

TABLE S4 Primers used in this study.

Primers for cloning/mutagenesis	Sequence (5' --> 3')
<i>pla</i> 5' -500 PstI	CTGCAGCGCCTGCTGGCTGCACTTGTCTGGTTGCCGG
<i>pla</i> (<i>gfp</i>) 3' +27	GTGAAAAGTTCTTCTCCTTTACTCATGGTTGCCACAATAGAACTTTTCTTCAT
<i>gfp</i> (<i>pla</i>) 5' 1	ATGAAGAAAAGTTCTATTGTGGCAACCATGAGTAAAGGAGAAGAAGTTTTCAC
<i>gfp</i> 3'717 Bam	CGGGATCCTTATTTGTATAGTTCATCCATGCCATGTG
P _{tet} -405 5' Pst	AACTGCAGCTTTCACCAGCGTTTCTGGGGT
P _{tet} (<i>pla</i> 5' UTR) 3' -tx	CTCTCTGCATGAACGAAAACATATTATTTGTATCTGATTATCTAATTTTCATTTTTAATAATGTGTGCTCAGTATCTCTATCACTGATA
<i>gfp</i> (<i>pla</i> 5' UTR) 5' 1	CAAATAATATGTTTTCGTTCATGCAGAGAGATTAAGGGTGTCTAATGAAGAAAAGTTCTATTGTGGCAACCATGAGTAAAGGAGAA GAAC
<i>gfp</i> - <i>crp</i> 5'	ATGGTTCTCGGTAAGCCACAAACAGACATGAGTAAAGGAGAAGAAGTTTTCAC
<i>crp</i> 5' UTR- <i>gfp</i> 3'	GTGAAAAGTTCTTCTCCTTTACTCATGTCTGTTTGTGGCTTACCGAGAACCAT
P _{tet} - <i>crp</i> (-tx) 3'	CCCTGTACGGCGTGCTTATAGAATACAGTGCTCAGTATCTCTATCACTGATA
<i>crp</i> 5' -496 PstI	CTGCAGGACACGACATCAATGGCGCTACACCCCCCGC
<i>crp</i> 5' 128 SpeI	GACTAGTCGTGAAAGGCTCCGTTGCGGTGCTG
<i>crp</i> 3' 630 HA	AGCGTAATCTGGAACATCGTATGGGTAAGCGTAATCTGGAACATCGTATGGGTAACGGGTGCCGTAACGACGATCGT
<i>crp</i> 5' 631 HA	TACCACATACGATGTTCCAGATTACGCTTACCACATACGATGTTCCAGATTACGCTTAATTCCCTCTAAAACCGGCGTTAAAGC
<i>crp</i> 3' +502 SpeI	GACTAGTCGCCATACCAAAGCGTCCCCACGGC
<i>crp</i> 5' 1 (P _{tet})	CGAATTCATTAAGAGGAGAAAGGTACCCATGGTTCTCGGTAAGCCACAAACAGAC
<i>crp</i> -HA 3' 633 BamHI	GGATCCTTAAGCGTAATCTGGAACATCGTATGGGTA
P _{tet} 3' 1 (<i>crp</i>)	GTCTGTTTGTGGCTTACCGAGAACCATGGGTACCTTTCTCCTCTTTAATGAATTCCG
<i>crp</i> 3' +500	AGAGCAAGGGTGCCTGCCAGATAG
P4 <i>crp</i> 5' 632	GGTCGACGGATCCCCGGAATAATTCCCTCTAAAACCGGCGTTAAAGCTG
<i>crp</i> 5' -500	AATAGACACGACATCAATGGCGCTAC
P1 <i>crp</i> 3'3	GAAGCAGCTCCAGCCTACACCATTGCTGTTATCCTCTGTTGTTATCGCTG
Primers for qRT-PCR	Sequence (5' --> 3')
<i>pla</i> 5' 397	GACCTCAATGTGAAAGGCTGGTTACACC
<i>pla</i> 3' 498	ACCACCTGTAGCTGTCCAACCTGAAAC
<i>crp</i> 5' 2	TGGTTCTCGGTAAGCCACAAACAGAC
<i>crp</i> 3' 146	GCAACGGAGCCTTTCACGATGTAG
<i>gyrB</i> .r.3'	ATTGGTAAAGGTCTGGAAACTTGGCC
<i>gyrB</i> .f.5'	TCGCCGTGAAGGTAAAGTTC
<i>proS</i> 5' 745	ACAGAAGAGCTGCGAATCGTTGACAC
<i>proS</i> 3' 887	AGCGCAACCAACTTATGACCACTCTC
16S rRNA 5' 993	TAAAGTGCCTTCGGGAACTGTGAG
16S rRNA 3' 1109	ACGTGCTGGCAACAAAGGATAAGG
Primers for RACE	Sequence (5' --> 3')
<i>crp</i> 5' 464	CTTCCTCATCTTTGATCAGCAC
<i>crp</i> 3' 444	GCAAAGAGATGATCCTCTCC