

Sequence type, (Isolate ID)	GI No. 1 upstream <i>dusC</i> , approx. pos. 38,000	GI No. 2 downstream <i>dusC</i> , approx. pos. 38,000	GI No. 3, approx. pos. 40-80,000	GI No. 4, approx. pos. 170,000,	GI No. 5, approx. pos. 280,000	GI No. 6 (ssI/lpf), approx. position 400,000	GI No. 7, approx. pos. 900,000	GI No. 8, pos. 1,500,000	GI No. 9 (<i>egc</i> and <i>lukD/E</i>) approx. pos. 1,800,000	GI No. 10 (<i>sspP/sspS</i>), approx. pos. 2,000,000	GI No. 11, approx. pos. 2,500,000	GI No. 12, approx. pos. 2,530,000	GI No. 13 (st 'xanthin'), approx. position 2,650,000	GI No. 14, approx. pos. 2,770,000
ST2990 (27-G-H)	Q7A899, <i>dusC</i> -trnc, DUF81-GI, <i>cstR/A/B</i> - GI, <i>sqr</i> , <i>xyiE2</i>	Q6GKK6, Q7A890	<i>lpl</i> , <i>hysA</i> , 5 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, DUF600, a 2 nd copy of <i>essC</i> , 2 copies of DUF600, Q1Y4R3, DUF600, ADI96784, DUF4064, DUF5084, 2 copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /06, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	negative	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, <i>fstA</i> , AJP22988, DUF1433, DUF1433, Q6GFP5, A0A0E1X646, Q6GFP3, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /E, <i>sp</i> /D1, <i>sp</i> /D2-1, <i>sp</i> /D2- 2, <i>sp</i> /C, <i>sp</i> /A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD/E</i> , <i>sprC1</i> , Q2FXA9, Q2FXA8, Q7A4X2	<i>sspP/sspS</i>	teg131, 3 copies of <i>lpl</i> sauUSI, <i>nudix</i>	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	<i>cna</i>
ST3268 (Ma2/A14043)	mcrB, mcrC, tnp_IS1, tnp_IS232, Q6GD44, <i>dusC</i> -trnc, DUF81-GI, <i>cstR/A/B</i> -GI, <i>sqr</i>	Q6GKK6, <i>yycY</i> , G7ZTC1, Q2YUT3	3 copies of <i>lpl</i>	Q5HUH7-trnc	Transposon with the penicillinase operon, followed by 3copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /06, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, <i>lip7</i> , <i>istB2</i> IS232, tnp_IS232	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	Q2YY95, <i>sagD</i> , <i>sagC</i> , <i>nfnB</i> , D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i> , tnp_IS232, tnp_IS200	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, DUF1433, <i>hysA</i> , DUF1433, ERS140266 RS01930, <i>hsdS/M</i> - <i>spl</i> , <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD/E</i> , <i>sprC1</i> , Q2FXA9, Q2FXA8, Q99T51, A6U2S7, Q7A4X2, <i>seg/n/u/i/m/o</i>	<i>sspP/sspS</i>	teg131, sau961, dcm_Sau961	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST4168 (16CS0209)	tnp_ISSha1, C2G7A0, Q6GKL3, C2G798, UPI0001C11B31, Q6GKL1, tnp_IS232, Q6GD44	Q6GKK6, Q7A890	<i>lpl</i>	Q5HUH7-trnc	A0A0E0VL30, DUF5082, UPI0001C11B8A, Q1Y4R2, Q6GK19, DUF5085, Q1Y4R3, 2 copies of DUF600, Q1Y4R2, Q1Y4R3, Q9L3N6, 4 copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 4 copies of <i>lpl</i> , <i>lipC3</i> ,	<i>oppC</i> -GI-RF122, <i>oppD</i> -GI-RF122, <i>oppF</i> -GI-RF122, <i>oppA</i> -GI-RF122	Q931R4	Q5HEX7, A5ITW8, Q5HEX3, <i>sprA</i> , <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /E, <i>sp</i> /D2, <i>sp</i> /C/B/A, <i>ear2</i> , <i>epiG</i> , ERS140266 RS14190, <i>epiF</i> , <i>epiD</i> , <i>epiC</i> , <i>epiB</i> , <i>epiA</i> , <i>lukD/E</i> , <i>sprC1</i> , Q6GFN4, tnp_IS1, Q7A4X2, <i>seg/n/u/i/m/o</i>	<i>sspP/sspS</i>	teg131	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	<i>cna</i>
ST7687 (01-RR-86)	Q7A899, <i>dusC</i> -trnc, DUF81-GI, <i>cstR/A/B</i> - GI, <i>sqr</i> , <i>xyiE2</i> (as in CC5/8)	Q6GKK6, Q7A890	3 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	A0A0E0VL30, DUF5082, UPI0001C11B8A, Q6GK19, DUF5085, Q1Y4R3, Q9L3N6, DUF600, Q1Y4R2, 3 copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /06, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 6 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	Q2YY95, <i>sagD</i> , <i>sagC</i> , <i>nfnB</i> , D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i>	Q5HEX7, A5ITW8, DUF955, Q5HEX3, DUF1433, <i>sprA</i> , AJP22988, 3 copies of DUF1433, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD/E</i> , <i>sprC1</i> , Q2FXA9, Q2FXA8, Q7A4X2	<i>sspP/sspS</i>	teg131, Q5HD68, D2NA72, <i>lipC3</i> , 3 copies of <i>lpl</i> , sauS0385, dcm_SauS0385	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST7688 (05-RR-90)	Q7A899, <i>dusC</i> -trnc, DUF81-GI, <i>cstR/A/B</i> - GI, <i>sqr</i> , <i>xyiE2</i>	Q5HJT2, Q6GD34, A6QD75, A6QD76, A8YZ18, Q6GKK6, <i>yycY</i> , G7ZTC1, Q7A890, Q2YUT3	3 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, Q1Y4R2, Q1Y4R3, and 8 copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 3 copies of <i>lpl</i> , tnp_IS232, <i>istB2</i> IS232, 3 copies of <i>lpl</i> , <i>lipC3</i> SAV0446	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i>	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, 4 copies of DUF1433, <i>hysA</i> , Q5HE01, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /E, <i>sp</i> /D1, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD</i> , <i>lukE</i>	<i>sspP/sspS</i>	teg131, 2 copies of <i>lpl</i> , Q5HD68, D2NA72, <i>lipC3</i> , 2 copies of <i>lpl</i> , dcm_Sau3A1, sau3A1	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST7689 (08-G-E)	Q6GD44	Q6GKK6, Q7A890	2 copies of <i>lpl</i> , <i>lipC3</i>	Q5HUH7, Q9RL82, A6QDI2	<i>essC</i> , <i>essB</i> , <i>esaE</i> , <i>essD</i> , <i>essD</i> , Q1Y4R2, Q1Y4R3, 3 copies of DUF600, Q1Y4R2, Q1Y4R3, 2 copies of DUF600, ADI96784, DUF4064, DUF5084, 3 copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 6 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	negative	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, tnp_IS3, <i>sprA</i> , DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /D1, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , ASITY2, <i>lukD/E</i> , Q2FXA9, Q2FXA8, Q99T51, A6U2S7, Q7A4X2, <i>seg/n/u/i/m/o</i>	<i>sspP/sspS</i>	teg131, 2 copies of <i>lpl</i> , Q5HD68, D2NA72, <i>lipC3</i> , 3 copies of <i>lpl</i> , sauUSI, <i>nudix</i>	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST7690 (09-G-F)	Q7A899, <i>dusC</i> -trnc, DUF81-GI, <i>cstR/A/B</i> - GI, <i>sqr</i> , <i>xyiE2</i>	Q5HJT2, Q6GD34, A6QD75, A6QD76, A8YZ18, Q6GKK6, <i>yycY</i> , tnp-Q6GH59, tnp-Q6GJF3, Q7A890	3 copies of <i>lpl</i> , <i>hysA</i> , 3 copies of <i>lpl</i>	Q5HUH7, tnp- Q6GH59, tnp- Q6GJF3	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, 4 copies of DUF600, Q1Y4R2, Q1Y4R3, 4 copies of DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 6 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	negative	Q6GFQ3, DUF955, Q5HEX3, <i>istB2</i> IS232, tnp_IS232, <i>sprA</i> , DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /E, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD/E</i> , Q2FXA9, Q2FXA8, Q2FXA8, Q7A4X2	<i>sspP/sspS</i>	teg131, 4 copies of <i>lpl</i> <i>istB2</i> IS232, tnp_IS232, Q5HD68, D2NA72, <i>lipC3</i> , <i>lpl</i> , sauS0385, dcm_SauS0385	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	<i>cna</i>
ST7691 (13-G-52)	Q6GD44	Q6GKK6, Q7A890	2 copies of <i>lpl</i> , <i>lipC3</i>	Q5HUH7, Q9RL82, A6QDI2	<i>essC</i> , <i>essB</i> , <i>esaE</i> , <i>essD</i> , <i>essD</i> , TIGR01741, Q1Y4R2, Q1Y4R3, DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /06, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 6 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	negative	Q6GFQ3, DUF955, Q5HEX3, tnp_IS3, <i>sprA</i> , DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /D1, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , ASITY2, <i>lukD/E</i> , Q2FXA9, Q2FXA8, Q99T51, A6U2S7, Q7A4X2, <i>seg/n/u/i/m/o</i>	<i>sspP/sspS</i>	teg131, <i>lpl</i> , Q5HD68, D2NA72, <i>lipC3</i> , 3 copies of <i>lpl</i> sauUSI, <i>nudix</i>	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	<i>cna</i>
ST7692 (17-H-61)	Q6GD44	Q5HJT2, Q6GD34, A6QD75, A6QD76, A8YZ18, Q6GKK6, <i>yycY</i> , G7ZTC1, Q7A890, Q2YUT3	4 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q9L3N6, 2 copies of DUF600, Q1Y4R3, DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 5 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	Q2YY95, <i>sagC</i> , <i>nfnB</i> , D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i>	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, AJP22988, DUF1433, DUF1433, DUF1433, DUF1433, <i>hysA</i> , A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /E, <i>sp</i> /D1, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD/E</i> , A6U2S7, Q7A4X2, Q99T51, <i>int-2b</i>	<i>sspP/sspS</i>	teg131, 4 copies of <i>lpl</i> Q5HD68, D2NA72, <i>lipC3</i> , SACOL2496, 2 copies of <i>lpl</i> dcm_Sau3A1, sau3A1	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST7693 (29-P-01)	negative	Q6GKK6, <i>yycY</i> , G7ZTC1, Q7A890, Q2YUT3	<i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	<i>essC</i> , <i>essB</i> , <i>esaE</i> , <i>essD</i> , <i>essD</i> , TIGR01741, Q1Y4R2, Q1Y4R3, DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 6 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appB</i> , <i>appC</i>	negative	Q6GFQ3, DUF955, Q5HEX3, <i>sprA</i> , DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /E, <i>sp</i> /D1, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , ASITY2, <i>lukD/E</i> , A6U2S7, Q7A4X2, Q99T51, <i>int-2b</i>	<i>sspP/sspS</i> <i>sspP/sspS</i>	teg131, Q5HD68, D2NA72, <i>lipC3</i> , <i>lpl</i> , tx universal3, dcm_SauRF122, sauRF122, zoo28neu_002509	Q5HJU6, <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST7694 (40-B-50)	negative	Q6GKK6, <i>yycY</i> , G7ZTC1, Q7A890, Q2YUT3	5 copies of <i>lpl</i>	Q5HUH7-trnc	<i>essC</i> , <i>essB</i> , <i>esaE</i> , <i>essD</i> , <i>essD</i> , TIGR01741, Q1Y4R2, Q1Y4R3, DUF600	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /03, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i>	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, <i>sprA</i> , DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /D1, <i>sp</i> /C/B/A, A6QHZ7, <i>ear2</i> , <i>sprB</i> , <i>epiG</i> , ASITY2, ASITY3, <i>lukD/E</i> , <i>sprC1</i> , Q2FXA9, Q2FXA8, Q7A4X2	<i>sspP/sspS</i>	teg131, 4 copies of <i>lpl</i> sauUSI, <i>nudix</i>	A8YYP8, <i>sasG</i> , <i>sarT</i> , <i>sarU</i>	<i>cbiQ</i> , <i>cbiO</i> , UPF0397, DUF62	negative
ST7695 (16CS0212)	Q6GKL6	Q5HJT2, Q6GD34, A6QD75, A6QD76, A8YZ18, Q6GKK6, <i>yycY</i> , G7ZTC1	<i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	<i>essC</i> , <i>essB</i> , <i>esaE</i> , <i>essD</i> , <i>essD</i> , Q1Y4R2, Q1Y4R3, Q1Y4R3, Q9L3N6, DUF600, ADI96784, DUF4064, DUF5084	<i>ss</i> /01, <i>ss</i> /02, <i>ss</i> /04, <i>ss</i> /05, <i>ss</i> /07, <i>ss</i> /08, <i>ss</i> /09, <i>ss</i> /10, <i>hsdM/S</i> - <i>ssI</i> , <i>ss</i> /11, <i>slafP</i> , Q2G0X2, 5 copies of <i>lpl</i> , <i>lipC3</i>	<i>appA</i> , <i>appD</i> , <i>appF</i> , <i>appB</i> , <i>appC</i>	negative	Q5HEX7, A5ITW8, <i>sprA</i> , DUF1433, <i>hysA</i> , DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M</i> - <i>spl</i> , <i>sp</i> /F, <i>sp</i> /D2, <i>sp</i> /C/B/A, <i>ear2</i> , <i>epiG</i> , ERS140266 RS14190, <i>epiE</i> /P/D/C/B/A, <i>bsaX</i> , <i>lukD/E</i> , <i>sprC1</i> , Q2FXA9, Q2FXA8	<i>sspP/sspS</i>	teg131, BDS-53E_002366, <i>lpl</i> , Q5HD68, D2NA72, <i>lipC3</i> , 2 copies of <i>lpl</i> dcm_Sau3A1, sau3A1	negative	negative	negative

Sequence type, (Isolate ID)	GI No. 1 upstream <i>dusC</i> , approx. pos. 38,000	GI No. 2 downstream <i>dusC</i> , approx. pos. 38,000	GI No. 3, approx. pos. 40-80,000	GI No. 4, approx. pos. 170,000,	GI No. 5, approx. pos. 280,000	GI No. 6 (<i>ssI/lpl</i>), approx. position 400,000	GI No. 7, approx. pos. 900,000	GI No. 8, pos. 1,500,000	GI No. 9 (<i>egc</i> and <i>lukD/E</i>) approx. pos. 1,800,000	GI No. 10 (<i>sspP/sspS</i>), approx. pos. 2,000,000	GI No. 11, approx. pos. 2,500,000	GI No. 12, approx. pos. 2,530,000	GI No. 13 (st'xanthin), approx. position 2,650,000	GI No. 14, approx. pos. 2,770,000
ST7745 (03-RR-88)	<i>lrpC</i> , Q6GD44	Q6GKK6, Q7A890	5 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	Q6GK23, Q6GK22, Q6GK21, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, TIGR01741, 2 copies of DUF600, AD196784, DUF4467, DUF4064, DUF5084, DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI06, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, lpl, locus-0450, tnp IS232, istB2 IS232, lipC3_SAA6159_00398</i>	<i>appA, appD, appF, appB, appC</i>	Q2YY95, <i>sagD, sagC, nfbB, D9RHF6, Q2YYD5, D9RHF8, stsA</i>	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, DUF1433, <i>sprA</i> , AJP22988, 2 copies of DUF1433, <i>hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E, sprC1, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	<i>teg131, Q5HD68, D2NA72, lipC3, lpl, dcm SauRF122, sauRF122</i>	negative	<i>cbiO, cbiO, UPF0397, DUF62</i>	<i>cna</i>
ST7746 (07-G-D)	Q7A899, <i>dusC</i> -trnc, DUF81-GI, <i>csrR/A/B-GI, sgr, yxiE2</i>	Q5HJT2, Q6GD34, A6QD75, A6QD76, A8YZ18, Q6GKK6, <i>ycjY, G7ZTC1, Q7A890</i>	6 copies of <i>lpl</i>	A5IP65, A5IP66, A5IP67, A5IP68, A5IP69, A5IP70, A5IP71, A5IP72, A6TXY8, A5IP73, A6QDI2	A0A0Z1N098, A0A0U0VFZ6, A0A122YW60, AD196784, DUF4064, DUF5084, Q1Y4R2, Q1Y4R3, Q9L3N6, 2 copies of DUF600, Q1Y4R2, Q1Y4R3, 2 copies of DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI06, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2</i>	<i>appA, appD, appF, appB, appC</i>	negative	DUF1433, 2 copies of DUF1433, Q6GFP5, A0A0E1X646, <i>hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E, sprC1, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	<i>teg131, 2 copies of lpl, Q5HD68, D2NA72, lipC3, lpl, tnp IS232, istB2 IS232, 2 copies of lpl, D2NA72, lipC3, lpl, sauS0385, dcm_SauS0385</i>	negative	<i>cbiQ, cbiO, UPF0397, DUF62</i>	<i>cna</i>
ST7747 (12-G-51)	H0CD41, D2N3D1, C2G7A0, Q6GKL3, C2G798, UPI0001C11B31, D2N3E0, Q6GKL1, tnp IS232, Q6GD44 <i>Similar as in CC398 (except for some tnp)</i>	Q6GKK6, Q7A890	<i>lpl, hysA, 2 copies of lpl, istB2 IS232, tnp IS232, lpl</i>	Q5HUH7, Q9RL82, A6QDI2	A0A0E0VL30, DUF5082, UPI0001C11B8A, Q1Y4R2, Q6GK19, DUF5085, Q1Y4R3, Q9L3N6, DUF600, Q1Y4R2, Q1Y4R3, 3 copies of DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, istB2 IS232, tnp IS232, 3 copies of lpl</i>	<i>appA, appD, appF, appB, appC</i>	Q2YY95, <i>sagD, sagC, nfbB, D9RHF6, Q2YYD5, D9RHF8, stsA</i>	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, DUF1433, <i>hysA, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, ERS140266 RS14190, epiE/F/P/D/C/B/A, bsaX, epiA, lukD/E, Q2Y2TQ0, Q6GFN4, tnp IS1, A6U2S7, Q7A4X2, Q99T51</i>	<i>sspP/sspS</i>	<i>teg131, lpl, Q5HD68, D2NA72, lipC3, lpl, sauS0385, dcm_SauS0385</i>	A8YYP8	negative	negative
ST7748 (15-G-54)	Q6GD44	Q6GKK6, Q7A890	2 copies of <i>lpl, lipC3</i>	Q5HUH7, Q9RL82, A6QDI2	<i>exsC, exsB, esaE, exsD, essD, TIGR01741, Q1Y4R2, Q1Y4R3, DUF600</i>	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, 6 copies of lpl, lipC3</i>	<i>oppB-GI-RF122, oppC-GI-RF122, oppD-GI-CIG93, oppF-GI-RF122, oppA-GI-RF122</i>	negative	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, tnp IS3, tnp IS3, r122, <i>sprA, DUF1433, Q2FXD2, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, lukD/E, Q2FXA9, Q2FXA8, Q99T51, A6U2S7, Q7A4X2, seg/n/u/i/m/o</i>	<i>sspP/sspS</i>	<i>teg131, BDS-53E 002366, lpl, Q5HD68, D2NA72, lipC3, 2 copies of lpl, dcm_Sau3AI, sau3AI</i>	Q5HJU6, sarT, sarU	<i>cbiQ, cbiO, UPF0397, DUF62</i>	<i>cna</i>
ST7749 (18-H-62)	<i>hsdM5, hsdR5, Q6GKL3, C2G798, D2N3E0, Q6GKL1, tnp IS232, Q6GD44</i>	Q6GKK6, Q7A890	3 copies of <i>lpl, hysA, 3 copies of lpl</i>	Q5HUH7, Q9RL82, A6QDI2	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, 2 copies of DUF600, Q1Y4R2, Q1Y4R3, 2 copies of DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, 6 copies of lpl, lipC3</i>	<i>appA, appD, appF, appB, appC</i>	D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i>	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, <i>sprA, DUF1433, Q6GFP5, A0A0E1X646, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, ERS140266 RS14190, epiE/F/P/D/C/B/A, lukD/E, sprC1, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	<i>teg131, BDS-53E 002366, 2 copies of lpl Q5HD68, D2NA72, lipC3 SACOL2496, 2 copies of lpl dcm_Sau3AI, sau3AI</i>	A8YYP8	<i>cbiQ, cbiO, UPF0397, DUF62</i>	negative
ST7750 (26-G-G)	H0CD41, D2N3D1, Q6GD44	Q6GKK6	3 copies of <i>lpl, lipC3</i>	A5IP65, A5IP66, A5IP67, A5IP68, A5IP69, A5IP70, A5IP71, A5IP72, A6TXY8, A5IP73	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, 3 copies of DUF600, AD196784, DUF4467, DUF4064, DUF5084, 4 copies of DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, 3 copies of lpl, lipC3</i>	<i>appA, appD, appF, appB, appC</i>	Q2YY95, <i>sagD, sagC, nfbB, D9RHF6, Q2YYD5, D9RHF8, stsA</i>	DUF1433, <i>hysA, DUF1433, DUF1433, Q6GFP5, A0A0E1X646, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E, sprC1, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	<i>teg131, BDS-53E 002366, 2 copies of lpl Q5HD68, D2NA72, lipC3, 2 copies of lpl, dcm_Sau3AI, sau3AI</i>	A8YYP8, <i>sasG, sarT, sarU</i>	<i>cbiQ, cbiO, UPF0397, DUF62</i>	<i>cna</i>
ST7751 (28-G-1)	C2G7A0, Q6GKL3, C2G798, UPI0001C11B31, D2N3E0, Q6GKL1, Q6GD44	Q6GKK6, <i>ycjY, G7ZTC1, Q7A890</i>	<i>lpl, hysA, 4 copies of lpl</i>	Q5HUH7, Q9RL82, A6QDI2	<i>exsC, exsB, esaE, exsD, essD, Q1Y4R2, Q1Y4R3, Q9L3N6, 2 copies of DUF600, Q1Y4R2, Q1Y4R3, TIGR01741, DUF600</i>	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, 3 copies of lpl</i>	<i>appA, appD, appF, appB, appC</i>	negative	Q5HEX7, A5ITW8, DUF955, Q5HEX3, DUF1433, <i>sprA, DUF1433, DUF1433, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	<i>teg131, lpl, Q5HD68, D2NA72, lipC3, 3 copies of lpl dcm_Sau3AI, sau3AI</i>	Q5HJU6, sarT, sarU	<i>cbiQ, cbiO, UPF0397, DUF62</i>	negative
ST7752 (30-P-10)	Q7A899, <i>dusC</i> -trnc, DUF81-GI, <i>csrR/A/B-GI, sgr, yxiE2</i>	Q5HJT2, Q6GD34, A6QD75, A6QD76, A8YZ18, Q6GKK6, <i>ycjY, G7ZTC1</i>	5 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	A0A0Z1N098, A0A0U0VFZ6, A0A122YW60, AD196784, DUF4064, DUF5084, Q1Y4R2, Q1Y4R3, Q9L3N6, 2 copies of DUF600, Q1Y4R2, Q1Y4R3, 4 copies of DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI06, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, 5 copies of lpl, lipC3</i>	<i>appA, appD, appF, appB, appC</i>	negative	Q5HEX7, A5ITW8, Q6GFQ3, DUF955, Q5HEX3, <i>fstAT, AJP22988, DUF1433, DUF1433, Q6GFP5, A0A0E1X646, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E</i>	<i>sspP/sspS</i>	<i>teg131, lpl-locus-2500, dcm_Sau3AI, sau3AI</i>	A8YYP8	<i>cbiQ, cbiO, UPF0397, DUF62</i>	negative
ST7753 (32-T-13)	<i>mcrC, dhIC</i> -trnc, Q6GD44	Q6GKK6, Q7A890	6 copies of <i>lpl</i>	Q5HUH7, Q9RL82, A6QDI2	<i>exsC, exsB, esaE, exsD, essD, Q1Y4R2, Q1Y4R3, 3 copies of DUF600, Q1Y4R2, Q1Y4R3, 2 copies of DUF600, AD196784, DUF4064, DUF5084, 3 copies of DUF600</i>	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, tnp IS232, istB2 IS232, 4 copies of lpl,</i>	<i>appA, appD, appF, appB, appC</i>	D9RHF6, Q2YYD5, D9RHF8, <i>stsA</i>	Q6GFQ3, DUF955, Q5HEX3, <i>sprA, DUF1433, Q6GFP5, A0A0E1X646, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	<i>teg131, Q5HD68, D2NA72, lipC3, lpl, tx universal3, dcm_SauRF122, sauRF122</i>	A8YYP8	<i>cbiQ, cbiO, UPF0397, DUF62</i>	<i>cna</i>
ST7754 (39-B-49)	C1PH94, C1PH95, istB2 IS232, tnp IS232, <i>lrpC, Q6GD44, tnp IS1272, DUF81-GI, csrR/A/B-GI, sgr, yxiE2</i>	Q6GKK6, Q7A890	<i>lpl, lipC3</i>	Q5HUH7, Q9RL82, A6QDI2	Q6GK23, Q6GK22, Q6GK21, Q2G192, Q6GK19, Q6GK18, Q6GK17, Q9L3N6, 2 copies of DUF600	<i>ssI01, ssI02, ssI03, ssI04, ssI05, ssI07, ssI08, ssI09, ssI10, hsdM/S-ssI, ssI11, slaP, Q2G0X2, 3 copies of lpl, lipC3</i>	<i>appA, appD, appF, appB, appC</i>	Q931R4	Q6GFQ3, DUF955, Q5HEX3, tnp IS3, tnp IS3, r122, <i>sprA, DUF1433, Q2FXD2, hsdS/M-spl, spIF, spIC/B/A, A6QHZ7, ear2, sprB, epiG, A5ITY2, A5ITY3, lukD/E, sprC1, Q2FXA9, Q2FXA8, Q7A4X2</i>	<i>sspP/sspS</i>	Q5HD68, D2NA72, <i>lipC3, 2 copies of lpl sauUSI, nudix</i>	negative	<i>cbiQ, cbiO, UPF0397, DUF62</i>	<i>cna</i>