

logo

NTDB id 1036735 ACB350 RS00675 WP 002986552.1		MINQWNN	RHKKL	KGFTL	LEML	LVIL	VISVL	MLLFVPNL	SKQK	DRVT	ETG	NAAVVK	LVEN	QAELYEL	SKG	72					
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	SLV	KIHGRK	LCQKK	QKAF	TL	IEML	IVLA	IISIL	ILLFVPNL	IK	KAQV	QKT	GEA	AVVKV	VESQA	QLYEL	DH	DN	DK	80
NTDB id 460 GPW51 RS00790 WP 011921720.1																					68
NTDB id 502 SM12261 RS00955 WP 000738641.1																					72
NTDB id 272 KZH43 RS09255 WP 000738626.1																					72
NTDB id 231 SPD RS09885 WP 000738626.1																					72
NTDB id 197 SPR RS09420 WP 000738626.1																					72
NTDB id 162 SP RS10370 WP 000738627.1																					72
NTDB id 530 SMSK321 RS09230 WP 000738643.1																					72
NTDB id 431 SGO RS09410 WP 012130940.1																					69
NTDB id 396 FSA28 RS09395 WP 002263437.1																					68
NTDB id 387 SMU RS09025 WP 002263437.1																					68
consensus			**	*	**	*	**	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	

logo

NTDB id 1036735 ACB350 RS00675 WP 002986552.1		PSLSQL	KADGS	ITEK	QEKAY	QDYD	KHKNEKAR	LSN	108
NTDB id 588 KW2 RS10845 WP 051303741.1		PNLSELL	SAGMIT	QKQVT	AYDD	YDQ	NKNEQ	RNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1		ATVADL	QAKGY	ITEK	QAKQY	ATAKK			93
NTDB id 502 SM12261 RS00955 WP 000738641.1		ASLSKL	QADGR	ITEEQ	AKAYK	EYHAK	QNTS	QTVAD	107
NTDB id 272 KZH43 RS09255 WP 000738626.1		ASLRKL	QADGR	ITEEQ	AKAYK	EYHD	KNGGAN	RKVND	108
NTDB id 231 SPD RS09885 WP 000738626.1		ASLRKL	QADGR	ITEEQ	AKAYK	EYHD	KNGGAN	RKVND	108
NTDB id 197 SPR RS09420 WP 000738626.1		ASLRKL	QADGR	ITEEQ	AKAYK	EYHD	KNGGAN	RKVND	108
NTDB id 162 SP RS10370 WP 000738627.1		ASLRKL	QADGR	ITEEQ	AKAYK	EYND	KNGGAN	RKVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1		ASLSKL	QADGR	ITEEQ	AKAYN	EYYAK	KNGGAN	RKVND	108
NTDB id 431 SGO RS09410 WP 012130940.1		ATLSKL	VAAAGNI	SQKQAD	SYKAYY	GKNNSET	QAVAN		105
NTDB id 396 FSA28 RS09395 WP 002263437.1		ATLSKL	VASGD	ITQEQAN	SYKDYY	AKHPSKTR	RVAN		104
NTDB id 387 SMU RS09025 WP 002263437.1		ATLSKL	VASGD	ITQEQAN	SYKDYY	AKHPSKTR	RVAN		104
consensus									

- X non conserved
- X similar
- X ≥ 50% conserved