

Supplementary Table S3. Oligonucleotide primers used in the present study.

I. Primers used for constructing *dgcX::lacZ* and *yneF::lacZ* on pJL28¹:

<i>dgcX</i> -d-412(<i>Eco</i> RI)	5'- GCAAAGA AAT TCTGTGCCTCAGTTTTGTC-3'
<i>dgcX</i> -u-18(<i>Hind</i> III)	5'- ATAGA AAGCT TGGTTGATAATCATGTACGCACC-3'
<i>yneF</i> -d-373(<i>Eco</i> RI)	5'- GGCGGA AAT TCGTCGCTAAACGTC-3'
<i>yneF</i> -u-14(<i>Hind</i> III)	5'- GGGTA AAGCT TGCATTAATTATCTGTCCATAC-3'

II. Primers used for PCR analysis of the *pgaA-ycdT* and *dgcX* regions in the chromosome:

<i>pgaA</i> -E1(-2890)-for-	5'-CAGGGTTAAAATTTGTCC-3'
<i>ycdT</i> -E2(+1446)-rev	5'-CCTCAGTCCGGAACAATT-3'
<i>dgcX</i> -d-(-62)	5'-GTAAGGAATTCTCGTTATTTGTAGCGATACATATTA-3'
<i>dgcX</i> -u-1310	5'-AAAAGGAGATCTTTTTGAGTGAGTTATCACC-3'

III. Primers used for the construction of insertion-deletion mutations in *bcsE*, *bcsF* and *bcsG*²:

<i>bcsE</i> -H1P1-pKD13	5'-GATAAGTTTTAATTTCAATGGTAGGTTTATTTCTTAGCTTTTCGCTA GGTGTAGGCTGGAGCTGCTTC-3'
<i>bcsE</i> -H2P4-pKD13	5'-GCCAGTGGCGACCATCGTGCTCGGCATTGATGACCGGTTTAGAACTT TGCATTCCGGGGATCCGTCGACC-3'
<i>bcsF</i> -H1P1-pKD13	5'-GAAATTATTGTCGTTTGCGCACTGATATTTTCCCGCTGGGCTATCTG GCGTGTAGGCTGGAGCTGCTTC-3'
<i>bcsF</i> -H2P4-pKD13	5'-CCGTGCGGCGTAACGTCCCGGCCGGTTTAACATAACGAGGTTTAGCA AAGATTCCGGGGATCCGTCGACC-3'
<i>bcsG</i> -H1P1-pKD13	5'-GGCGCGGCCTTTCCGGCTGGAACCTTCTATTTTCTGGTTAAGTTCGGC CTGGTGTAGGCTGGAGCTGCTTC-3'
<i>bcsG</i> -H2P4-pKD13	5'-TTACTGCGGGTAAGGCACCCAGTCGCCGCCGTTTCAGGCGAACGTA CGGATTCCGGGGATCCGTCGACC-3'

IV. Primers used for cloning into *csgD* into pQE60³:

<i>csgD</i> - <i>Nco</i> I	5'-TTCATGCCAT GG TTAATGAAGTCCATAGTATTCATGGTC-3'
<i>csgD</i> - <i>Bgl</i> II	5'-AATGGA AGATCT TCGCCTGAGGTTATCGTTTGCCAGGAAACC-3'

¹ Relevant restriction sites are given in **bold**.

² Sequences complementary to pKD13 (providing the resistance cassette) are given in *italics*.

³ Relevant restriction sites are given in **bold**.