMTFLKKA	KVKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	NDKCK	AAVVKV	62
NTDB id 272 KZH43 RS09255 WP 000738626.1		MKKMTFLKKA	KVKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	NDKCK	AAVVKV	62
NTDB id 231 SPD RS09885 WP 000738626.1		MKKMTFLKKA	KVKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	NDKCK	AAVVKV	62
NTDB id 197 SPR RS09420 WP 000738626.1		MKKMTFLKKA	KVKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	NDKCK	AAVVKV	62
NTDB id 162 SP RS10370 WP 000738627.1		MKKMTFLKKA	KVKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	NDKCK	AAVVKV	62
NTDB id 530 SMSK321 RS09230 WP 000738643.1		MKKMTFLKKA	KVKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	NDKCK	AAVVKV	62
NTDB id 431 SGO RS09410 WP 012130940.1		MNKLKKLR	VKAFTL	VEMLVVL	LIISVL	LLFVPNL	TKQKEAV	SDTCN	AAVVKV	58
NTDB id 396 FSA28 RS09395 WP 002263437.1		MKKTKRVS	VRFTL	IEMLVVL	LIISL	LLFVPNL	TKQK	DKVKDT	GSAVVKV	58
NTDB id 387 SMU RS09025 WP 002263437.1		MKKTKRVS	VRFTL	IEMLVVL	LIISL	LLFVPNL	TKQK	DKVKDT	GSAVVKV	58
consensus		**	*	***	*****	!!!!	*!!!!	*!!!!	*!!!!	*

logo

NTDB id 183097 LLUC06 RS12070 WP 012898622.1	QLYELDHDD	EKPSLSELLS	SAGMITQKQIS	AYDNYYDQ	NKNEERN	NFND	127
NTDB id 588 KW2 RS10845 WP 051303741.1	QLYELDHDD	NKPNSELLS	SAGMITQKQVT	AYDDYYDQ	NKNEQR	NFND	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ELYELEHD	EEATVADL	QAKGYITE	KQAKQYATA	KK.....		93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ELYSLDKN	EDASLSKLQ	ADGRITEEQAK	AYKEYHAK	QNTSQT	VD	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ELYSLEKN	EDASLRKLQ	ADGRITEEQAK	AYKEYHDK	KNGGAN	NRKVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ELYSLEKN	EDASLRKLQ	ADGRITEEQAK	AYKEYHDK	KNGGAN	NRKVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ELYSLEKN	EDASLRKLQ	ADGRITEEQAK	AYKEYHDK	KNGGAN	NRKVND	108
NTDB id 162 SP RS10370 WP 000738627.1	ELYSLEKN	EDASLRKLQ	ADGRITEEQAK	AYKEYYND	KNGGAN	NRKVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ELYSLDKN	EDASLSKLQ	ADGRITEEQAK	AYNEYYA	KNGGAN	NRKVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ELYELKNTG	DQATLSKL	VAAAGNISQK	QADSYKAY	YGKMNSET	QAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ELYELNKT	DSATLSKL	VASGDITQE	QANSYKDY	YAKHPSK	TRHVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ELYELNKT	DSATLSKL	VASGDITQE	QANSYKDY	YAKHPSK	TRHVAN	104
consensus	***	!*	*****	*****	*****	**	* * *

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved