

TABLE S1 Strains and plasmids used in this study.

Strains/plasmids	Number	Used in figures	Characteristics	Source or reference
<i>Pseudomonas aeruginosa</i>				
PA14		3, 4, 5, 6, S3, S4, S5	Clinical isolate UCBPP-PA14	(1)
PA14 Δ <i>ldhA</i>	LD2729	3, 5, S3	PA14 with a deletion <i>ldhA</i>	This study
PA14 Δ <i>lldDE</i>	LD2735	3, 4, 5, S3, S4, S5	PA14 with a deletion <i>lldD-E</i>	This study
PA14 Δ <i>lldA</i>	LD2844	3, 4, 5, S3, S5	PA14 with a deletion <i>lldA</i>	This study
PA14 Δ <i>lldDE</i> Δ <i>lldA</i>	LD2759	3, 4, S5	PA14 with deletions of <i>lldD-E</i> and <i>lldA</i>	This study
PA14 Δ <i>lldDE</i> Δ <i>lldA</i> :: <i>lldDE</i>	LD2904	3	PA14 Δ <i>lldDE</i> Δ <i>lldA</i> with complementation of <i>lldD-E</i>	This study
PA14 <i>attB</i> ::MCS- <i>gfp</i>	LD2820	2, 7, S2, S4	PA14 with MCS- <i>gfp</i> inserted at the <i>attB</i> site using pLD2722 (pSEK103)	This study
PA14 <i>attB</i> :: <i>ldhAp-gfp</i>	LD3095	S2	PA14 with <i>ldhAp-gfp</i> inserted at the <i>attB</i> site using pLD2814 (pSEK103- <i>gacSp</i>)	This study
PA14 <i>attB</i> :: <i>gacSp-gfp</i>	LD2815	5, 7, S2	PA14 with <i>gacSp-gfp</i> inserted at the <i>attB</i> site using pLD2814 (pSEK103- <i>gacSp</i>)	This study
PA14 <i>attB</i> :: <i>lldPp-gfp</i>	LD2798	2, 4, 5, 6, 7, S4	PA14 with <i>lldPp-gfp</i> inserted at the <i>attB</i> site using pLD2797 (pSEK103- <i>lldPp</i>)	This study
PA14 <i>attB</i> :: <i>lldAp-gfp</i>	LD2868	2, 4, 5, 6	PA14 with <i>lldAp-gfp</i> inserted at the <i>attB</i> site using pLD2867 (pSEK103- <i>lldAp</i>)	This study
<i>Escherichia coli</i>				
UQ950	LD44		<i>E. coli</i> DH5 α λ (<i>pir</i>) strain for cloning; F Δ (<i>argF-lac</i>) 169 ϕ 80d <i>lacZ</i> 58(Δ M15) <i>glnV44</i> (AS) <i>rjbD1 gyrA96</i> (Nal ^R) <i>recA1 endA1 spoT thi-1 hsdR17 deoR</i> λ <i>pir</i> ⁺	D. Lies
BW29427	LD661		Donor strain for biparental conjugation; <i>thrB1004 pro thi rpsL hsdS lacZ</i> Δ M15RP4-1360 Δ (<i>araBAD</i>)567 Δ <i>dapA1341</i> ::[<i>erm pir</i> (wt)]	B. Wanner
S17-1	LD2901		Str ^R , Tp ^R , F ⁻ RP4-2-Tc::Mu <i>aphA</i> ::Tn7 <i>recA</i> λ <i>pir</i> lysogen	(2)
<i>Saccharomyces cerevisiae</i>				
InvSc1	LD622		<i>MATa/MATα leu2/leu2 trp1-289/trp1-289 ura3-52/ura3-52 his3-Δ1/his3-Δ1</i>	Invitrogen
Plasmids				
pMQ30	LD621		Yeast-based allelic-exchange vector; <i>sacB</i> ⁺ , CEN/ARSH, URA3 ⁺ , Gm ^R	(3)
pLD2728 (pMQ30- Δ <i>ldhA</i>)	LD2728		Δ <i>ldhA</i> flanking fragments introduced into pMQ30 by gap repair cloning in yeast strain InvSc1	This study
pLD2734 (pMQ30- Δ <i>lldDE</i>)	LD2734		Δ <i>lldDE</i> flanking fragments introduced into pMQ30 by gap repair cloning in yeast strain InvSc1	This study
pLD2758 (pMQ30- Δ <i>lldA</i>)	LD2758		Δ <i>lldA</i> flanking fragments introduced into pMQ30 by gap repair cloning in yeast strain InvSc1	This study
pLD2903 (pMQ30- <i>lldDE</i> -comp)	LD2903		<i>lldDE</i> complementation flanking fragments introduced into pMQ30 by gap repair cloning in yeast strain InvSc1	This study

pYL122	LD742		Amp ^R <i>rhlA-gfp</i> transcription fusion in mini-CTX- <i>lacZ</i>	(4)
pUC18-mini-Tn7			Amp ^R ColE1 replicon mini-Tn7 base vector	(5)
pLD844 (pSEK101)	LD844		<i>rhlA</i> promoter of pYL122 was removed (XhoI and EcoRI) and replaced with a multiple cloning site (MCS) from pUC18-mini-Tn7	(6)
pLD2722 (pSEK103)	LD2722		<i>aacC1</i> (gentamicin resistance cassette) from pMQ30 was inserted at the BspDI site of pLD844	(6)
pLD3094 (pSEK103- <i>ldhA</i> p)	LD3094		494 bp of <i>ldhA</i> promoter sequence inserted at the MCS (SpeI and EcoRI) of pLD2722	This study
pLD2814 (pSEK103- <i>gacS</i> p)	LD2814		297 bp of <i>gacS</i> promoter sequence inserted at the MCS (SpeI and XhoI) of pLD2722	This study
pLD2797 (pSEK103- <i>lldP</i> p)	LD2797		324 bp of <i>lldP</i> promoter sequence inserted at the MCS (SpeI and XhoI) of pLD2722	This study
pLD2867 (pSEK103- <i>lldA</i> p)	LD2867		356 bp of <i>lldA</i> promoter sequence inserted at the MCS (SpeI and XhoI) of pLD2722	This study
pFLP2	LD743		Site-specific excision vector with <i>cl857</i> -controlled FLP recombinase; <i>sacB</i> ⁺ , Amp ^R	(7)

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