

logo

NTDB id 1250 GC085 RS13065 WP 027219989.1
NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 49534 SPYALAB49 RS04335 WP 002995009.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus



logo

NTDB id 1250 GC085 RS13065 WP 027219989.1
NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 49534 SPYALAB49 RS04335 WP 002995009.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus



logo

NTDB id 1250 GC085 RS13065 WP 027219989.1
NTDB id 599 KW2 RS05940 WP 021037268.1
NTDB id 525 SMSK321 RS06295 WP 000705298.1
NTDB id 267 KZH43 RS05590 WP 000705306.1
NTDB id 226 SPD RS05990 WP 000705306.1
NTDB id 192 SPR RS05715 WP 000705306.1
NTDB id 157 SP RS06205 WP 000705306.1
NTDB id 497 SM12261 RS05565 WP 000705318.1
NTDB id 49534 SPYALAB49 RS04335 WP 002995009.1
NTDB id 407 SMU RS04605 WP 002262865.1
consensus



logo

NTDB id 1250 GC085 RS13065 WP 027219989.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	313
NTDB id 599 KW2 RS05940 WP 021037268.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 525 SMSK321 RS06295 WP 000705298.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 267 KZH43 RS05590 WP 000705306.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 226 SPD RS05990 WP 000705306.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 192 SPR RS05715 WP 000705306.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 157 SP RS06205 WP 000705306.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 497 SM12261 RS05565 WP 000705318.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	282
NTDB id 49534 SPYALAB49 RS04335 WP 002995009.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	278
NTDB id 407 SMU RS04605 WP 002262865.1	SGSLITCERAMEEGRDVFALPGSILDLGSLDGGCHHLIQEGAKLVTSQDVLAEFEF	280
consensus	!!!!!!*****!***!*****!*!*****!***!***!*****!*!*!***	

logo

NTDB id 1250 GC085 RS13065 WP 027219989.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	361
NTDB id 599 KW2 RS05940 WP 021037268.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 525 SMSK321 RS06295 WP 000705298.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 267 KZH43 RS05590 WP 000705306.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 226 SPD RS05990 WP 000705306.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 192 SPR RS05715 WP 000705306.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 157 SP RS06205 WP 000705306.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 497 SM12261 RS05565 WP 000705318.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	282
NTDB id 49534 SPYALAB49 RS04335 WP 002995009.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	278
NTDB id 407 SMU RS04605 WP 002262865.1	GFETTTIDQIIDRSQYGMQVTSGLAELELKGAVIAVPGGYIRCEYER	280
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved