

[illegible]

NTDB id	498	SM12261	RS01980	WP	000282464.1	NGQMN YVTEVL V TG FQLLESRAQRAMRENNAGQD.....LADLVL	124
NTDB id	526	SMSK321	RS08825	WP	000282440.1	NGQMN YVAEVL V TG FQLLESRAQRAMRENNAGQD.....LADLVL	124
NTDB id	158	SP	RS09595	WP	000282442.1	NGQMN YVTEVL V TG FQLLESRAQRAMRENNAGQD.....LADLVL	124
NTDB id	268	KZH43	RS08520	WP	000282467.1	NGQMN YVTEVL V TG FQLLESRAQRAMRENNAGQD.....LADLVL	124
NTDB id	227	SPD	RS09110	WP	000282467.1	NGQMN YVTEVL V TG FQLLESRAQRAMRENNAGQD.....LADLVL	124
NTDB id	193	SPR	RS08685	WP	000282467.1	NGQMN YVTEVL V TG FQLLESRAQRAMRENNAGQD.....LADLVL	124
NTDB id	583	DLJ51	RS09560	WP	002962590.1	DGGTH YVTEI L AQS FQLLESRAQRLRENNGASD.....LTDLVL	124
NTDB id	409	SMU	RS08940	WP	002263454.1	DGVTH FVTEVL GHSFQLLESRAQRAMRENNSEND.....LADLVL	124
NTDB id	624	LCA	RS00040	WP	011373726.1	QGGRV YVTEVV VDN FSLL ESRTTTEQR QG DGASQNFNNSNQSGS.....QQSGFTSPQQTGNAPAAANNTQADPFANNGQAID	163
NTDB id	112	BSU	36310	NP	391512.2	EGVN YVTEVL ADTVRFMDPKPREKAAD.....YHQQRQT.....QTGNNPFDN...TTAT	113
NTDB id	273769	RLO6	RS11600	WP	000934385.1	VGQRV FVTEV ADSV QFL E PKNN NQPPMNN.....YHQQRQT.....QTGNNPFDN...TTAT	135
NTDB id	113	BSU	40900	NP	391970.1	QGGRV FVTEV QAESV QFL E PK NGGC SGSGGYNENSGGGGYFGGGQNDNPFGGNQNNQRRNQGSNFDDPFANDGKPID	165
consensus						*!* *!***** ***** ***** *	

NTDB id 498 SM12261 RS01980 WP 000282464.1	EEEELPF	131
NTDB id 526 SMSK321 RS08825 WP 000282440.1	EEEELPF	131
NTDB id 158 SP RS09595 WP 000282442.1	EEEELPF	131
NTDB id 268 KZH43 RS08520 WP 000282467.1	EEEELPF	131
NTDB id 227 SPD RS09110 WP 000282467.1	EEEELPF	131
NTDB id 193 SPR RS08685 WP 000282467.1	EEEELPF	131
NTDB id 583 DLJ51 RS09560 WP 002962590.1	EEEELPF	131
NTDB id 409 SMU RS08940 WP 002263454.1	EEEELPF	131
NTDB id 624 LCA RS00040 WP 011373726.1	SDDDLPF	170
NTDB id 112 BSU 36310 NP 391512.2		113
NTDB id 273769 RL06 RS11600 WP 000934385.1	TDDDLPF	142
NTDB id 113 BSU 40900 NP 391970.1	SDDDLPF	172
consensus	*****	

- X non conserved
- X similar
- X >50% conserved