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

Bacterial strain or plasmid	Description	Reference or source
<u>*P. aeruginosa*</u>		
PAO1	Wild type strain	(1)
PAO1 Δ*lasR*	PAO1 containing an unmarked, in-frame *lasR* deletion	(2)
PAO1 Δ*lasI*	PAO1 containing an unmarked, in-frame *lasI* deletion	(3)
PAO1 Δ*nuh*	PAO1 containing an unmarked, in-frame *nuh* deletion	This work
PAO1 ΔPA0148	PAO1 containing an unmarked, in-frame *nuh* deletion	This work
<u>PAO1 variants</u>		
A	Isolated after daily transfer in 0.75% adenosine / 0.25% casein for 25 days	(4)
B	Isolated after daily transfer in 0.75% adenosine / 0.25% casein for 25 days	(4)
C	Isolated after daily transfer in 0.9% adenosine / 0.1% casein for 14 days	This work
D	Isolated after daily transfer in 0.9% adenosine / 0.1% casein for 14 days	This work
E	Isolated after daily transfer in 0.9% adenosine / 0.1% casein for 14 days	This work
E. coli DH5α	F^- Φ80*lac*ZΔM15 Δ(*lac*ZYA-*arg*F) U169 *rec*A1 *end*A1 *hsd*R17(r_k^-, m_k^+) *pho*A*sup*E44 *thi*-1 *gyr*A96 *rel*A1 λ^-	Invitrogen
<u>Plasmids</u>		
pJN105	araC-P_{Bad} cassette cloned in pBBR1MCS-5, Gm^r	(5)
pJN105.nuh	Arabinose inducible *nuh* in pJN105, Gm^r	This work
pPROBE-GT	Broad-host-range pVS1/p15a GFP reporter, Gm^r	(6)
pBBR1MCS-5	Medium copy, BHR plasmid vector (pBBR), Gm^r	(7)
Pnuh-wt	pPROBE-GT with the -91 through 15 of the 5' region of *nuh* from PAO1 inserted with HindIII and BamHI, Gm^r	This work
Pnuh-var	pPROBE-GT with the -91 through 15 of the 5' region of *nuh* from variant A inserted with HindIII and BamHI, Gm^r	This work
pJT3	pBBR1MCS with PA0148 inserted with HindIII and SacI, Gm^r	This work
pJT4	pBBR1MCS with *nuh* inserted with HindIII and SacI, Gm^r	This work
pJT5	pBBR1MCS with PA0144 inserted with HindIII and SacI, Gm^r	This work
pJT7wt	pBBR1MCS with PA0142 through PA0148 assembled from PAO1, Gm^r	This work
pJT7var	pBBR1MCS with PA0142 through PA0148 assembled from Variant A, Gm^r	This work
pJT10	pBBR1MCS with PA0143 and PA0148 assembled from PAO1, Gm^r	This work

Primers	
Pnuh.BamHI R	5’ – AAAAAGGATCCCAAAAGCGATTGCATGGGCG – 3’
Pnuh.HindIII F	5’ – TTTTTAAGCTTGTTGATGCGGGTTCCGGTA G – 3’
nuhF300	5’ – CTCGGCGCTGCTGCTGTGCG – 3’
nuhR300	5’ – GGACAGCGGCACCGCCTTCG – 3’
pJT3 F.PA0148.HindIII	5’ – TTCAAGCTTCTACTCCATGCCGTTCTTCG – 3’
pJT3 R.PA0148.SacI	5’ – TTCGAGCTCCAGAAGCGCAGGAACAGTTC – 3’
pJT4 F.nuh.HindIII	5’ – TTCAAGCTTCTCGGCGCTGCTGCTGTGCG – 3’
pJT4 R.nuh.SacI	5’ – TTCGAGCTCGGACAGCGGCACCGCCTTCG – 3’
pJT5 F.PA0144.HindIII	5’ – TTCAAGCTTTATATGCCTACCTCAGCGATG – 3’
pJT5 F.PA0144.SacI	5’ – TTCGAGCTCCTACGTCCAGTACGACATCAA – 3’
pJT7.F	5’ – GGGATGTGCTGCAAGGCGATTAAGTTGGGTAACGCCAGGACGCCTCGAACATGATCCTC – 3’
pJT7.R	5’ – CACACAGGAAACAGCTATGACCATGATTACGCCAAGCGCGCATCAATACCGAGAAGCCGG – 3’
pJT7_V.F	5’ – GCAAGGCGTTTCGTCGTCGCCGGCTTCTCGGTATTGATGCGCGCTTGGCGTAATCATGG– 3’
pJT7_V.R	5’ – ACAGGGCCTGGCGCGCCTCGAGGATCATGTTCGAGGCGTCCTGGCGTTACCCAACTTAA – 3’
pJT10_PA0148 F	5’ – GCATAGCCGACCTCGAAGGCGGTGCCGCTGTCCCTACTCCATGCCGTTCTTCGC – 3’
pJT10_nuh R	5’ – GTCCATGTGCGGTTCGGCGAAGAACGGCATGGAGTAGGGACAGCGGCACCGCCTTCG – 3’
pJT11.D1	5’ – TGTAAAGCAAGCTTCTGCAGGTCGACTCTAGAGGATCGCCTGCTGGTGGAACAGATC – 3’
pJT11.D2	5’ – GCTTTTCAGCGCAGGCTCCAGAAACCCAGGCGATGCCGAACACGTCCGGCTC – 3’
pJT11.D3	5’ – GCCGCCTACGAGCCGGACGTGTTCGGCATCGCCTGGGTTTCTGGAGC – 3’
pJT11.D4	5’ – CGTGGAAATTAATTAAGGTACCGAATTCGAGCTCGAGGCCGCTGCTGGTCTTCATG – 3’
pJT11_V.F	5’ – GACCCGGCGGATTTTCCTGCTCAATCGCATCCACGGCTCGAGCTCGAATTCGGTACC – 3’
pJT11_V.R	5’ – GCTACGGGTGATATCGATCTTCAGCTTCTCTTCGACGGGGATCCTCTAGAGTCGAC – 3’

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