

Supplementary material

Table S1. Strains, plasmids and primers used in this study.

Strain/Plasmid/Primer	Relevant characteristics ^a	Source
<i>Escherichia coli</i>		
TOP10	F ⁻ <i>mcrA</i> Δ(<i>mrr-hsdRMS-mcrBC</i>) φ80lacZΔM15 Δ <i>lacX74 recA1 araD139</i> Δ(<i>ara-leu</i>) 7697 <i>galU galK rpsL</i> (Str ^R) <i>endA1 nupG λ-</i>	Invitrogen
OmniMAX	F' { <i>proAB+ lacI^q lacZΔM15 Tn10(Tet^R) Δ(ccdAB)</i> } <i>mcrA</i> Δ(<i>mrr-hsdRMS-mcrBC</i>) Φ80lacZΔM15 Δ(<i>lacZYA-argF</i>) U169 <i>endA1 recA1 supE44 thi-1 gyrA96 relA1 tonA panD phoA20 thi-1 rpsE rpoB argE</i> (Am) <i>recA1 Rf' (lpir)</i>	Invitrogen
CC118λpir	Host strain for pKNG101; D (<i>ara-leu</i>) <i>araD DlacX74 galE galK-phoA20 thi-1 rpsE rpoB argE</i> (Am) <i>recA1 Rf' (lpir)</i>	(1)
1047	Helper strain for conjugation of plasmids harbouring pRK2013, Km ^R	Lab collection
S17-1	F ⁻ , lambda (-), <i>endA</i> , <i>thi</i> , <i>pro</i> , <i>recA</i> , restriction (-) modification (+), RP4 derivative integrated into the chromosome with <i>Tet::Mu</i> , <i>Km::Tn7</i>	Lab collection
<i>Pseudomonas aeruginosa</i>		
PAK	Wild-type	(2)
PAKΔ <i>retS</i>	PAK with a <i>retS</i> deletion	(2)
PAKΔ <i>pelG</i>	PAK with a <i>pelG</i> deletion	(3)
PAKΔ <i>retSΔpelF</i>	PAK with deletions of <i>retS</i> and <i>pelF</i>	(4)
PAKΔ <i>retSΔPA0338</i>	PAK with deletions of <i>retS</i> and <i>PA0338</i>	This study
PAKΔ <i>retSΔsadC</i>	PAK with deletions of <i>retS</i> and <i>sadC</i>	This study
PAKΔ <i>sadC</i>	PAK with a <i>sadC</i> deletion	This study
PAKΔ <i>retSΔsadB</i>	PAK with deletions of <i>retS</i> and <i>sadB</i>	This study
PAKΔ <i>retSΔsadCΔbifA</i>	PAK with deletions of <i>retS</i> , <i>sadC</i> and <i>bifA</i>	This study
PAKΔ <i>hptB</i>	PAK with a <i>hptB</i> deletion	(5)
PAKΔ <i>hptBΔsadC</i>	PAK with deletions of <i>hptB</i> and <i>sadC</i>	This study
PAKΔ <i>hptBΔsadB</i>	PAK with deletions of <i>hptB</i> and <i>sadB</i>	This study
PAKΔ <i>rsmA</i>	PAK with a <i>rsmA</i> deletion	(6)
PAKΔ <i>rsmAΔsadC</i>	PAK with deletions of <i>rsmA</i> and <i>sadC</i>	This study
PAKΔ <i>rsmYZ</i>	PAK with deletions of <i>rsmY</i> and <i>rsmZ</i>	(5)
PA01	Wild-type	Lab collection
PA01 PA0338-Flag	PA01 containing PA0338 with a chromosomal C-terminal Flag-tag	This study
PA01 PA0847-Flag	PA01 containing PA0847 with a chromosomal C-terminal Flag-tag	This study
PA01 PA4332-Flag	PA01 containing PA4332 with a chromosomal C-terminal Flag-tag	This study
PA01 PA5487-Flag	PA01 containing PA5487 with a chromosomal C-terminal Flag-tag	This study
<i>Plasmids</i>		
pCR2.1-TA	Cloning plasmid, Ap ^R , Km ^R	Invitrogen
pBBR1MCS-4	Broad host range vector, Ap ^R	(7)
pBBR1MCS-4- <i>sadC</i>	<i>sadC</i> cloned into pBBR1MCS-4 (Apal/BamHI)	This study
pBBR1MCS-4- <i>ladS</i>	<i>ladS</i> cloned into pBBR1MCS-4	(6)
pBBR1MCS-4- <i>hsbR</i>	<i>hsbR</i> cloned into pBBR1MCS-4	(5)

Strain/Plasmid/Primer	Relevant characteristics^a	Source
pKNG101	Suicide vector carrying <i>sacB</i> , <i>SmR</i>	(8)
pKNG101- Δ PA0338	Mutator fragment for deletion of PA0338 (from amino acid 2 to 367), <i>Sm^R</i>	This study
pKNG101- Δ sadC	Mutator fragment for deletion of <i>sadC</i> (from amino acid 87 to 414), <i>Sm^R</i>	This study
pKNG101- Δ sadB	Mutator fragment for deletion of <i>sadB</i> (from amino acid 4 to 467), <i>Sm^R</i>	This study
pKNG101- Δ bifA	Mutator fragment for deletion of <i>bifA</i> (from amino acid 4 to 657), <i>Sm^R</i>	This study
P _{cdrA-gfp}	Plasmid expressing <i>gfp</i> from the promoter of <i>cdrA</i> , <i>Ap^R</i> , <i>Gm^R</i>	(9)
pME3087	Tet ^R , suicide vector for allelic replacement; <i>ColeI</i> -replicon, <i>IncP-1</i> , <i>Mob</i>	(10)
pME3087-0338-Flag	pME3087 with the cassette for introducing a c-terminal Flag-tag on PA0338 (BamHI/EcoRI)	This study
pME3087-0847-Flag	pME3087 with the cassette for introducing a c-terminal Flag-tag on PA0847 (BamHI/HinDIII)	This study
pME3087-4332-Flag	pME3087 with the cassette for introducing a c-terminal Flag-tag on PA4332 (BamHI/HinDIII)	This study
pME3087-5487-Flag	pME3087 with the cassette for introducing a c-terminal Flag-tag on PA5487 (BamHI/EcoRI)	This study
pME6918	Tet ^R , <i>rsmY</i> under Ptac promoter at position +1 in pME6032	(11)
pME3849	Gm ^R , <i>rsmA</i> under Plac promoter in pME6001 (BamHI/SmaI)	(12)
<i>Primers for qRT-PCR</i>		
qGyrAF	AACGACTGGAACAAGCCCTACA	This study
qGyrAR	GCGCACGATGGTGTGTA	This study
qPcrVF	AGTGGGATCTGCGCGAGTT	This study
qPcrVR	TGGGTCTGCAGGACATCCTT	This study
qHsiA1F	ATCGCCTGCTCGAGTATTACG	This study
qHsiA1R	GGGATGAGATTCTCACGATT	This study
qPA0338F	GAGCGTGGAGACCTGGAAAGA	This study
qPA0338R	GCAGTATTCCGGGCTTCG	This study
<i>Primers for deletion mutants</i>		
1PA0338F	GAECTCCTTCGGTCGGTTC	This study
2PA0338R	ATCGCGACGCACGACGAAGCGTCC	This study
3PA0338F	TTCGTCGTGCGTCGCGATTGCGTG	This study
4PA0338R	CCATCAGCTCTGGCACTACA	This study
5PA0338F	ATCCTGCTGCCGATGATCT	This study
6PA0338R	ACTATTCCGGCTTCAACCTG	This study
1SadCF	TATCGTCACGCTCATGGTA	This study
2SadCR	TTCCAGCTGAGCGGACCGCGGATTATC	This study
3SadCF	CGGTCCGCTCAGCTGGAAAGCTGTTGCGA	This study
4SadCR	TCAGGATCGAAAGGCTGCAA	This study
5SadCF	ATGGTCGGTACGGTTTCAG	This study
6SadCR	CTCCACAGCAGCAGGTGCT	This study
1SadBF	ACTGCGTGCCTACACCAC	This study
2SadBR	TCACCCGGTTCTGTCATGACGAGACCATG	This study

Strain/Plasmid/Primer	Relevant characteristics^a	Source
3SadBF	ATGACAGAACCGGGGTGACCGGGTAG	This study
4SadBR	GGCAAGGACCTGGTGTCCAGTT	This study
5SadBF	CAACGCCAGCCTGATGAT	This study
6SadBR	CGTCTCTCCTGTGGATGCT	This study
1BifAF	GCTGGCTGGTGAAGAAGG	This study
2BifAR	GCTGTACAGCAGTTCAAGGGGCCTCCT	This study
3BifAF	TTGAAACTGCTGTACAGCAAGCCGCTGC	This study
4BifAR	GTCGCCAGGGACTCGATG	This study
5BifAF	GGCTCCTTCGACAAGTTCAA	This study
6BifAR	CTCTACCGCTACACCGAGGA	This study
<i>Primers for cloning of sadC</i>		
sadCF	TTCACACAGGAAAAGGAGATGAACTGCAGGGC	This study
sadCR	GATCGAAAGGCTGCAACAC	This study
<i>Primers introduction of Flag-tag</i>		
PA0338Flag up F BamH1	GACCTGGAAGAGGATCCTCCATCC CTATTATCGTCGTATCTTGATGATCATGATCTTT	This study
PA0338Flag up R	ATAATCACCGTCATGGTCTTGATGTCGAAGCAACAGG CCACGCAATCGC GAATACAAAGACCATGACGGTGATTATAAGATCATGATA	This study
PA0338Flag down F	TCGACTACAAAGATGACGACGATAATAGTAAAGGTC AGCCTGGCGGCAGCACCAAG	This study
PA0338Flag down R EcoRI	CCAACCGTGACGAATTCTATGCC	This study
PA0847Flag up F BamHI	CCATCCCTACTAGGATCCGCTC	This study
PA0847Flag up R	CTATTATCGTCGTATCTTGATGATCATGATCTTT ATAATCACCGTCATGGTCTTGATGTCGAAGGCTGGCG CCTGGTACAGGCAATAG GAATACAAAGACCATGACGGTGATTATAAGATCATGATA	This study
PA0847Flag down F	TCGACTACAAAGATGACGACGATAATAGTAAAGCAGCG GACAAACAGG	This study
PA0847Flag down R HinDIII	CATCCAGATGCCGAAAAGCTTCCACAG	This study
PA4332Flag up F BamHI	CGTTCATCGCTTTCCGGATCCTACCTG	This study
PA4332Flag up R	CTATTATCGTCGTATCTTGATGATCATGATCTTT ATAATCACCGTCATGGTCTTGATGTCGAAGGCACTGG TGACCTCCCAGG GAATACAAAGACCATGACGGTGATTATAAGATCATGATA	This study
PA4332 Flag down F	TCGACTACAAAGATGACGACGATAATAGTAAAGCAGCG CATACGGGTCGG	This study
PA4332Flag down R HinDIII	CAGGTGAAGCTTCTGGATCG	This study
PA5487Flag up F BamHI	GTCAGTGGCCTGGATCCCAGCGTGC	This study
PA5487Flag up R	CTATTATCGTCGTATCTTGATGATCATGATCTTT ATAATCACCGTCATGGTCTTGATGTCGAAGGCCACTT CCAGGCGGTCCC	This study
PA5487Flag down F	GAATACAAAGACCATGACGGTGATTATAAGATCATGATA	This study

Strain/Plasmid/Primer	Relevant characteristics ^a	Source
PA5487Flag down R EcoRI	TCGACTACAAAGATGACGACGATAAATAGTAAAGGGCGAC CAGCGCACATTC CGAGGGAATTCTGGAGGAAC	This study

^aAp^R, Ampicillin resistance; Str^R, Streptomycin resistance; Km^R, Kanamycin resistance; Gm^R, Gentamicin resistance; Tet^R, Tetracyclin resistance.

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