

**Table S1.** Strains and plasmids used in this study.

Strain or Plasmid	Genotype	Source or Reference
Strains		
<i>S. Typhimurium</i> SL1344	Wild-type strain	ATCC (W Köster, U of S)
<i>S. Enteritidis</i> ATCC 4931	Wild-type strain	ATCC (D Korber, U of S)
<i>S. Enteritidis</i> 301 (Sal18)	Wild-type strain	W Köster [1]
<i>S. Typhi</i> E02-2759	Wild-type strain	G. Dougan [2]
<i>S. Typhi</i> E03-9804	Wild-type strain	G. Dougan [2]
<i>S. Typhi</i> ISP03-07467	Wild-type strain	G. Dougan [2]
<i>S. Typhi</i> ISP04-06979	Wild-type strain	G. Dougan [2]
<i>S. Typhi</i> 8(04)N	Wild-type strain	G. Dougan [2]
<i>S. Typhi</i> CT18	Wild-type strain	G. Dougan [3]
<i>S. Enteritidis</i> D7795	Wild-type strain	G. Dougan [4]
-47 T>C	<i>PcsgD</i> from 14028	This study
-47 T>T	Native <i>PcsgD</i> <sup>a</sup>	This study
<i>S. Typhimurium</i> D23580	Wild-type strain	G. Dougan [5]
-80 A>C	<i>PcsgD</i> SNP from 14028	This study
-80 A>C, -189 A>G	<i>PcsgD</i> SNPs from 14028	This study
<i>S. Typhimurium</i> 14028	Wild-type strain	ATCC
-47 C>T	<i>PcsgD</i> from D7795	This study
-47 C>C	Native <i>PcsgD</i> <sup>a</sup>	This study
<i>bcsG</i> <sub>(D23580)</sub>	<i>bcsG</i> from D23580	This study
<i>bcsG</i> <sub>(14028)</sub>	Native <i>bcsG</i> <sup>a</sup>	This study
<i>csgD</i> <sub>(CT18)</sub>	<i>csgD</i> from CT18	This study
<i>csgD</i> <sub>(14028)</sub>	Native <i>csgD</i> <sup>a</sup>	This study
Plasmids		
pCS26, pU220	Bacterial luciferase	[6]
pCS26- <i>csgB</i>	<i>csgBAC</i> promoter	[7]
pU220- <i>csgD</i>	<i>csgDEFG</i> promoter	[7]
pCS26-sig38H4	Synthetic, RpoS-responsive promoter	[7]
pCS26- <i>adrA</i>	<i>adrA</i> promoter	[7]
pCS26- <i>cpxR</i>	<i>cpxR</i> promoter	This study
pFLAG-CTC		Sigma-Aldrich
p3xFLAG	Modified to have 3xFLAG at C-terminus	This study
p3xFLAG/ <i>csgD</i> <sup>CT18</sup>	<i>csgD</i> from <i>S. Typhi</i> CT18	This study
p3xFLAG/ <i>csgD</i> <sup>14028</sup>	<i>csgD</i> from <i>S. Typhimurium</i> 14028	This study

<sup>a</sup> These strains were selected at the end of the genome engineering procedure, and possessed promoter sequences or alleles that were the same as the parent strain

## SUPPLEMENTARY REFERENCES

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