

logo

NTDB id 1037947 ACB335 RS00665 WP 002986552.1		MINQWNN	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	IVLA	IISIL	ILLFVPNL	IK	EKAQV	QKT	GEA	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																				

logo

NTDB id 1037947 ACB335 RS00665 WP 002986552.1	P	S	L	S	Q	L	K	A	D	G	S	I	T	E	K	Q	E	K	A	Y	Q	D	Y	D	K	H	K	N	E	K	A	R	L	S	N	108
NTDB id 588 KW2 RS10845 WP 051303741.1	P	N	L	S	E	L	L	S	A	G	M	I	T	Q	K	Q	V	T	A	Y	D	D	Y	D	Q	N	K	N	E	Q	R	N	F	D	116	
NTDB id 460 GPW51 RS00790 WP 011921720.1	A	T	V	A	D	L	Q	A	K	G	Y	I	T	E	K	Q	A	K	Q	Y	A	T	A	K	.	.	.	.	.	.	.	.	.	.	93	
NTDB id 502 SM12261 RS00955 WP 000738641.1	A	S	L	S	K	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	A	K	N	T	S	Q	T	V	A	D	107	
NTDB id 272 KZH43 RS09255 WP 000738626.1	A	S	L	R	K	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	D	K	N	G	G	A	N	R	K	V	N	108
NTDB id 231 SPD RS09885 WP 000738626.1	A	S	L	R	K	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	D	K	N	G	G	A	N	R	K	V	N	108
NTDB id 197 SPR RS09420 WP 000738626.1	A	S	L	R	K	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	H	D	K	N	G	G	A	N	R	K	V	N	108
NTDB id 162 SP RS10370 WP 000738627.1	A	S	L	R	K	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	K	E	Y	N	D	K	N	G	G	A	N	R	K	V	N	108
NTDB id 530 SMSK321 RS09230 WP 000738643.1	A	S	L	S	K	L	Q	A	D	G	R	I	T	E	E	Q	A	K	A	Y	N	E	Y	A	K	N	G	G	A	N	R	K	V	N	108	
NTDB id 431 SGO RS09410 WP 012130940.1	A	T	L	S	K	L	V	A	A	G	N	I	S	Q	K	Q	A	D	S	Y	K	A	Y	G	K	N	N	S	E	T	Q	A	V	A	N	105
NTDB id 396 FSA28 RS09395 WP 002263437.1	A	T	L	S	K	L	V	A	S	G	D	I	T	Q	E	Q	A	N	S	Y	K	D	Y	A	K	H	P	S	K	T	R	H	V	A	N	104
NTDB id 387 SMU RS09025 WP 002263437.1	A	T	L	S	K	L	V	A	S	G	D	I	T	Q	E	Q	A	N	S	Y	K	D	Y	A	K	H	P	S	K	T	R	H	V	A	N	104
consensus																																				

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