

NTDB id 25 SA RS06185 WP 000593194.1	MIK	LFL	LK	L	Y	A	H	F	S	T	K	Q	I	H	Q	F	L	M	A	P	N	V	I	K	E	E	G	R	K	K	D	S	Y	C	E	W	VNREENV	L	L	R	K	Y	Y	A	F	I	K	L	D	H	N	66													
NTDB id 1250 GC085 RS13065 WP 027219989.1	..	M	N	L	Q	P	L	L	A	N	R	M	K	..	R	I	G	P	R	T	V	L	K	L	Q	K	R	W	P	D	L	N	L	M	F	Q	L	S	S	V	D	L	E	E	A	G	L	P	S	W	L	A	Q	...T	I	K	N	F	D	P	61				
NTDB id 599 KW2 RS05940 WP 021037268.1	..	M	I	T	N	F		L	F	R	W	..	K	A	G	M	T	N	L	G	V	N	K	..	L	L	K	F	R	K	Y	D	R	K	I	S	L	R	Q	M	G	..Q	V	A	Q	V	K	S	I	P	N	F	I	E	...Q	Y	K	N	Q	D	V	61			
NTDB id 1069091 ACF1RT RS05465 WP 002262865.1	..	M	D	N	F	Q		L	F	K	L	K	..	K	A	G	L	T	N	L	N	I	L	I	I	D	Y	E	E	R	T	Q	K	S	L	S	R	D	M	A	..V	S	K	N	K	K	P	L	I	F	M	E	..H	Y	K	N	L	D	S	K	60				
NTDB id 407 SMU RS04605 WP 002262865.1	..	M	D	N	F	Q		L	F	K	L	K	..	K	A	G	L	T	N	L	N	I	L	I	I	D	Y	E	E	R	T	Q	K	S	L	S	R	D	M	A	..V	S	K	N	K	K	P	L	I	F	M	E	..H	Y	K	N	L	D	S	K	60				
NTDB id 525 SMSK321 RS06295 WP 000705298.1		M	K	I	T	N	Y	E		I	Y	K	L	K	..	K	S	G	L	T	N	Q	Q	I	L	K	..V	L	E	Y	G	E	N	V	D	Q	E	L	L	G	D	I	A	..D	I	S	G	C	R	N	P	A	V	F	M	E	..R	Y	F	Q	I	D	D	A	62
NTDB id 267 KZH43 RS05590 WP 000705306.1		M	K	I	T	N	Y	E		I	Y	K	L	K	..	K	S	G	L	T	N	Q	Q	I	L	K	..V	L	E	Y	G	E	N	V	D	Q	E	L	L	G	D	I	A	..D	I	S	G	C	R	N	P	A	V	F	M	E	..R	Y	F	Q	I	D	D	A	62
NTDB id 226 SPD RS05990 WP 000705306.1		M	K	I	T	N	Y	E		I	Y	K	L	K	..	K	S	G	L	T	N	Q	Q	I	L	K	..V	L	E	Y	G	E	N	V	D	Q	E	L	L	G	D	I	A	..D	I	S	G	C	R	N	P	A	V	F	M	E	..R	Y	F	Q	I	D	D	A	62
NTDB id 192 SPR RS05715 WP 000705306.1		M	K	I	T	N	Y	E		I	Y	K	L	K	..	K	S	G	L	T	N	Q	Q	I	L	K	..V	L	E	Y	G	E	N	V	D	Q	E	L	L	G	D	I	A	..D	I	S	G	C	R	N	P	A	V	F	M	E	..R	Y	F	Q	I	D	D	A	62
NTDB id 157 SP RS06205 WP 000705306.1		M	K	I	T	N	Y	E		I	Y	K	L	K	..	K	S	G	L	T	N	Q	Q	I	L	K	..V	L	E	Y	G	E	N	V	D	Q	E	L	L	G	D	I	A	..D	I	S	G	C	R	N	P	A	V	F	M	E	..R	Y	F	Q	I	D	D	A	62
NTDB id 497 SM12261 RS05565 WP 000705318.1		M	K	I	T	N	Y	E		I	Y	K	L	K	..	K	S	G	L	T	N	Q	Q	V	L	A	..V	L	E	Y	G	E	N	V	D	Q	E	L	L	G	D	I	A	..E	I	S	G	C	R	N	P	A	V	F	M	E	..R	Y	F	Q	I	D	D	A	62
consensus		*****																																																															

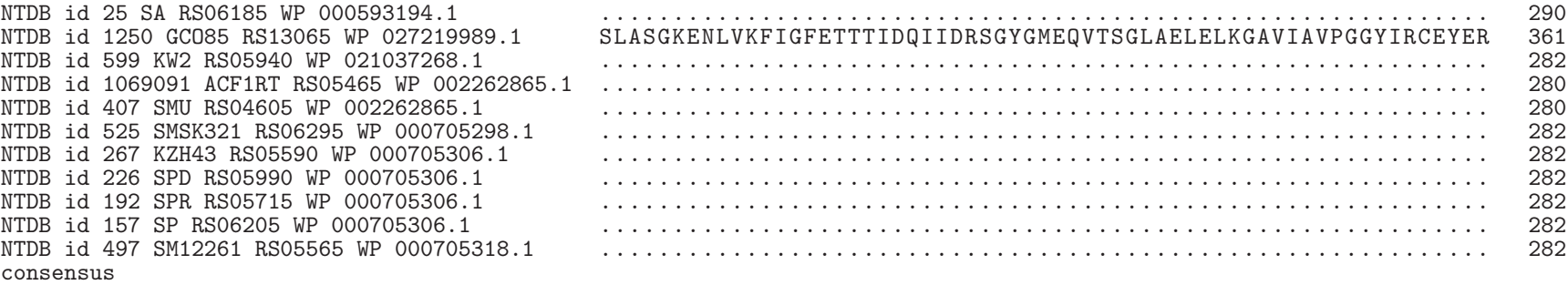
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NTDB id	25 SA RS06185 WP 000593194.1	YLTIVSGLAQGADAMAHQIALKYNLP TIAVLAFGHQTHYPKSTLALRNKTEEGKLVISEYPPHTPIAKYRFPERNRIISG	222
NTDB id	1250 GC085 RS13065 WP 027219989.1	GVSIVSGLALGIDAAHLCCLGGGKTIAVLGTGIDCIYPRRHCLKAEQITENGILLSEFPLKSPPIAGHFPLNRRIISG	220
NTDB id	599 KW2 RS05940 WP 021037268.1	SFTIVSGLAKGIDTASHLSAIKTKTP TIAVIGTGLDIFYPLENRRKIQEYLAKYQLVLSEYSLGEEKPLKYHFFPERNRIIAG	213
NTDB id	1069091 ACF1RT RS05465 WP 002262865.1	QFVIISGLARGIDTA AHLASLSKSGGA TIAVIGTGLDVHYPKENRRLLQDYIAKNHLLSEYEAAQS QPLKYHFFPERNRIIAG	212
NTDB id	407 SMU RS04605 WP 002262865.1	QFVIISGLARGIDTA AHLASLSKSGGA TIAVIGTGLDVHYPKENRRLLQDYIAKNHLLSEYEAAQS QPLKYHFFPERNRIIAG	212
NTDB id	525 SMSK321 RS06295 WP 000705298.1	ELVIVSGLAKGIDTA AHMAALQNGGKTIAVIGTGLDVFYPKANKRLQDYYISNDHVLVSEYGPGEQPLKFHF PARNRIIAG	214
NTDB id	267 KZH43 RS05590 WP 000705306.1	ELVIVSGLAKGIDTA AHMAALQNGGKTIAVIGTGLDVFYPKANKRLQDYYISNDHVLVSEYGPGEQPLKFHF PARNRIIAG	214
NTDB id	226 SPD RS05990 WP 000705306.1	ELVIVSGLAKGIDTA AHMAALQNGGKTIAVIGTGLDVFYPKANKRLQDYYISNDHVLVSEYGPGEQPLKFHF PARNRIIAG	214
NTDB id	192 SPR RS05715 WP 000705306.1	ELVIVSGLAKGIDTA AHMAALQNGGKTIAVIGTGLDVFYPKANKRLQDYYISNDHVLVSEYGPGEQPLKFHF PARNRIIAG	214
NTDB id	157 SP RS06205 WP 000705306.1	ELVIVSGLAKGIDTA AHMAALQNGGKTIAVIGTGLDVFYPKANKRLQDYYISNDHVLVSEYGPGEQPLKFHF PARNRIIAG	214
NTDB id	497 SM12261 RS05565 WP 000705318.1	ELVIVSGLAKGIDTA AHMAALQNGGKTIAVIGTGLDVFYPKANKRLQDYYISNDHVLVSEYGPGEQPLKFHF PARNRIIAG	214
consensus		***!*****!!**!*!!!!!!*	

logo



logo



X non conserved
X similar
X ≥ 50% conserved