

**Table S1.** Bacterial strains and plasmids used in this study

<b>Strain or plasmid</b>	<b>Relevant characteristics</b>	<b>Source</b>
<b><i>E. coli</i></b>		
DH5 $\alpha$	<i>F</i> <sup>-</sup> $\varphi 80lacZ \Delta M15 \Delta(lacZYA-argF)U169 recA1 endA1 hsdR17(rk^-, mk^+)phoA supE44 thi-1 gyrA96 relA1 tonA$	Stratagene
<b><i>P. aeruginosa</i></b>		
PAO1	Wild type	This lab
<i>tspR</i> ::Tn	PA4857 transposon mutagenesis mutant	This study
<i>tspR</i> ::Gm	<i>tspR</i> (PA4857) insertion mutant of PAO1; Gm <sup>r</sup>	This study
<i>tspR</i> ::Tc	<i>tspR</i> (PA4857) insertion mutant of PAO1; Tc <sup>r</sup>	This study
<i>retS</i> ::Gm	<i>retS</i> insertion mutant of PAO1; Gm <sup>r</sup>	This study
<i>exsA</i> ::Gm	<i>exsA</i> insertion mutant of PAO1; Gm <sup>r</sup>	This study
<i>rsmY</i> ::Gm	<i>rsmY</i> insertion mutant of PAO1; Gm <sup>r</sup>	This study
<i>rsmZ</i> ::Gm	<i>rsmZ</i> insertion mutant of PAO1; Gm <sup>r</sup>	This study
<i>rsmY</i> ::Gm/ <i>tspR</i> ::Tc	<i>tspR</i> and <i>rsmY</i> double mutant of PAO1; Gm <sup>r</sup> , Tc <sup>r</sup>	This study
<i>rsmZ</i> ::Gm/ <i>tspR</i> ::Tc	<i>tspR</i> and <i>rsmZ</i> double mutant of PAO1; Gm <sup>r</sup> , Tc <sup>r</sup>	This study
<i>tspR</i> ::Gm/CTX- <i>tspR</i>	<i>tspR</i> ::Gm complemented strain, derived from <i>tspR</i> ::Gm and Mini-CTX- <i>tspR</i> ; Gm <sup>r</sup> , Tc <sup>r</sup>	This study
<i>tspR</i> ::Gm /p- <i>tspR</i>	<i>tspR</i> ::Gm complemented strain, derived from <i>tspR</i> ::Gm and pAK1900- <i>tspR</i> ; Gm <sup>r</sup> , Cb <sup>r</sup>	This study
<i>retS</i> ::Gm/p- <i>retS</i>	<i>retS</i> ::Gm complemented strain, derived from <i>retS</i> ::Gm and pAK1900- <i>retS</i> ; Gm <sup>r</sup> , Cb <sup>r</sup>	This study
<i>retS</i> ::Gm/p- <i>tspR</i>	<i>retS</i> ::Gm strain carrying plasmid pAK1900- <i>tspR</i> ; Gm <sup>r</sup> , Cb <sup>r</sup>	This study
<b>Plasmids</b>		
pBT20	Mini-TnM delivery vector, Ap <sup>r</sup> , Gm <sup>r</sup>	(Kulasekara, <i>et al.</i> , 2005)
pEX18Ap	<i>oriT</i> <sup>+</sup> <i>sacB</i> <sup>+</sup> gene replacement vector with multiple-cloning site from pUC18; Ap <sup>r</sup>	(Hoang, <i>et al.</i> , 1998)
pEX18Tc	<i>oriT</i> <sup>+</sup> <i>sacB</i> <sup>+</sup> gene replacement vector with multiple-cloning site from pUC18; Tc <sup>r</sup>	(Hoang, <i>et al.</i> , 1998)
pPS858	Source plasmid of Gm <sup>r</sup> cassette; Gm <sup>r</sup> , Ap <sup>r</sup>	(Hoang, <i>et al.</i> , 1998)
pRK2013	Broad-host-range helper vector; Tra <sup>+</sup> , Kn <sup>r</sup>	(Ditta, <i>et al.</i> , 1980)
pMS402	Expression reporter plasmid carrying the promoterless <i>luxCDABE</i> gene; Kn <sup>r</sup> , Tmp <sup>r</sup>	(Duan, <i>et al.</i> , 2003)
pAK1900	<i>E. coli</i> - <i>P. aeruginosa</i> shuttle cloning vector carrying <i>plac</i> upstream of MCS; Ap <sup>r</sup> , Cb <sup>r</sup>	(Poole, <i>et al.</i> , 1993)
<i>exsA</i> -FLAG-A	<i>exsA</i> genes with Ptac and FLAG	(Li, <i>et al.</i> , 2013)
<i>exsA</i> -FLAG-S	<i>exsA</i> genes with Ptac, promoter region and FLAG	(Li, <i>et al.</i> , 2013)

mini-CTX- <i>lacZ</i>	Integration plasmid; Tc <sup>r</sup>	(Becher & Schweizer, 2000)
mini-CTX- <i>lux</i>	Integration plasmid; Tc <sup>r</sup>	(Becher & Schweizer, 2000)
pEX- <i>tspR</i> <sub>Gm</sub>	<i>tspR</i> deletion plasmid, pEX18Tc with 1959 bp upstream region, Gm <sup>r</sup> cassette from pPS858 and 1463 bp downstream region of <i>tspR</i> ; Tc <sup>r</sup> , Gm <sup>r</sup>	This study
pEX- <i>tspR</i> <sub>Tc</sub>	<i>tspR</i> deletion plasmid, pEX18Ap with 1959 bp upstream region, Tc <sup>r</sup> cassette from mini-CTX-lacZ and 1463 bp downstream region of <i>tspR</i> ; Tc <sup>r</sup>	This study
pEX- <i>retS</i>	<i>retS</i> deletion plasmid, pEX18Ap with 1958 bp upstream region, Gm <sup>r</sup> cassette from pPS858 and 1981 bp downstream of <i>retS</i> ; Ap <sup>r</sup> , Gm <sup>r</sup>	This study
pEX- <i>exsA</i>	<i>exsA</i> deletion plasmid, pEX18Ap with 1963 bp upstream region, Gm <sup>r</sup> cassette from pPS858 and 1972 bp downstream region of <i>exsA</i> ; Ap <sup>r</sup> , Gm <sup>r</sup>	This study
pEX- <i>rsmY</i>	<i>rsmY</i> deletion plasmid, pEX18Ap with 1963 bp upstream region, Tc <sup>r</sup> cassette from mini-CTX-lacZ and 1979 bp downstream of <i>rsmY</i> ; Ap <sup>r</sup> , Tc <sup>r</sup>	This study
pEX- <i>rsmZ</i>	<i>rsmZ</i> deletion plasmid, pEX18Ap with 1974 bp upstream region, Tc <sup>r</sup> cassette from mini-CTX-lacZ and 1972 bp downstream of <i>rsmZ</i> ; Ap <sup>r</sup> , Gm <sup>r</sup>	This study
p- <i>tspR</i>	pAK1900 with the entire <i>tspR</i> gene; Ap <sup>r</sup>	This study
p- <i>retS</i>	pAK1900 with the entire <i>retS</i> gene; Ap <sup>r</sup>	This study
<i>pexoS-lux</i>	pMS402 containing <i>exoS</i> promoter region; Kn <sup>r</sup> , Tmp <sup>r</sup>	(Duan, et al., 2003)
<i>PexoY-lux</i>	pMS402 containing <i>exoY</i> promoter region; Kn <sup>r</sup> , Tmp <sup>r</sup>	(Duan, et al., 2003)
<i>PexoT-lux</i>	pMS402 containing <i>exoT</i> promoter region; Kn <sup>r</sup> , Tmp <sup>r</sup>	(Duan, et al., 2003)
<i>PtspR-lux</i>	pMS402 containing <i>tspR</i> promoter region; Kn <sup>r</sup> , Tmp <sup>r</sup> . The promoter region of <i>tspR</i> is -522 to +77 from <i>tspR</i> translational starting site.	This study
<i>PretS-lux</i>	pMS402 containing <i>retS</i> promoter region; Kn <sup>r</sup> , Tmp <sup>r</sup> . The promoter region of <i>retS</i> is -378 to +139 from <i>retS</i> translational starting site.	This study
<i>PexsC-lux</i>	pMS402 containing <i>exsC</i> promoter region; Kn <sup>r</sup> , Tmp <sup>r</sup>	(Kong, et al., 2013)
CTX- <i>tspR-lux</i>	Integration plasmid, CTX6.1 with a fragment of pKD- <i>tspR</i> ; Kn <sup>r</sup> , Tc <sup>r</sup>	This study
CTX- <i>exoS-lux</i>	Integration plasmid, CTX6.1 with a fragment of pKD- <i>exoS</i> ; Kn <sup>r</sup> , Tc <sup>r</sup>	(Liang, et al., 2011)

CTX-exoT-lux	Integration plasmid, CTX6.1 with a fragment of pKD-exoT; Kn <sup>r</sup> , Tc <sup>r</sup>	(Liang, et al., 2011)
CTX-exoY-lux	Integration plasmid, CTX6.1 with a fragment of pKD-exoY; Kn <sup>r</sup> , Tc <sup>r</sup>	(Liang, et al., 2011)

## Reference

- Becher A & Schweizer HP (2000) Integration-proficient *Pseudomonas aeruginosa* vectors for isolation of single-copy chromosomal *lacZ* and *lux* gene fusions. *BioTechniques* **29**: 948-953.
- Ditta G, Stanfield S, Corbin D & Helinski DR (1980) Broad host range DNA cloning system for gram-negative bacteria: construction of a gene bank of *Rhizobium meliloti*. *Proc Natl Acad Sci U S A* **77**: 7347-7351.
- Duan K, Dammel C, Stein J, Rabin H & Surette MG (2003) Modulation of *Pseudomonas aeruginosa* gene expression by host microflora through interspecies communication. *Mol Microbiol* **50**: 1477-1491.
- Hoang TT, Karkhoff-Schweizer RR, Kutchma AJ & Schweizer HP (1998) A broad-host-range Flp-FRT recombination system for site-specific excision of chromosomally-located DNA sequences: application for isolation of unmarked *Pseudomonas aeruginosa* mutants. *Gene* **212**: 77-86.
- Kong W, Chen L, Zhao J, Shen T, Surette MG, Shen L & Duan K (2013) Hybrid sensor kinase PA1611 in *Pseudomonas aeruginosa* regulates transitions between acute and chronic infection through direct interaction with RetS. *Mol Microbiol* **88**: 784-797.
- Kulasekara HD, Ventre I, Kulasekara BR, Lazdunski A, Filloux A & Lory S (2005) A novel two-component system controls the expression of *Pseudomonas aeruginosa* fimbrial cup genes. *Mol Microbiol* **55**: 368-380.
- Li K, Xu C, Jin Y, et al. (2013) SuhB is a regulator of multiple virulence genes and essential for pathogenesis of *Pseudomonas aeruginosa*. *MBio* **4**: e00419-00413.
- Liang H, Duan J, Sibley CD, Surette MG & Duan K (2011) Identification of mutants with altered phenazine production in *Pseudomonas aeruginosa*. *J Med Microbiol* **60**: 22-34.
- Poole K, Neshat S, Krebes K & Heinrichs DE (1993) Cloning and nucleotide sequence analysis of the ferripyoverdine receptor gene *fpvA* of *Pseudomonas aeruginosa*. *J Bacteriol* **175**: 4597-4604.