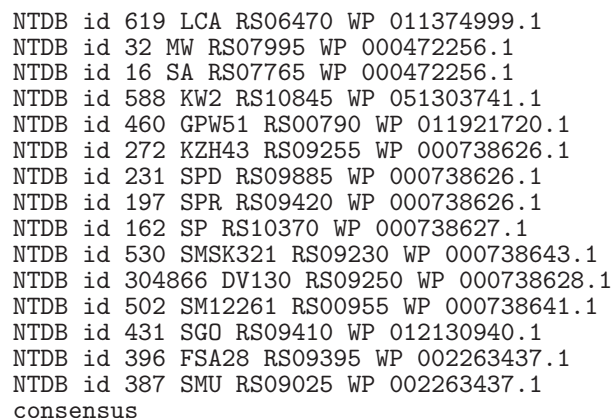


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NTDB id 619 LCA RS06470 WP 011374999.1
NTDB id 32 MW RS07995 WP 000472256.1
NTDB id 16 SA RS07765 WP 000472256.1
NTDB id 588 KW2 RS10845 WP 051303741.1
NTDB id 460 GPW51 RS00790 WP 011921720.1
NTDB id 272 KZH43 RS09255 WP 000738626.1
NTDB id 231 SPD RS09885 WP 000738626.1
NTDB id 197 SPR RS09420 WP 000738626.1
NTDB id 162 SP RS10370 WP 000738627.1
NTDB id 530 SMSK321 RS09230 WP 000738643.1
NTDB id 304866 DV130 RS09250 WP 000738628.1
NTDB id 502 SM12261 RS00955 WP 000738641.1
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

```



ASL	SK	QAD	R	TEE	QAK	YKEY	YK	KN	GGAN	RK	VND				
KSL	DEL	TD	KYT	ISK	DQL	KH	AKE	GK	GIT	IE	GGT	TV	KQK		
SS	IED	LI	AD	GF	I	KEA	QK	TS	G	ET	IT	IS	NGEAVAN		
SS	IED	LI	AD	GF	I	KEA	QK	TS	G	ET	IT	IS	NGEAVAN		
PN	LS	ELL	S	AG	MIT	CT	KQ	VT	AY	DD	Y	YD	QNKNEQRNFDD		
AT	VAD	L	QAK	G	Y	I	TE	KQAK	Q	Y	A	T	AKK.....		
AS	L	R	K	L	Q	A	D	G	R	I	T	E	E	QAKAYKEYHDKNGGANRKVND	
AS	L	R	K	L	Q	A	D	G	R	I	T	E	E	QAKAYKEYHDKNGGANRKVND	
AS	L	R	K	L	Q	A	D	G	R	I	T	E	E	QAKAYKEYHDKNGGANRKVND	
AS	L	R	K	L	Q	A	D	G	R	I	T	E	E	QAKAYKEYNDKNGGANRKVND	
AS	L	S	K	L	Q	A	D	G	R	I	T	E	E	QAKAYNEYAKNGGANRKVND	
AS	L	R	K	L	Q	D	D	G	R	I	T	E	E	QAKAYKEYHDKQNTSQIVKD	
AS	L	S	K	L	Q	A	D	G	R	I	T	E	E	QAKAYKEYHAKQNTSQTVAD	
AT	L	S	K	L	V	A	A	G	N	I	S	Q	K	QADS	
AT	L	S	K	L	V	A	A	G	N	I	S	Q	K	QADS	
AT	L	S	K	L	V	A	S	G	D	I	T	E	Q	E	QANSYKDYAKHPSKTRHVN
AT	L	S	K	L	V	A	S	G	D	I	T	E	Q	E	QANSYKDYAKHPSKTRHVN

- ☐ non conserved
- ☒ similar
- ☒ > 50% conserved