

logo

NTDB id 1037071 ACB340 RS00660 WP 002986552.1		MINQWNNL	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	IVL	AIIS	IL	ILLFVPNL	IK	EKAQV	QKT	GEA	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 1037071 ACB340 RS00660 WP 002986552.1	PSLSQLKADGS	ITEKQ	EKAYQ	DYD	KHKNEKAR	LSN	108
NTDB id 588 KW2 RS10845 WP 051303741.1	PNLSELL	SAGMIT	QKQVT	AYDD	YDQ	NKNEQRNFDD	116
NTDB id 460 GPW51 RS00790 WP 011921720.1	ATVADLQAKGY	ITEKQAKQY	ATAKK				93
NTDB id 502 SM12261 RS00955 WP 000738641.1	ASLSKLQADGR	ITEEQAKAYKE	YHAK	QNTS	QTVAD		107
NTDB id 272 KZH43 RS09255 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDK	KNGGANR	KVND		108
NTDB id 231 SPD RS09885 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDK	KNGGANR	KVND		108
NTDB id 197 SPR RS09420 WP 000738626.1	ASLRKLQADGR	ITEEQAKAYKE	YHDK	KNGGANR	KVND		108
NTDB id 162 SP RS10370 WP 000738627.1	ASLRKLQADGR	ITEEQAKAYKE	YND	KNGGANR	KVND		108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	ASLSKLQADGR	ITEEQAKAYNE	YYAK	KNGGANR	KVND		108
NTDB id 431 SGO RS09410 WP 012130940.1	ATLSKLVAAGNI	SQKQADS	SYKAYYG	KNNSETQ	AVAN		105
NTDB id 396 FSA28 RS09395 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAK	HPSKTRH	VAN		104
NTDB id 387 SMU RS09025 WP 002263437.1	ATLSKLVASGD	ITQEQANSYK	DYYAK	HPSKTRH	VAN		104
consensus							

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