

logo

NTDB id 25 SA RS06185 WP 000593194.1	SHITIDFALEQNRNVYVLPGSMFNPMTKGNLLRIQEGAKVVLNANDIFEDYYI	290
NTDB id 1250 GC085 RS13065 WP 027219989.1	SLITARMALEQNRDVLAI PGSIHNPLARGCHYL LQQGAKLVTSVADVLDDELKIEHHQFTSNKPIFSLASGKENLVKFIGFETTTI	320
NTDB id 599 KW2 RS05940 WP 021037268.1	SLITCERALEEGRDIFAIPGNIADGTSDGCNHLIQQGAKLVYQAQDILEEYLYN	282
NTDB id 407 SMU RS04605 WP 002262865.1	SLITCERAMEEGRDVFVVPGNILDGQSEGCHHLIQEGAKCITSGFDILNEFNF	280
NTDB id 61464 SCRE RS04440 WP 020997792.1	SLITCERAMEEGRDVF AIPGSILDGRSDGCHHLIQEGAKC IASGADVLS EFQF	280
NTDB id 525 SMSK321 RS06295 WP 000705298.1	SLITCERAMEEGRDVF AIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLA EF EF	282
NTDB id 497 SM12261 RS05565 WP 000705318.1	SLITCERAMEEGRDVF AIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLA EF EF	282
NTDB id 267 KZH43 RS05590 WP 000705306.1	SLITCERAMEEGRDVF AIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLA EF EF	282
NTDB id 226 SPD RS05990 WP 000705306.1	SLITCERAMEEGRDVF AIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLA EF EF	282
NTDB id 192 SPR RS05715 WP 000705306.1	SLITCERAMEEGRDVF AIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLA EF EF	282
NTDB id 157 SP RS06205 WP 000705306.1	SLITCERAMEEGRDVF AIPGSILDGLSDGCHHLIQEGAKLVTSGQDVLA EF EF	282
consensus	! * ! ! * * * ! * ! * * ! * * * * ! ! * * * * * * * ! * * * * * ! * ! ! ! * * * * * ! * * * * * *	

logo

NTDB id 25 SA RS06185 WP 000593194.1	DQ I I D R S G Y G M E Q V T S G L A E L E L K G A V I A V P G G Y I R C E Y E R	290
NTDB id 1250 GC085 RS13065 WP 027219989.1	DQ I I D R S G Y G M E Q V T S G L A E L E L K G A V I A V P G G Y I R C E Y E R	361
NTDB id 599 KW2 RS05940 WP 021037268.1		282
NTDB id 407 SMU RS04605 WP 002262865.1		280
NTDB id 61464 SCRE RS04440 WP 020997792.1		280
NTDB id 525 SMSK321 RS06295 WP 000705298.1		282
NTDB id 497 SM12261 RS05565 WP 000705318.1		282
NTDB id 267 KZH43 RS05590 WP 000705306.1		282
NTDB id 226 SPD RS05990 WP 000705306.1		282
NTDB id 192 SPR RS05715 WP 000705306.1		282
NTDB id 157 SP RS06205 WP 000705306.1		282
consensus		

- non conserved
- similar
- ≥ 50% conserved