

Supplementary Table 1. *E. coli* strains and plasmids used in this study.

Strains	Relevant genotype or phenotype ^a	Reference
W3110	W3110 derivative with functional RpoS (<i>rpoS396(Am)</i>)	(Serra et al., 2013a)
AR3110	W3110 derivative with functional cellulose synthase <i>bcsQ</i>	(Serra et al., 2013b)
VM388	W3110 $\Delta fliC$ Km ^s	This work
VM392	W3110 $\Delta csgA::Km^r$	This work
VM335	AR3110 $\Delta fliC::Km^r$	This work
VM661	AR3110 $\Delta csgA$ Km ^s	This work
VM391	W3110 $\Delta pgaC$ Km ^s	This work
VM482	W3110 $\Delta wcaF::Km^r$	This work
VS824	W3110 Δflu Km ^s	This work
OB301	W3110 $\Delta fimA::Km^r$	This work
OB148	W3110 $\Delta sfmH::Km^r$	This work
OB153	W3110 $\Delta ybgP::Km^r$	This work
OB154	W3110 $\Delta yraH::Km^r$	This work
OB156	W3110 $\Delta yehD::Km^r$	This work

OB166	W3110 Δ ypjA::Km ^r	This work
OB135	W3110 <i>fliC::fliC-RBS-mCherry codon optimized</i> Km ^s	This work
OB136	W3110 <i>csgA::csgA-RBS-sfGFP codon optimized</i> Km ^s	This work
OB53	W3110 <i>csgA::csgA-RBS-sfGFP</i> Km ^s , <i>fliC::fliC-RBS-mCherry opt</i> Km ^r	This work
OB54	W3110 <i>csgA::csgA-RBS-sfGFP</i> Km ^s , <i>osmY-mCherry</i> Km ^r	This work
OB55	W3110 <i>csgA::csgA-RBS-sfGFP</i> Km ^s , <i>rplL::rplL-RBS-mCherry opt</i> Km ^r	This work
OB191	AR3110 <i>csgA::csgA-RBS-sfGFP codon optimized</i> Km ^r	This work

Plasmids	Relevant genotype or phenotype ^a	Reference
pTrc99a	Amp ^r ; Expression vector; pBR ori; pTrc promoter; IPTG-inducible	(Amann et al., 1988)
pVM42	Amp ^r ; <i>eGFP</i> in pTrc99a, IPTG-inducible	This work
pOB2	Amp ^r ; <i>mCherry</i> in pTrc99a, IPTG-inducible	This work
pOB44	Amp ^r ; TIMER ^{bac} in pBR322	(Claudi et al., 2014)
pCP20	Amp ^r ; provides <i>flp</i> recombinase required for Km ^r removal	(Cherepanov and Wackernagel, 1995)

^a Amp^r and Km^r indicate ampicillin and kanamycin resistance, respectively; Km^s indicate kanamycin sensitivity.

Supplementary references:

Amann, E., Ochs, B., and Abel, K. J. (1988). Tightly regulated tac promoter vectors useful for the expression of unfused and fused proteins in *Escherichia coli*. *Gene* 69, 301–315.

Cherepanov, P. P., and Wackernagel, W. (1995). Gene disruption in *Escherichia coli*: TcR and KmR cassettes with the option of FLP-catalyzed excision of the antibiotic-resistance determinant. *Gene* 158, 9–14.

Claudi, B., Spröte, P., Chirkova, A., Personnic, N., Zankl, J., Schürmann, N., Schmidt, A., and Bumann, D. (2014). Phenotypic Variation of *Salmonella* in Host Tissues Delays Eradication by Antimicrobial Chemotherapy. *Cell* 158, 722–733.

Serra, D. O., Richter, A. M., Klauck, G., Mika, F., and Hengge, R. (2013a). Microanatomy at cellular resolution and spatial order of physiological differentiation in a bacterial biofilm. *MBio* 4. e103-e113. doi: 10.1128/mBio.00103-13

Serra, D. O., Richter, A. M., and Hengge, R. (2013b). Cellulose as an architectural element in spatially structured *Escherichia coli* biofilms. *J. Bacteriol.* 195, 5540–5554.