

## Supplemental Table 1. List of Primers.

Primer	Sequence	
Primers for Gram-negative deletions	( <b>Boldface</b> characters indicate the 20 nucleotides homologous to the FRT-PGK-gb2-neo-FRT cassette.)	deletion
Ecoli-pgaA-del5F	CTGTAATTAGATATAGAGAGAGATTTTGGCAATACATGGAGTAAT ACAGGA <b>ATTAACCCTCACTAAAGGGCGG</b>	<i>pgaA</i>
Ecoli-pgaA-del3R	ACTCACCAGCATCAGGATATATTTATTTCCATTACGTAACATATTT ATCCTA <b>ATACGACTCACTATAGGGCTC</b>	<i>pgaA</i>
Ecoli-pgaB-del5F	CACAACCTTATACGTTGAATTCGATATGACATTCAGATTTTAAGGAT AAATA <b>ATTAACCCTCACTAAAGGGCGG</b>	<i>pgaB</i>
Ecoli-pgaB-del3R	ATGCATAACACCAGACATAATATAAAAAACGATACGATGCGATTAA TCATTA <b>ATACGACTCACTATAGGGCTC</b>	<i>pgaB</i>
Ecoli-pgaC-del5F	CCAGTAAACGTAAGTGGTGTGCGGGTCGTAATAATTAATTTGTT CATAA <b>ATTAACCCTCACTAAAGGGCGG</b>	<i>pgaC</i>
Ecoli-pgaC-del3R	ACCTTATTCGTCCTGAGTTTTCAACAGCCTGGTATCCGAAAAATGA TTAATA <b>ATACGACTCACTATAGGGCTC</b>	<i>pgaC</i>
Ecoli-pgaD-del5F	GCAACGCGCCCGTTGGGTAAGTCCCAGATCGCGGGATTCTGAGAGGT TAATA <b>ATTAACCCTCACTAAAGGGCGG</b>	<i>pgaD</i>
Ecoli-pgaD-del3R	AGTGTATTATCGGGCAGAGCCCAGGCGAACCAGGGCTTTGTTTTGG GTGT <b>TAATACGACTCACTATAGGGCTC</b>	<i>pgaD</i>
Kp-pgaC-del5F	TGATATCTCCCGTATCAGGCCGCAATTTTCTTCCTGGTGGTACCCTG ACCA <b>ATTAACCCTCACTAAAGGGCGG</b>	<i>pgaC</i>
Kp-pgaC-del3R	AGCCGGGGCAGCACACGGTGTTCGGTTAAGATGAGCGTATTTTCAT TCATTA <b>ATACGACTCACTATAGGGCTC</b>	<i>pgaC</i>
Kpdelpga locUP	AGATCACCTTATTCAAATACGTTATTGTTTAATCGCAACACGAAAC TCTCA <b>ATTAACCCTCACTAAAGGGCGG</b>	<i>pga</i> locus
Kpdelpga locDOWN	GGTAACGCGTGGCTTTTAAATCAATAATCTGCCCGATTTGATCGT GGTATA <b>ATACGACTCACTATAGGGCTC</b>	<i>pga</i> locus
Primers for <i>B. subtilis</i>		
3971	AGGAGGA <b>ATTCTT</b> AAAAAGGATTTGACTTCATGGACA	<i>epsA</i> up
3972	CTCCT <b>CTCGAG</b> CATATTCTCATTTCATGTATTCATAGC	<i>epsA</i> up
3973	AGGAG <b>CTCGAG</b> AAACATTTTCGGGGAGTGAAGCGA	<i>epsA</i> down
3974	CTCCT <b>GGATCCA</b> ATAGTGTTCCGATTTTTTTCATTTTCT	<i>epsA</i> down
3975	AGGAGGA <b>ATTCAGA</b> ATTATATGCGATTGTCAGACACA	<i>epsB</i> up
3976	CTCCT <b>CTCGAG</b> TTTTCTAAAGATCACTCGCTTCACT	<i>epsB</i> up
3977	AGGAG <b>CTCGAG</b> GAACACTATTCCTACTAGTTTTTTGTA	<i>epsB</i> down
3978	CTCCT <b>GGATCC</b> AGTCCCACCTGAGCCCCGCC	<i>epsB</i> down
3979	AGGAGGA <b>ATTCAA</b> ATCAGTTGTTGCGGAACAATATC	<i>epsC</i> up
3980	CTCCT <b>CTCGAG</b> CAGCGCAATAATCATTGACAGTCT	<i>epsC</i> up
3981	AGGAG <b>CTCGAG</b> GCTGCCAGCGTTCATTAGGGG	<i>epsC</i> down
3982	CTCCT <b>GGATCCC</b> CTCAAGCACATTCATACCGAGG	<i>epsC</i> down
1278	AGGAG <b>GGATCC</b> GGCGGCCCGGTGACAGTAACA	<i>epsD</i> up
1279	CTCCT <b>GTGAC</b> AGTCGCGCAAAACAATATCTTTTTTC	<i>epsD</i> up
1280	AGGAG <b>GTGAC</b> ATGGATATGGATACAAAGGAGAAAAG	<i>epsD</i> down
1281	CTCCT <b>GAATT</b> CGCGAAGCCACAAATCAATATCTTC	<i>epsD</i> down

1282	AGGAGGGATCCTTGGCACCGGCATGCTTGTGTTT	epsF up
1283	CTCCTGTCGACATGGAGCACGCGCTTTTGGCTGC	epsF up
1284	AGGAGGTCGACAACATCAGCACGGAAAAGGACCAT	epsF down
1285	CTCCTGAATTCCCATGTATTGCTGATGCCGTTAA	epsF down
2357	AGGAGGAATTCGTGCGCTTATGATGAAGAGATTTTAT	epsG up
2358	CTCCTCTCGAGCAAATAAATAACAATCCCCATATTGAC	epsG up
2359	AGGAGCTCGAGCCGGTCGATTCATCGGTTCTGC	epsG down
2360	CTCCTGGATCCACACCGGGTCCGCGCATATCAC	epsG down
2114	AGGAGGAATTCCAGACGTACACGCTGTTGTACGA	epsH up
2115	CTCCTCTCGAGATAAACAGCGACTAACAGACTAACC	epsH up
2116	AGGAGCTCGAGGCTGTGTATCAGCGGGTGATCG	epsH down
2117	CTCCTGGATCCTTAAGCCATTGATAGCCCTCCATC	epsH down
3983	AGGAGGGTACCTCAGAACGTGTCTCGAATCACTG	epsI up
3984	CTCCTCTCGAGCGATTGTAACGACATGTCCTCAC	epsI up
3985	AGGAGCTCGAGAGCGGTGAAGCGCAATGATCCC	epsI down
3986	CTCCTAAGCTTGCTTGATGAGAAAAGTGAGCCGC	epsI down
1286	AGGAGGGATCCCGCAGAACCGGAAGGGGTTTATA	epsJ up
1287	CTCCTGTCGACCGGGACAATAATGCTGACGAGCG	epsJ up
1288	AGGAGGTCGACCGGGACAATGAAAGGCAGTGCGAA	epsJ down
1289	CTCCTGAATTCCTGAATGGAAGTGGTGATATAAAG	epsJ down
4068	AGGAGGGTACCTAAACATGCTCACGAAACGAGATT	epsK up
4069	CTCCTCTCGAGTATCGTGAATTCATGCCTGCTTC	epsK up
4070	AGGAGCTCGAGACGGGAGTGACAGTCAGCTTTTG	epsK down
4071	CTCCTAAGCTTACCAGTTGTCAACGTACCAAACAT	epsK down
2361	AGGAGGAATTCAAATGCTTATACGGGTGCTGTCTG	epsL up
2362	CTCCTCTCGAGCAAAAAAATGGCGGCCGTCAGATC	epsL up
2363	AGGAGCTCGAGCATGTGACCGCGGAACGGTTTAC	epsL down
2364	CTCCTGGATCCCATATAGAATCGGATTGGCGGTTG	epsL down
1487	AGGAGGAATTCTACGTATCCCGCATTACAGGCTAC	epsM up
1488	CTCCTCTCGAGTATCAGCTCTCTGATCACCTTTCC	epsM up
1489	AGGAGCTCGAGGTGCTCCGGCACGCATCATTTCT	epsM down
1490	CTCTGGATCCCCGTTATCAAGTGTGAGCGTGGT	epsM down
1491	AGGAGGAATTCGAATGTCCTGAAGGGCGATCTG	epsN up
1492	CTCCTCTCGAGCTCCCTGCCGCTCATATGAGG	epsN up
1493	AGGAGCTCGAGGTCATTGAAGTGCTACTGCACTTA	epsN down
1494	CTCCTGGATCCCTTCCGCTCACATGTGAGCAGGA	epsN down
3991	AGGAGGAATTCTGAAATCGTAAGCCTGTGTGATGC	epsO up
3992	CTCCTCTCGAGTTTCTGTTTCAGGCTGATCATCGA	epsO up
3993	AGGAGCTCGAGCTCATCCAGCACGCCTCAAAGAA	epsO down
3994	CTCCTGGATCCGGCCGATACAGATAATCGCCGT	epsO down

Cassette **Boldface** characters indicate restriction sites

primers:

Complementa  
tion

Ecoli-pgaA-  
compF TTTTTTCTAGACGGATTATGAGGTGCAAAAATATCTTTCTTTTCAGTTACC

Ecoli-pgaA-  
compR TTTTTCTGCAGGAGTGAGCATAATTATACTCACCAGCATCAGG

Ecoli-pgaB- TTTTTTCTAGATTATGACGGTGACAGAGAACAACACTTATACG

compF  
Ecoli-pgaB-  
compR TTTTTCCTGCAGGTACGCATAGGGGGATGCATAACACC

Ecoli-pgaC-  
compF TTTTTGAGCTCAGTGAAAACTATGGTTATTATCCCGACAATTTTCTGC

Ecoli-pgaC-  
compR TTTTTTCTAGATTCTGGCCTCGCTTTGCCAGTAATAACCC

Ecoli-pgaD-  
compF TTTTTTCTAGATGTTGATGCCTAAAAAGCAACGCGC

Ecoli-pgaD-  
compR TTTTTCCTGCAGGCGGGCGAACC GGGCTTTGTTTTG

Bsub-epsH-  
compF TTTTTTCTAGAATGCCGGTCGATTCATCGGTTCTG

Bsub-epsH-  
compR TTTTTCCTGCAGGTTTAGCAGCAGCCATTCTGCAAAATTGATTTTCAACG

Bsub-epsI-  
compF TTTTTTCTAGAGCGAAATACAGAATGATTCCGTTTCGTCGC

Bsub-epsI-  
compR TTTTTCCTGCAGGTACATCGGGACAATAATGCTGACGAGCG

Bsub-epsJ-  
compF TTTTTTCTAGAAAAAGAAACATCCCGTGCAGGCTTTCCG

Bsub-epsJ-  
compR TTTTTCCTGCAGGAAGCCGTGAGATTCGCGCTG

Bsub-epsK-  
compF TTTTTTCTAGATTAGCACTGTGTAAATGGCGGCTCAC

Bsub-epsK-  
compR TTTTTCCTGCAGGCCGGCCTTACTTGCTTAAAGAAGA

