

logo

NTDB id 1037127 ACB360 RS00605 WP 002986552.1		MINQWNN	RHKKL	KGFTLL	EMLLV	ILVISVL	MLLFVPNL	SKQKDRVT	ETGNA	AAVVKL	VENQA	ELYEL	SLSGG	.SK	72	
NTDB id 588 KW2 RS10845 WP 051303741.1	MSKALS	SLV	KIHGRK	LCQKK	QKKAFTL	IEMLIV	LAIIIS	ILILLFVPNL	IKKKAQV	QKTGE	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			**	*	**	**	*****	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	!!!!!!	**

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NTDB id 1037127 ACB360 RS00605 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	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