

**Table S3:** Oligonucleotides used as primers

Primer	Sequence (5'-3')	Function
1	TTTCTAGAGAATAGGAACTTCTCGACGGAGGCCT GTAGCAGTTGAGGAATTATAG	Cloning of pFuseA- <i>npt-5VNTR-clbRp-lux</i>
2	GTTGGTATGTAAGCAAAAAGTTTCCAAATTTTCAT GTTTCCCCCATCCTG	Cloning of pFuseA- <i>npt-5VNTR-clbRp-lux</i>
3	GCGCGAATTCACCATGAGTAATATCAGTTTGTAT TG	Cloning of pBAD24- <i>clbQ-rrnBt</i>
4	GCGCTCTAGAGCCATATCTATTGCTCCTTG	Cloning of pBAD24- <i>clbQ-rrnBt</i>
5	CCATCGAATGGCCAGATGAT	Cloning of pBAD24- <i>tetAp-clbQ-rrnBt</i>
6	GCGCACATGTAGCATTATCAGGGTTATTGTCT	Cloning of pBAD24- <i>tetAp-clbQ-rrnBt</i>
7	TGGAACAAACGCCAGCAACG	Cloning of pBAD24- <i>tetAp-clbQ-rrnBt</i>
8	TTCACTTTTCTCTATCACTGATAG	Cloning of pBAD24- <i>tetAp-clbQ-rrnBt</i>
9	GTGATAGAGAAAAGTGAAATGAGTAATATCAGTT TGTATTGTTTG	Cloning of pBAD24- <i>tetAp-clbQ-rrnBt</i>
10	TGCTCATGGTAGTTACTCCTTTTC	Cloning of <i>clbR</i> deletion
11	CAATCCTCATGTTTCCCCCATCCTGAATGGTATC	Cloning of <i>clbR</i> deletion
12	GGGGGGAAACATGAGGATTGATATATTAATTGGA C	Cloning of <i>clbR</i> deletion
13	CGCCATTTAAATGAAGTTCCTATTCTCAATTCTG CCCATTGACG	Cloning of <i>clbR</i> deletion
14	GCGCCACGTCGTACGTGCAAATATGGCAAACC	Cloning of <i>clbR</i> deletion
15	GTATAGGAACTTCTCAATTCTGCCATTGACG	Cloning of <i>clbR</i> deletion
16	GGGCAGAATTGAGAAGTTCCTATACTTTCTAGAG AATAG	Cloning of <i>clbR</i> deletion
17	CCTGTCTCTTGATCAGATCTTGATC	Cloning of <i>clbR</i> deletion
18	GGTGGTTGCTCTTCCGCAGATAATCTCATTCTGT TAGC	Cloning of <i>clbR</i> deletion/ pTXB1- <i>clbR</i>
19	GGTGGTCATATGGATAAGTTCAAAGAAAAAAACC	Cloning of pTXB1- <i>clbR</i>
20	TGTTTCTCCATACCCGTTTTTTTTGGGCTAGCAGGA GGAATTCACCATGGATAAGTTCAAAGAAAAAA CC	Amplification of <i>clbR</i> for cloning into pBAD24- <i>clbR</i>
21	TTCTGTTTTATCAGACCGCTTCTGCGTTCTGATTT AATCTGTATCATTAGATAATCTCATTCTGTAGC	Amplification of <i>clbR</i> for cloning into pBAD24- <i>clbR</i>
22	GCTAACAGGAATGAGATTATCTAATGATACAGAT TAAATCAGAACGC	Amplification of <i>rrnBt</i> for cloning into pBAD24- <i>clbR</i>
23	CAAAGAGTTTGTAGAAACGC	Amplification of <i>rrnBt</i> for cloning into pBAD24- <i>clbR</i>
24	TTTCGCCACCTCTGACTT	Amplification of <i>tetAp</i> for cloning into pGEM®- T Easy- <i>tetAp-clbR</i>
25	GGTTTTTTTCTTTGAACTTATCCATTTCACTTTTCT CTATCACTGATAGG	Amplification of <i>tetAp</i> for cloning into pGEM®- T Easy- <i>tetAp-clbR</i>
26	ATGGATAAGTTCAAAGAAAAAAACC	Amplification of <i>clbR-rrnBt</i> from pBAD24- <i>clbR</i>
27	AGCATTTATCAGGGTTATTGTCT	Amplification of <i>clbR-rrnBt</i> from pBAD24- <i>clbR</i>
28	CCATCGAATGGCCAGATGAT	Fusion PCR to fuse the <i>tetAp</i> and <i>clbR-rrnBt</i> amplicons

Primer	Sequence (5'-3')	Function
29	CAAAAGAGTTTGTAGAAACGC	Fusion PCR to fuse the <i>tetAp</i> and <i>clbR-rrnBt</i> amplicons
30	AATGCCATCTGGTATCACTTAAAGGTATTA CAACTTTTTGTCTTTTTACCTCATATGAATATCCT CCTTAGTTCC	Cloning of M1/5 <i>rpsLK42R</i> $\lambda$ <i>attB</i> ::FRT
31	AAGCCAATGCCAGCGCCAGACGGGAAACTGAAA ATGTGTTACAGGTTGCTCCGGGCTATGTGTAGG CTGGAGCTGCTC	Cloning of M1/5 <i>rpsLK42R</i> $\lambda$ <i>attB</i> ::FRT
32	TTCCCGTGATGGATAAATAAG	Amplification <i>frr</i> promoter region
33	GTTACGAATCCTTGAAAACCT	Amplification <i>frr</i> promoter region
34	TATGCAGCAGCGACATAAGG	Fusing <i>luxA</i> with <i>frr</i> promoter region
35	AAGTTTTCAAGGATTCGTAACATGAAATTTGGAA ACTTTTTGCTTACATACCAAC	Fusing <i>luxA</i> with <i>frr</i> promoter region
36	GACGGAAGATCACTTCGCAG	Amplification of <i>cat</i> cassette
37	ACACCAGGATTTATTTATTCTGCGAAGTGATCTTC CGTCTTCCCGTGATGGATAAATAAG	Fusing <i>luxA</i> overlap with <i>frr</i> promoter region, adding <i>cat</i> cassette overlap
38	GAGATAACGGGTTTTTTTCTTTGAACTTATCCATG TTTCCCCCATCCTGTTACGCCCCGCCCTGCCACT	Scarless VNTR cloning
39	GCCTGTAGCAGTTGAGGAATTATAGCATCAATAC CGTCATCCTGCGGGACGCTAGCTTCTTCGTCTGTT T	Scarless VNTR cloning
40	GAGATAACGGGTTTTTTTCTTTGAACTTAT	Scarless VNTR cloning
41	GCCTGTAGCAGTTGAGGAATTATAG	Scarless VNTR cloning
42	ACGAGTATCGAGATGGCACA	Control PCR for chromosomal <i>clbR</i> deletion
43	ATGGCTCTTGTATCTATC	Control PCR for chromosomal <i>clbR</i> deletion
44	GATAAGTTCAAAGAAAAAACCCTTATCTCTGC GTGAAAGACAAGTATTGCGCATGCTGTGTAGGCT GGAGCTGCTTC	Chromosomal <i>clbR</i> deletion
45	ATATGAAAATCAATATTATCGACGGCTCAGAAGT GTCTAGATTATCCGTGGCGATCATATGAATATCCT CCTTAGTTCC	Chromosomal <i>clbR</i> deletion
46	ACCGTTGTATCGTAAATTCCTC	Chromosomal <i>clbR</i> deletion
47	GATTGTATTGAATTTCAATTAAC	Chromosomal <i>clbR</i> deletion
48	TGCTCATGGTAGTTACTCCTTTTC	General primer probe generation for EMSA with <i>clbB</i> upstream region
49	CATCCTGAATGGTATCTGTGTATC	547-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 1)
50	ATCTGTGTATCTGTTGTTTTGGCAG	478-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 2)
51	TTATGTAAATGGGAATTACGC	422-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 3)
52	TGTATGTAATGCAATGACTTACATG	355-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 4)
53	AACATGTAATCAGAATATAAGG	294-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 5)
54	CAATATGATTATATGAATACGC	238-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 6)
55	ATGTAAATAATCTATAAATCC	171-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 7)

<b>Primer</b>	<b>Sequence (5'-3')</b>	<b>Function</b>
56	GAGATTTTCCATGTAATAAGATGG	115-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 8)
57	TGAATACGCAATAAAAACTATACTTG	225-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 9)
58	AAAAACTATACTTGCGGATAGGTG	212-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 10)
59	TGCGGATAGGTGTGCCATTTAG	200-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 11)
60	CCATTTAGAATAATCATGTAAATAATC	186-bp probe generation for EMSA with <i>clbB</i> upstream region (probe 12)
61	ACCGTTGTATCGTAAATTCCTC	General primer probe generation for EMSA with <i>clbR</i> upstream region
62	CTTTCACGCAGAGATAACGGG	176-bp probe generation for EMSA with <i>clbR</i> upstream region (probe 13)
63	GGTTTTTTTTCTTTGAACTTATC	157-bp probe generation for EMSA with <i>clbR</i> upstream region (probe 14)
64	CATGTTTCCCCCATCCTG	135-bp probe generation for EMSA with <i>clbR</i> upstream region (probe 15)
65	CATCCTGAATGGTATCTGTGTATC	123-bp probe generation for EMSA with <i>clbR</i> upstream region (probe 16)
66	CCGATTCATTAATGCAGCTG	229-bp probe generation for EMSA <i>lacZ</i> control
67	CCAGTCACGACGTTGTAATACG	229-bp probe generation for EMSA <i>lacZ</i> control
68	GTTGTCGCTGAAGCAACTGG	RT-qPCR <i>gapA</i>
69	AGCGTTGGAAACGATGTCCT	RT-qPCR <i>gapA</i>
70	CGTGGTGATAAAGTTGGGAC	RT-qPCR <i>clbA</i>
71	CTCCACAGGAAGCTACTAAC	RT-qPCR <i>clbA</i>
72	AGCGTGATTTCGTATTCCGAG	RT-qPCR <i>clbR</i>
73	CCGTTATCTCTGCGTGAAAG	RT-qPCR <i>clbR</i>
74	CGACGTCACACCTTTCAGCACG	RT-qPCR <i>clbB</i>
75	TGCGGGGGTGATACGATAATCG	RT-qPCR <i>clbB</i>
76	ATCGTCAAGCTGGGCGGAACTC	RT-qPCR <i>clbQ</i>
77	AGTCCGCGTCTTTCGTGTGGTG	RT-qPCR <i>clbQ</i>