

logo

NTDB id 619 LCA RS06470 WP 011374999.1
NTDB id 32 MW RS07995 WP 000472256.1
NTDB id 16 SA RS07765 WP 000472256.1
NTDB id 588 KW2 RS10845 WP 051303741.1
NTDB id 460 GPW51 RS00790 WP 011921720.1
NTDB id 502 SM12261 RS00955 WP 000738641.1
NTDB id 530 SMSK321 RS09230 WP 000738643.1
NTDB id 98540 R8611 RS09590 WP 317647938.1
NTDB id 162 SP RS10370 WP 000738627.1
NTDB id 272 KZH43 RS09255 WP 000738626.1
NTDB id 231 SPD RS09885 WP 000738626.1
NTDB id 197 SPR RS09420 WP 000738626.1
NTDB id 431 SGO RS09410 WP 012130940.1
NTDB id 396 FSA28 RS09395 WP 002263437.1
NTDB id 387 SMU RS09025 WP 002263437.1
consensus

[illegible]

logo

```

NTDB id 619 LCA RS06470 WP 011374999.1
NTDB id 32 MW RS07995 WP 000472256.1
NTDB id 16 SA RS07765 WP 000472256.1
NTDB id 588 KW2 RS10845 WP 051303741.1
NTDB id 460 GPW51 RS00790 WP 011921720.1
NTDB id 502 SM12261 RS00955 WP 000738641.1
NTDB id 530 SMSK321 RS09230 WP 000738643.1
NTDB id 98540 R8611 RS09590 WP 317647938.1
NTDB id 162 SP RS10370 WP 000738627.1
NTDB id 272 KZH43 RS09255 WP 000738626.1
NTDB id 231 SPD RS09885 WP 000738626.1
NTDB id 197 SPR RS09420 WP 000738626.1
NTDB id 431 SGO RS09410 WP 012130940.1
NTDB id 396 FSA28 RS09395 WP 002263437.1
NTDB id 387 SMU RS09025 WP 002263437.1
consensus

```

QAD	TR	TEE	AKA	YKEY	PKN	GGAN	NRK	VND	
L	TDKY	TSKD	QLK	HAK	EKG	ITIE	GGTV	KQK	99
LI	ADFI	KEBA	KTCK	SGET	ITIS	.NGE	AVAN		103
LI	ADFI	KEBA	KTCK	SGET	ITIS	.NGE	AVAN		103
LL	SAGM	ITGK	QVTAY	DDYD	QNK	NEQR	RFDD		116
LQ	AKCY	ITEB	QAKQ	YATA	KK				93
LQ	ADGR	ITEE	QAKA	YKEY	HAKQ	NTSQT	VAD		107
LQ	ADGR	ITEE	QAKA	YNEE	YAKN	GGAN	RRKVND		108
LK	DDGR	ITEE	QVKA	YKEH	NDK	GGAN	RRKVND		108
LQ	ADGR	ITEE	QAKA	YKEY	NDK	GGAN	RRKVND		108
LQ	ADGR	ITEE	QAKA	YKEY	HDK	GGAN	RRKVND		108
LQ	ADGR	ITEE	QAKA	YKEY	HDK	GGAN	RRKVND		108
LQ	ADGR	ITEE	QAKA	YKEY	HDK	GGAN	RRKVND		108
LQ	ADGR	ITEE	QAKA	YKEY	HDK	GGAN	RRKVND		108
LV	AAGN	ISGK	QADS	YKAY	YGKN	NNSET	QAVAN		105
LV	ASGD	ITGE	QANS	YKDY	YAKH	PSKTR	RRHVAN		104
LV	ASGD	ITGE	QANS	YKDY	YAKH	PSKTR	RRHVAN		104

-  non conserved
-  similar
-  >50% conserved