

Table S1. Oligonucleotides used in this study

Primer	5'-3' sequence	Use
Strains construction		
Δ gcvB ::tetfor	GTCCGTTGAGCTTCTACCAGCAAATACCTATAGTGGCGGCACctagacatcattaattccta	Construction <i>ΔgcvB</i> :: <i>tet</i> allele
Δ gcvB ::tetrev	CGATCGCAAGGTAAAAAAAAGCACCGCAATTAGGCGGTGCTgaagctaaatcttctttatcg	
Δ gcvB ::kanfor	GTCCGTTGAGCTTCTACCAGCAAATACCTATAGTGGCGGCACaaagccacgttggtctcaa	Construction <i>ΔgcvB</i> :: <i>kan</i> allele
Δ gcvB ::kanrev	CGATCGCAAGGTAAAAAAAAGCACCGCAATTAGGCGGTGCTttagaaaaactcatcgagc	
5'Ptet-phoP	GATAGAGATTGACATCCCTATCAGTGATAGAGATACTGAGCACacactattttaataattaag acaggg	Construction Ptet- <i>phoP</i> - <i>lacZ</i> fusion
3'phoP-lacZ	TAACGCCAGGGTTTTCCCAGTCACGACGTTGTAAAACGACcgcattgtcttcaacaaccag	
phoPmutR3for	ACCTGACGCTTTTTATCGCAACTCTCTACTGTTTCTCCATacactattttaataattaactgctcgga gaaataaaaatgcgcgtactgg	Construction P _{BAD} - <i>phoPmutR3</i> - <i>lacZ</i> fusion
5'P1mgtA	GCGAAGCGGCATGCATTTACGTTGACACCATCGAATGGCGCgtgcattcagcaatgggtaaa gtc	Construction P1- <i>mgtA</i> - <i>lacZ</i> fusion
3'mgtA-lacZ	GTAACGCCAGGGTTTTCCCAGTCACGACGTTGTAAAACGACaatgagccgggtaaaaatttctt	
5'LivJ-lac	ACCTGACGCTTTTTATCGCAACTCTCTACTGTTTCTCCATagagtatgctgctaaagcacg	Construction P _{BAD} - <i>livJ</i> - <i>lacZ</i> fusion
3'LivJ-lac	TAACGCCAGGGTTTTCCCAGTCACGACGTTGTAAAACGACtatgttcatcctgagaatccc	
5'Cm-P _{BAD} - phoPQ	AAGATTATCCGCTTTTTATTTTTTCACTTTACCTCCCCTCCgtgtaggctggagctgct	Construction CmR-P _{BAD} - <i>phoPQ</i> allele
3'Cm-P _{BAD} - phoPQ	GCATTTTTATTCTCCCTGTCTTAATTATTTAAATAGTGTatggagaaacagtagagagt	
Plasmid construction		
pBR-for	ggaaaacgttcttcggggcg	
5'Hind-MicA- tet	cgtacaagcttatcgatgataagctgtcaaacatgagaattccacgcctgacagaaaagaaaaaggc	
GcvBmutR1 for	ggcttacggttgatgtgacaacatgtgttgcaattggctgctgcatc	
GcvBmutR1 rev	gaatcgagaccaattgcaaacacatggtgtcaacatcacaaccgtaagcc	

GcvBmutR3for	ttttcaattcctgtacatttacc gacagt gtccatagtgattaatgtagc	
GcvBmutR3rev	gctacattaatcactatggac actgtc ggtaaagtacagggaagtgaaaa	
Probes for Northern-Blot		
MgrR-probe	5'Bio-cagtaaaccggcggtgaatgcttgcattgatagatttg	
ompT-probe	5'Bio-gccgatatcatctctgaatccctcctcagaactgtagatatagg	
SsrA-probe	5'Bio-cgccactaacaactagcctgattaagttttaacgcttca	
GcvB-probe	5'Bio-cacaaccgtaagcctaaagttcaccagaacacgcattccg	
phoP-probe	5'Bio-gcatcttctgcgtcatcgacctgatgaccagcatctgaatc	
In vitro experiments		
5'T7MicA	<u>taatac</u> gactcactataggaaagacgcgcatttg	
5'T7MicAmut	<u>taatac</u> gactcactataggaaag gccc atttg	
3'T7MicA	aaaaggccactcgtgagtgg	
5'T7GcvB	<u>taatac</u> gactcactataggacttctgagccggaacg	
3'T7GcvB	ggtaaaaaaaagcaccgcaattaggcgg	
5'T7phoP	<u>taatac</u> gactcactataggacactattttaataattaagacaggg	
3'T7phoP	cgtctggcaatccgagatc	
phoP-Cy5-probe#1	5'Cy5-ccagcatcctgaatctgaac	
phoP-Cy5-probe#2	5'Cy5-cttctttggcatcttctgcg	

Nts in upper cases correspond to homology regions for recombineering

Underlined nts indicate restriction sites used for cloning or T7 promoter sequence, and bold nts mutations