

Table S3. Strains, plasmids and oligos used in this study.

| Strain/Plasmid | Description | Source |
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| <i>E. coli</i> strains | | |
| DH5 α | Cloning and plasmid maintenance strain | Lab stock |
| XL1-Blue | Cloning and plasmid maintenance strain | Lab stock |
| CC118 λ pir | Expresses the λ pir protein, allowing replication of suicide plasmids with the R6K origin of replication. | A. Filloux lab stock |
| SM10 λ pir | <i>P. aeruginosa</i> conjugation donor strain Expresses the λ pir protein, allowing replication of suicide plasmids with the R6K origin of replication. Km ^R | A. Filloux lab stock (Simon <i>et al.</i> , 1983) |
| T7 Express <i>lysY^{iq}</i> | Protein expression strain. | New England Biolabs, Ipswich, MA |
| T7 Express <i>lysY^{iq}</i> PA0709 | Protein expression strain harbouring pET15DG1 PA0709 | This study |
| T7 Express <i>lysY^{iq}</i> PA3390 | Protein expression strain harbouring pET15DG1 PA3390 | This study |
| <i>P. aeruginosa</i> strains | | |
| WT MPAO1 | Wild-type <i>P. aeruginosa</i> PAO1 strain (clinical, non-respiratory isolate) | Manoil Lab. (Holloway, 1955; Jacobs <i>et al.</i> , 2003; Stover <i>et al.</i> , 2000) |
| Δ phuR | Unmarked deletion of <i>phuR</i> in MPAO1 background. Mutant retains the first two and last two codons of the <i>phuR</i> gene | This study |
| Δ phuS | Unmarked deletion of <i>phuS</i> in MPAO1 background. Mutant retains the first two and last two codons of the <i>phuS</i> gene | This study |
| Δ phuUV | Unmarked deletion of <i>phuUV</i> in MPAO1 background. Mutant retains the first two codons of the <i>phuU</i> gene and the last two codons of the <i>phuV</i> gene | This study |
| Δ hasR | Unmarked deletion of <i>hasR</i> in MPAO1 background. Mutant retains the first two and last two codons of the <i>hasR</i> gene | This study |
| Δ hxC | Unmarked deletion of <i>hxC</i> in MPAO1 background. Mutant retains the first two and last two codons of the <i>hxC</i> gene | This study |
| Δ OMR | Unmarked deletions of <i>phuR</i> , <i>hasR</i> and <i>hxC</i> in MPAO1 background. Mutant retains the first two and last two codons of each of the deleted genes | This study |
| Δ cheY | Unmarked deletion of <i>cheY</i> in MPAO1 background. Mutant retains the first two and last two codons of the <i>cheY</i> gene | This study |
| Δ prfF1,2 | Unmarked deletion of the PrrF1/2 sRNAs in PAO1 background | A.Oglesby-Sherrouse (Wilderman <i>et al.</i> , 2004) |
| Δ PA0028 | Unmarked deletion of PA0028 in MPAO1 background. Mutant retains the first three and last three codons of the PA0149 gene | This study |
| Δ PA0149 | Unmarked deletion of PA0149 in MPAO1 background. Mutant retains the first three and last three codons of the PA0149 gene | This study |

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| ΔPA0527 | Unmarked deletion of PA0149 in MPAO1 background. Mutant retains the first three and last three codons of the PA0527 gene | This study |
| ΔPA0709 | Unmarked deletion of PA0709 in MPAO1 background. Mutant retains the first three and last three codons of the PA0709 gene | This study |
| ΔPA3390 | Unmarked deletion of PA3390 in MPAO1 background. Mutant retains the first three and last three codons of the PA3390 gene. | This study |

| Plasmids | Description | Source |
|--------------------|---|---|
| pSB109 | Medium copy number expression vector for expression of genes under control of the pBAD promoter. Gm ^R . | Bardy Lab Ketelboeter <i>et al.</i> 2017 |
| pIFP | pSB109 expressing IFP1.4 under the control of the pBAD promoter. Gm ^R . | This study |
| pIFPHO | pSB109 expressing the IFP+HO synthetic operon construct under the control of the pBAD promoter. Gm ^R . | This study |
| pIFPHO_AR | pIFPHO modified to express the <i>bla</i> gene from pUCP19. Gm ^R and Amp ^R . | This study |
| pBT20 | Mariner transposon mutagenesis vector. Gm ^R and Amp ^R . | Kulasekara <i>et al.</i> 2005 |
| pBT20_AS | pBT20 modified to lack the <i>bla</i> gene. Gm ^R and Amp ^S . | This study |
| pET-15b | Vector for expression of His ₆ -tagged proteins. Amp ^R . | Novagen - EMD Millipore |
| pET15DG1 | pET-15b modified to contain a TEV protease cleavage site in place of the thrombin cleavage site. Amp ^R . | This study |
| pET15DG1 PA0709 | pET15DG1 containing PA0709. Amp ^R . | This study |
| pET15DG1 PA3390 | pET15DG1 containing PA3390. Amp ^R . | This study |

| Oligo description | Oligo name | Nucleotide sequence (5'-3') |
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| Primers to poison pSB109 <i>NcoI</i> site | <i>NcoI</i> poison-F | GTTTCCACGGTGTGCGTCGATGGGCAAATATTAT ACGCAAG |
| | <i>NcoI</i> poison-R | CTTGCGTATAATATTTGCCATCGACGCACACCGT GGAAAC |
| Primers for construction of pIFP and pIFPHO | IFP F1 | CATGTCACGTGACCCACTTCCATTCTTC |
| | IFP F2 | TCACGTGACCCACTTCCATTCTTC |
| | IFP R1 | TATGTCATTTGTCGTCGTCGTCTTTGTAGTCGAAT TCAGCTTCTTTACGTTGAACTTGAC |
| | IFP R2 | TGTCATTTGTCGTCGTCGTCTTTGTAGTCGAATTC AGCTTCTTTACGTTGAACTTGAC |
| | IFPHO R1 | TATGTCATTTGTCGTCGTCGTCTTTGTAGTCACCT TCTGAAGTAGCAAGACCAACTTC |
| | IFPHO R2 | TGTCATTTGTCGTCGTCGTCTTTGTAGTCACCTTC TGAAGTAGCAAGACCAACTTC |

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| | pSB109 seq F | GAAAAGTCCACATTGATTATTTGCACG |
| | pSB109 seq R | CGGGCCTCTTCGCTATTACG |
| Primers for construction of MPAO1 Δ <i>phuR</i> mutant | phuR_primer 1 | CGATATCCGGCGAGACCACC |
| | phuR_primer 2 | TCAGATGTTCGAGCGGCATGTGGGACTC |
| | phuR_primer 3 | ATGCCGCTCGACATCTGACCCGTTACC |
| | phuR_primer 4 | CCAGCTCCACCCAGAGGC |
| | phuR_primer 5 | CCTTCAACGCGGCCAGC |
| | phuR_primer 6 | GCCGTATCGAAACGCTCC |
| Primers for construction of MPAO1 Δ <i>phuS</i> mutant | phuS_primer 1 | CGCTGAAGAAGTCGTTGG |
| | phuS_primer 2 | TCAGAGCGCGCTGCTCATCGACGGTTC |
| | phuS_primer 3 | ATGAGCAGCGCGCTCTGAGGAGAGATC |
| | phuS_primer 4 | CGCGATCCAGTCCGCCTGG |
| | phuS_primer 5 | CGAGGTCAACCAGGAGTGG |
| | phuS_primer 6 | CGGCTTCCGTACTIONCAGCG |
| Primers for construction of MPAO1 Δ <i>phuUV</i> mutant | phuUV_primer 1 | CGTCCTGCTGGTGATCGG |
| | phuUV_primer 2 | TCAACGGGCGACCAACACGAGCGAAAT |
| | phuUV_primer 3 | GTGTTGGTCCCGTTGAAGGAGTCGC |
| | phuUV_primer 4 | CGCATGATCTCGCTCCGCT |
| | phuUV_primer 5 | CTGCCGAGGGCGTGCTGG |
| | phuUV_primer 6 | GCATGGCGACTCCTTCAACG |
| Primers for construction of MPAO1 Δ <i>hxC</i> mutant | hxC_primer 1 | CCGTGGCGGTGAAACAGGG |
| | hxC_primer 2 | TCAGAAGAACAAGGTCATGCTGGAAC |
| | hxC_primer 3 | ATGACCTTGTTCTTCTGAGGGGCGTCA |
| | hxC_primer 4 | GCACCGCCAGTCTCGACC |
| | hxC_primer 5 | CCTCGCCTACGGCACGGC |
| | hxC_primer 6 | CCTGCTGCTGCTCAAGAC |
| Primers for construction of MPAO1 Δ <i>cheY</i> mutant | cheY_primer 1 | CGACGCGATCCGCGCGATC |
| | cheY_primer 2 | TCAGCCGTTAATTTTCATGTTCTTGTC |
| | cheY_primer 3 | ATGAAAATTAACGGCTGAAGCCGGCCA |
| | cheY_primer 4 | GAGGAACTGTTTCGATGCGC |
| | cheY_primer 5 | GGAGCAACTGATCCAGCG |

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| | cheY_primer 6 | GCTCCAGCGCGGCCTGGTG |
| Primers to check PA0028 mutant | PA0028 check F | GACTACCTCTGGTACCGCCT |
| | PA0028 check R | CGTTTCGATGCCAACCAGGAG |
| Primers to check PA0149 mutant | PA0149 check F | GACTACCTCTGGTACCGCCT |
| | PA0149 check R | CGTTTCGATGCCAACCAGGAG |
| Primers to check PA0527 mutant | PA0527 check F | GAAGTGAACCACATGAGCGGG |
| | PA0527 check R | CTTCGCGCTGTACGGGTTTTTC |
| Primers to check PA0709 mutant | PA0709 check F | GTGTGGCCGAACACATAGAG |
| | PA0709 check R | GCGAACCGACCTACTACTCG |
| Primers to check PA3390 mutant | PA3390 check F | GTTGCCGGATGCTAGGAGG |
| | PA3390 check R | CATCCGCTCCTGCTTCATGG |
| Primers to construct PA3390 mutant | PA3390 BamHI F | GGATCCCTTCAGGGAGAGGGGAGAG |
| | PA3390 Up R | GATTGAGCCTTCTCAGCGCAGGCAGTACACGGAA ACTTCCTTTTG |
| | PA3390 Down F | CAAAAGGAAGTTTCCGTGTA CTGCCTGCGCTGAGAAGGCTCAATC |
| | PA3390 SpeI R | ACTAGTGACAAGCCGCGACTGAAGG |
| Primers for construction of pUCP19-phuUV complement plasmid | phuUV_Forward | GATACAAGCTTCGAAACCGCTGAGTACG |
| | phuUV_Reverse | GCTAGGGATCCTCAACGGGCGACGATCAG |
| Primers for amplification of pUCP19 bla | pUCP19_bla_F | CTAGACTGCAGCGTCAGGTGGCACTTTTCG |
| | pUCP19_bla_R | CGATGCTGCAGGGATCTCAAGAAGATCCTTTGAT C |
| Primers used in <i>Tn</i> -seq analysis | pBT20-92_1 | GAGCATCGTTTGTTCGCCCGGCTTCTGTATGG |
| | olj376 | GTGACTGGAGTTCAGACGTGTGCTCTTCCGATCT GGGGGGGGGGGGGGGGGG |

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| | pBT20_PAIR_Am pF_2 | AATGATACGGCGACCACCGAGATCTACACTCTTT CGTCTAGAAGCCTGCTTTCTAGAGACCGGGGACT TATCAGC |
| | TdT_Index_1_AT CACG | CAAGCAGAAGACGGCATAACGAGATCGTGATGTGA CTGGAGTTCAGACGTGTGCTCTTCCGATCT |
| | TdT_Index_3_TT AGGC | CAAGCAGAAGACGGCATAACGAGATGCCTAAGTGA CTGGAGTTCAGACGTGTGCTCTTCCGATCT |
| | TdT_Index_4_TG ACCA | CAAGCAGAAGACGGCATAACGAGATTGGTCAGTGA CTGGAGTTCAGACGTGTGCTCTTCCGATCT |
| | pBT20_SEQ-6 | CCTGCTTTCTAGAGACCGGGGACTTATCAGCCAA C |
| Primers for construction of pET15DG1 | pET15b TEV F | CATGGGCAGCAGCCATCATCATCATCACAGC AGCGGCGAGAATCTTTATTTTCAGGGACA |
| | pET15b TEV R | TATGTCCCTGAAAATAAAGATTCTCGCCGCTGCTG TGATGATGATGATGATGGCTGCTGCC |
| Primers for construction of PA0709 and PA3390 expression plasmids | PA0709-F1 | TATGACCTACCACGTA CTGGTTC |
| | PA0709-F2 | TGACCTACCACGTA CTGGTTC |
| | PA0709-R1 | GATCCTCAACCCTCGACGCGGTAG |
| | PA0709-R2 | CTCAACCCTCGACGCGGTAG |
| | PA3390-F1 | TATGTACTGCATCTTTATCAAGGCC |
| | PA3390-F2 | TGTACTGCATCTTTATCAAGGCC |
| | PA3390-R1 | GATCCTCAGCGCAGCGGATTGAGC |
| | PA3390-R2 | CTCAGCGCAGCGGATTGAGC |