

logo

NTDB id 1038058 ACB338 RS00665 WP 002986552.1		MINQWNN	RHKKL	KGFTLL	EMLLV	ILVISVL	MLLFVPNL	SKQKDRVT	ETGNA	AAVVKL	VENQA	ELYEL	SLSG	.SK	72	
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	SLV	KIHGRK	LCQKK	QKKAFTL	IEMLIV	LAIIIS	ILILLFVPNL	IKKKAQV	QKTGE	AAVVKV	VESQA	QLYEL	DHDN	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			**	*	**	**	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!

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NTDB id 1038058 ACB338 RS00665 WP 002986552.1	PSLSQLKADGS	ITEKQEKAYQ	DYYDK	HKNEKAR	LSN	108
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELLS	SAGMITQ	KQVTAYD	DYYDQ	NKNEQ	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQAKQY	ATAKK			93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQAKAYKE	YHAKQ	NTSQT	VAD	107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDKN	GGANR	KVND	108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDKN	GGANR	KVND	108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDKN	GGANR	KVND	108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYKE	YNDKN	GGANR	KVND	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYNE	YYAKN	GGANR	KVND	108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNI	SQKQADS	SYKAYYG	KNNSET	QAVAN	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQANSY	KDYAK	HPSKTR	RVAN	104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQANSY	KDYAK	HPSKTR	RVAN	104
consensus						

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