

Supplemental Information for:

**Rcs phosphorelay activation in cardiolipin-deficient *Escherichia coli* reduces biofilm
formation**

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Running title: Biofilm reduction in cardiolipin-deficient *E. coli*

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Supplemental Methods

Determination of fim invertible element orientation. We grew *E. coli* cells in 96 well microplates and used sonication to harvest biofilm cells as described above. We used an Epicentre Master Pure Complete DNA and RNA Purification kit (Madison, WI, USA) to extract genomic DNA from cells. Multiplex PCR was performed using New England Biolabs Q5 High-Fidelity DNA Polymerase (Ipswich, MA, USA), following the manufacturer's instructions. Primers were included to amplify off-oriented *fimAp* (INV and FIME), on-oriented *fimAp* (INV and FIMA), and *ftsZ* (EcFtsZ 1 and EcFtsZ 2; loading control) (1, 2).

Tables

Table S1. Strains of *E. coli* used in this study

Strain name	Relevant genotype	Reference
MG1655	Wild-type <i>E. coli</i>	Laboratory strain
PO10	MG1655 Δ <i>clsABC</i> ::FRT	(3)
Δ <i>clsA</i>	MG1655 Δ <i>clsA</i> ::FRT	This work
Δ <i>clsB</i>	MG1655 Δ <i>clsB</i> ::FRT	This work
Δ <i>clsC</i>	MG1655 Δ <i>clsC</i> ::FRT	This work
Δ <i>clsBC</i>	MG1655 Δ <i>clsBC</i> ::FRT-kan-FRT	This work
Δ <i>clsAC</i>	MG1655 Δ <i>clsAC</i> ::FRT-kan-FRT	This work
Δ <i>clsAB</i>	MG1655 Δ <i>clsAB</i> ::FRT-kan-FRT	This work
Δ <i>flhD</i>	BW25113 Δ <i>flhD</i> ::FRT-kan-FRT	(4)
Δ <i>cls</i> Δ <i>rcaA</i>	MG1655 Δ <i>clsABC</i> ::FRT Δ <i>rcaA</i> ::FRT-cam-FRT	This work
Δ <i>cls</i> Δ <i>rcaC</i>	MG1655 Δ <i>clsABC</i> ::FRT Δ <i>rcaC</i> ::FRT-cam-FRT	This work
Δ <i>cls</i> Δ <i>rcaF</i>	MG1655 Δ <i>clsABC</i> ::FRT Δ <i>rcaF</i> ::FRT-cam-FRT	This work
DH5 α	F ⁻ <i>endA1 glnV44 thi-1 recA1 relA1 gyrA96 deoR nupG</i> Φ 80 <i>lacZ</i> Δ M15 Δ (<i>lacZYA-argF</i>)U169, <i>hsdR17</i> (r κ ⁻ m κ ⁺), λ ⁻	Laboratory strain
5-alpha F' I ^q	F' <i>proA</i> ⁺ <i>B</i> ⁺ <i>lacI</i> ^q Δ (<i>lacZ</i>)M15 <i>zzf</i> ::Tn10 (Tet ^R) / <i>fhuA2</i> Δ (<i>argF-lacZ</i>)U169 <i>phoA glnV44</i> Φ 80 Δ (<i>lacZ</i>)M15 <i>gyrA96 recA1 relA1 endA1 thi-1 hsdR17</i>	New England Biolabs

Table S2. Plasmids used in this study

Plasmid	Relevant genotype	Reference
pMP4641	TetR	(5)
pMP4658	TetR	(5)
pBad33	Low copy arabinose inducible expression plasmid	(6)
pBad33- <i>clsA</i>	<i>clsA</i> inserted between XbaI and SphI sites; CamR	This work
pBad33- <i>clsB</i>	<i>clsB</i> inserted between XbaI and HindIII sites; CamR	This work
pBad33- <i>ymdBC</i>	<i>ymdB-clsC</i> inserted between XbaI and HindIII sites; CamR	This work
pBad33- <i>clsA</i> (H224F H404F)	<i>clsA</i> with H224F and H404F substitutions	This work
pBad33- <i>clsB</i> (H113A)	<i>clsB</i> with H113A substitution	This work
pBad33- <i>clsB</i> (H290A)	<i>clsB</i> with H290A substitution	This work
pBad33- <i>ymdB-ymdC</i> (H130A)	<i>clsC</i> with H130A substitution, <i>ymdB</i> wild-type	This work
pBad33- <i>ymdB-ymdC</i> (H369A)	<i>clsC</i> with H369A substitution, <i>ymdB</i> wild-type	This work
pBad33- <i>phoA</i>	<i>phoA</i> inserted into X site; CamR	This work
pCP20	FLP recombinase expression plasmid; AmpR CamR; temperature sensitive	(7)
pKD3	AmpR CamR	(7)
pKD46	AmpR; temperature sensitive	(7)
pBad33- <i>ompA-6XHis</i>	CamR	This work

Table S3. Primers

Name	Sequence (5'→3')
INV ¹	GAGGTGATGTGAAATTAATTTAC
FIMA ¹	GATGCGGTACGAACCTGTCC
FIME ¹	GCAGGCGGTTTGTACGGGG
EcFtsZ 1 ²	TAGCGGTATCACCAAAGGACT
EcFtsZ 2 ²	GTGATCAGAGAGTTCACATGCT

¹Schwan *et al.* (1992), *J Bacteriol*

²Schwan *et al.* (2007), *J Bacteriol*

qPCR		
Gene	Forward (5'→3')	Reverse (5'→3')
clsA	ATATAATCGTCCTGAACGGCGGC	TGGTATGCAGTAACCCGCCTTCA
clsB	TAAAAAGTGGAACGCCAGCACG	AGTGACAAACTGAGCGGATCGAGA
clsC	CCGAAATGGCAGAGCATACGCTC	CGCTTGCTTGACAGTCATCCACG
cpsB	GTTGGCTCCTGGTCTTCATTA	CAGGCCAGATTCAGCATACA
cpsG	TGACGTGCTGGATATTGGTATG	GATTATGGCTGGCGGTAAC
degP	GCCTTCAGTGGTCAGCATT	GGCAGAACGGAGAATCATCA
htrA	TATCGCGCTGATCCAAATCC	ACCAATCGCTACGGTGTAATC
mdtA	CAAGCAGGTTGATGTTGGTAAC	CCGGCAGGGTAAAGACTAAAT
acrD	TGACCTCGCTGGCATTATC	CGAAATCATCCCGCCATTA
pspA	CTGATGATCCAGGAGATGGAAG	TGTTCAATACGGCGAGTCAG
pspC	CTGGTGGTGCTGTCGATTT	AAGGTAGCTGCTCACCAAAG
idnT ³	CTGTTTAGCGAAGAGGAGATGC	ACAAACGGCGGCGATAGC
gapA ⁴	GTGATCCGGCTAACCTGAAA	GTCTGGCCAGCATATTTGT

³Zhou *et al.* (2011), *BMC Mol Biol*

⁴Uchiyama *et al.* (2010), *FEMS Microbiol Lett*

Table S4. Minimum inhibitory concentration (MIC) of antimicrobial compounds in *E. coli* grown at 30°C in M9 minimal medium supplemented with amino acids. Values shown are the averages of 3 independent biological replicates and the standard error of the mean.

Compound	MG1655			<i>ΔclsABC</i>		
Cecropin A ^a	12.5	±	0	4.1	±	1.1
Ampicillin ^a	16.0	±	2.7	13.3	±	2.7
Kanamycin ^a	16.0	±	5.3	10.7	±	2.7
EDTA ^a	64.0	±	10.7	16.0	±	0
Polymyxin B ^a	8.7	±	4.1	4.0	±	0
Cefuroxime ^a	32.0	±	0	8.0	±	4.0

^a Values in µg/mL

Supplemental Figures

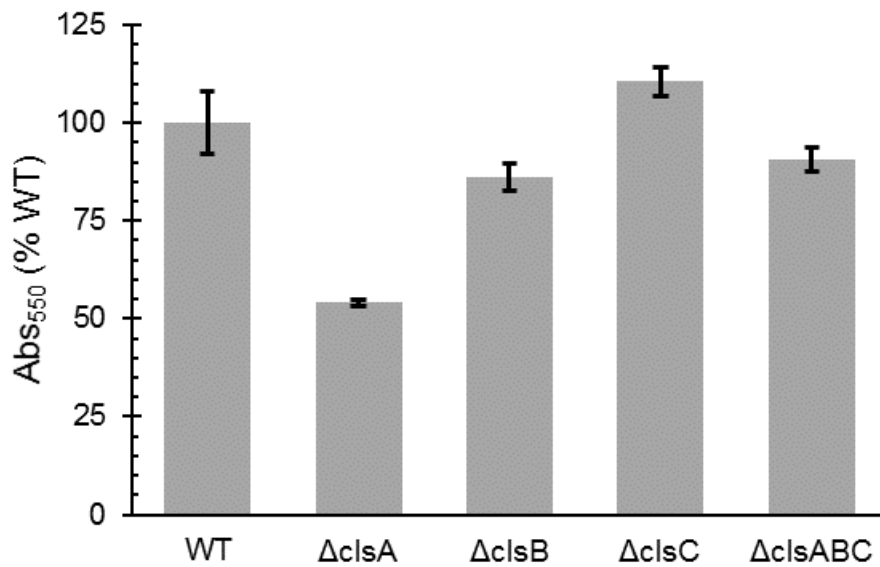


Figure S1. Surface attachment in rich nutrient medium. Cells were grown in lysogeny broth (LB) in microtiter plates for 24 h at 30°C without shaking. Adherent cells were stained with crystal violet (CV), and CV absorbance was measured at 550 nm. Error bars indicate standard error (N ≥ 6); differences between strains were not significant based on a Student's *t* test ($p > 0.05$), except WT vs. *ΔclsA* ($p < 0.0001$).

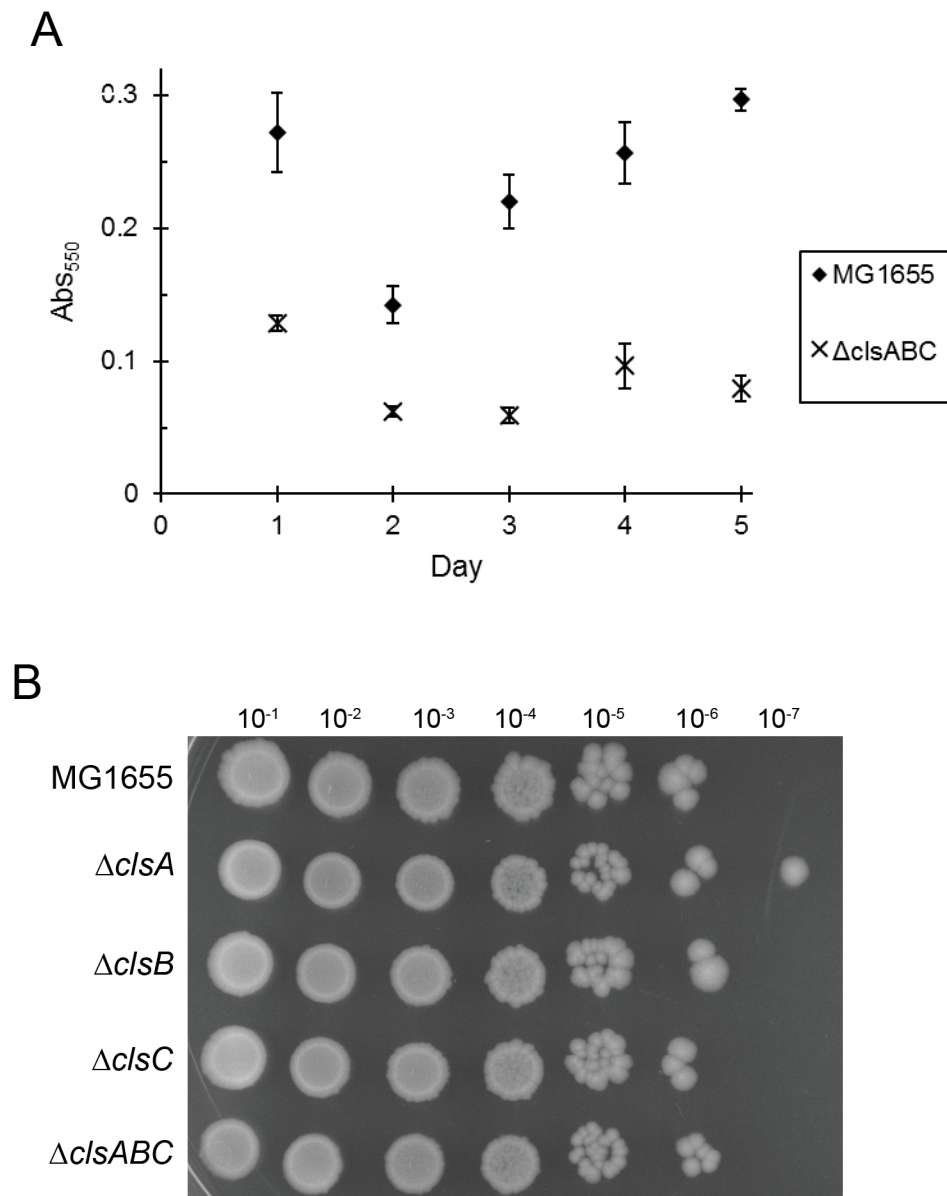


Figure S2. Surface attachment over multiple days. Cells were grown in microtiter plates for 24 h at 30°C without shaking. (A) Adherent cells were stained with crystal violet (CV), and CV absorbance was measured at 550 nm. Error bars indicate standard error (N ≥ 6). (B) Colony forming units were determined for 5 day-old cultures.

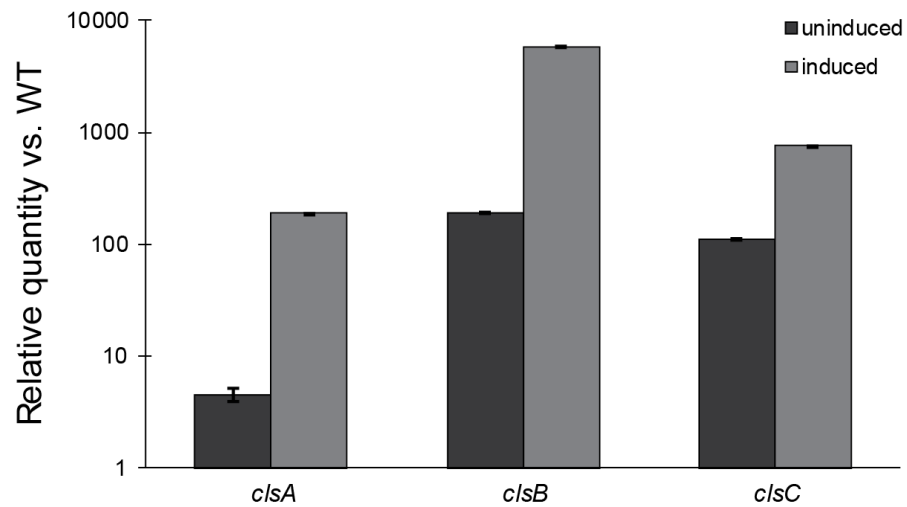


Figure S3. Induction level of *cls* transcripts. We extracted RNA from 24 h old cultures grown with (induced) or without (uninduced) 0.2% arabinose to stimulate protein expression. All strains lacked a chromosomal copy of one *cls* gene, which was replaced with a vector copy of the gene in the arabinose-inducible plasmid pBad33 (e.g. $\Delta clsA/pBad33-clsA$).

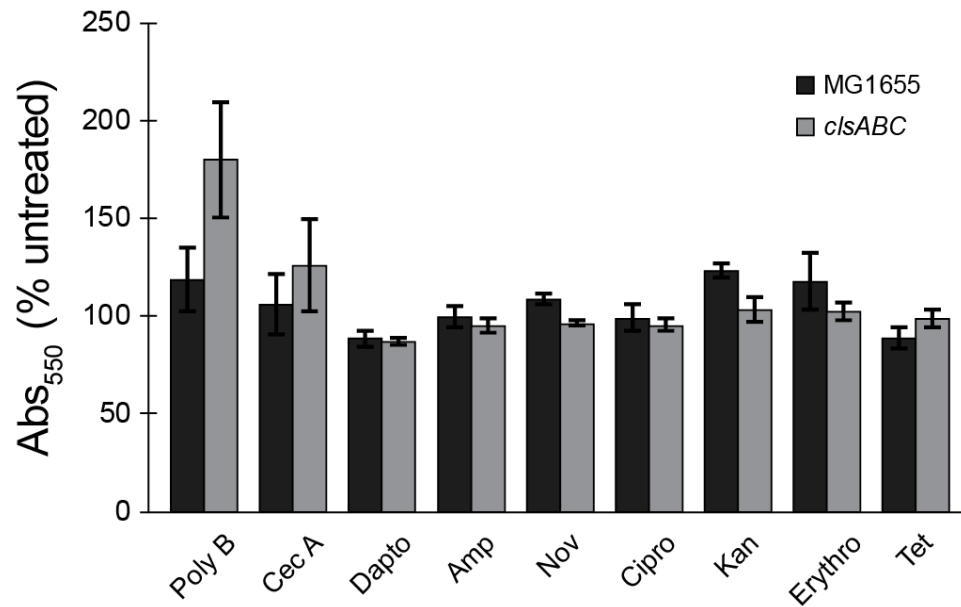


Figure S4. Effect of treatment with antimicrobials on biofilm formation in $\Delta clsABC$

cells. Cells were grown in M9 minimal medium in microtiter plates for 24 h at 37°C without shaking, then treated with antimicrobial compounds for 24 h as described in the Materials and Methods. Adherent cells were stained with CV, and CV absorbance was measured at 550 nm. Abs values were normalized to those obtained from untreated controls. Error bars indicate standard error (N ≥ 16); (Poly B, polymyxin B; Cec A, cecropin A; Dapto, daptomycin; Amp, ampicillin; Nov, novobiocin; Cipro, ciprofloxacin; Kan, kanamycin; Erythro, erythromycin; Tet, tetracycline).

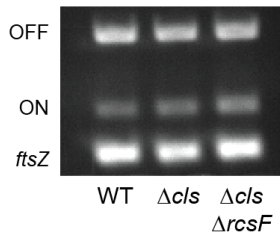
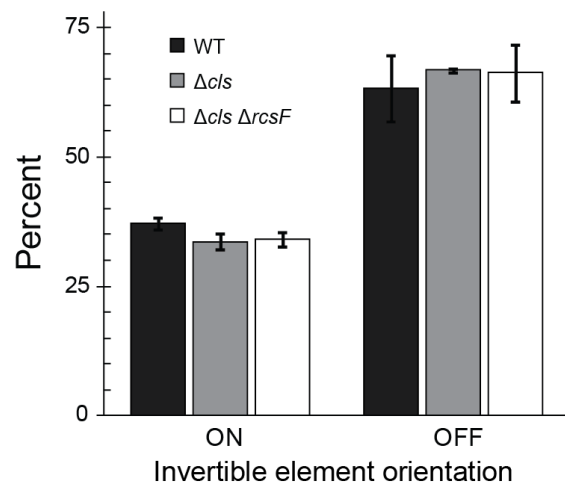
A**B**

Figure S5. Fimbriation of CL deficient cells. Genomic DNA was extracted from *E. coli* biofilms as described in the Materials and Methods and PCR was used to determine the orientation of the FimA invertible element. (A) INV and FIMA primers amplify phase-on oriented DNA (450 bp product), INV and FIME primers amplify phase-off oriented DNA (750 bp product), and EcFtsZ1 and 2 primers amplify a 302 bp segment of the *ftsZ* gene. (B) ImageJ was used to determine the intensity of DNA bands, which were normalized to the *ftsZ* loading control. Error bars indicate standard error (N = 3); differences between strains were not significant based on a Student's *t* test ($p > 0.05$).

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