

Table S1. Bacterial strains, plasmids, and oligonucleotides used in this study.

(A) Bacterial strains and plasmids

Name	Description	Reference
<i>S. enteric</i> serovar Typhimurium		
14028s	wild-type	Fields et al. (1986)
SM101	$\Delta phoU$	Choi et al. (2022)
SM323	<i>phoU</i> ^{Ala 147 Glu}	Choi et al. (2022)
SM108	pBAD33- <i>phoU</i> -His, pTGFP	This study
SM109	pBAD33- <i>phoU</i> -His, pTGFP- <i>phoR</i>	This study
SM343	pBAD33- <i>phoU</i> ^{Ala 147 Glu} -His, pTGFP- <i>phoR</i>	This study
SM454	<i>phoR</i> -8×Myc	Choi et al. (2022)
SM443	<i>phoU</i> ^{Ala 147 Glu} , <i>phoR</i> -8×myc::Km ^R	This study
SM457	<i>phoU</i> ^{Ala 147 Glu} , <i>phoR</i> -8×myc	This study
<i>Escherichia coli</i>		
DH5α	<i>fhuA2 lac(del)U169 phoA glnV44 Φ80' lacZ(del)M15 gyrA96 recA1 relA1 endA1 thi-1 hsdR17.</i>	Taylor, Walker et al. (1993)
SM329	DH5α/pBAD33- <i>phoU</i> ^{Ala 147 Glu} -His	This study
BTH101	<i>F⁻, cya-854, recA1, endA1, gyrA96 (Nal^r), thi1, hsdR17, spoT1, rfbD1, glnV44(AS)</i>	Karimova, et al. (1998)
LJ18	BTH101/pUT18- <i>mgtC</i> , pKT25- <i>mgtR</i>	Choi et al. (2019)
LJ27	BTH101/pUT18- <i>mgtC</i> , pKT25	Choi et al. (2019)
SM089	BTH101/pUT18- <i>phoU</i> , pKT25- <i>phoR</i>	Choi et al. (2022)
SM314	BTH101/pUT18c- <i>phoU</i> ^{Ala 147 Glu} , pKT25- <i>phoR</i>	Choi et al. (2022)
SM315	BTH101/pUT18c- <i>phoU</i> ^{Ala 147 Lys} , pKT25- <i>phoR</i>	Choi et al. (2022)
SM311	BTH101/pUT18c- <i>phoU</i> ^{Arg 148 Ala} , pKT25- <i>phoR</i>	Choi et al. (2022)
SM210	BTH101/pUT18- <i>phoU</i> , pKT25- <i>pstB</i>	This study
SM326	BTH101/pUT18c- <i>phoU</i> ^{Ala 147 Glu} , pKT25- <i>pstB</i>	This study
SM327	BTH101/pUT18c- <i>phoU</i> ^{Ala 147 Lys} , pKT25- <i>pstB</i>	This study
SM328	BTH101/pUT18c- <i>phoU</i> ^{Arg 148 Ala} , pKT25- <i>pstB</i>	This study
Plasmids		
pBAD33	pACYC184 <i>ori</i> Cm ^R	Guzman et al. (1995)
pBOP508	repR _{R6K} Ap ^R 8×myc FRT Km ^R FRT	Cho et al. (2006)
pKD46	rep _{pSC101} ^{ts} Ap ^R P _{araBAD} γ β <i>exo</i>	Datsenko and Wanner (2000)
pCP20	rep _{pSC101} ^{ts} Ap ^R Cm ^R <i>cl857</i> λP _R <i>flp</i>	Datsenko and Wanner (2000)
pUT18	p _{lac} ColEI <i>ori</i> Ap ^R	Karimova, et al. (2001)
pUT18c	p _{lac} ColEI <i>ori</i> Ap ^R	Karimova, et al. (2001)

pKT25	p _{lac} p15Aori Km ^R	Karimova, et al. (2001)
pTGFP	ColE1 <i>ori</i> Ap ^R 'gfp	Lee and Groisman (2012)

(B) Primers used in this study.

Name	Sequence (from 5' to 3')*	
Knockout, deletion, tagging		
KU452	TGA GCT GGA TAA GCT GCT GGC GGG GAA AGA TCC GAAAGA GAT CGG ATC CAG AAT TCG TGA T	<i>phoR</i> -8×Myc Km ^R insertion
KU453	GGC CCG GTA AGC GCA GCG CCA CCG GGC AAA AGA ATG AGA TGA GCT CGA TCC GTC GAC C	<i>phoR</i> -8×Myc Km ^R insertion
Cloning		
KHU767	CCC AAG CTT TTA GTG GTG ATG GTG ATG ATG CTC TTT CGG ATC TTT CCC CG	<i>phoU</i> -His-HindIII-R
KHU768	GCT CTA GAT AAT TCA GGA GTG CGT AAT GGA CAG TCT GAA CCT TAA	<i>phoU</i> -XbaI-F
For qRT-PCR		
KHQ015	CGCTGGGAAGCTGAGTTTG	<i>phoE</i> qPCR-F
KHQ016	CCAGGCCTCAACATCGTACA	<i>phoE</i> qPCR-R
KHQ097	CATATTTCCGGCCAGTTCAAC	<i>phoU</i> qPCR-F
KHQ098	CGGTGATAGCGTCAGAAAG	<i>phoU</i> qPCR-R
6970	CCAGCAGCCGCGGTAAT	<i>rrsH</i> qPCR -F
6971	TTTACGCCCAGTAATTCCGATT	<i>rrsH</i> qPCR -R

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