Table S4. Bacterial strains, plasmids, and primers used in this study.

Bacterial strain or plasmid	Description	Reference or source
P. aeruginosa strains		
PAO1	Wild type strain	(1)
PAO1 Δ*lasR*	PAO1 containing an unmarked, in-frame *lasR* deletion	(2)
PAO1 Δ*lasR* Δ*rhlR* Δ*qscR*	*lasR, rhlR, qscR triple* mutant in PAO1	(3)
Other strains		
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
Plasmids		
pSC11	Broad-host-range *lasI-lacZ* reporter, Ap^r	(4)
pJN105L	Arabinose inducible *lasR* in pJN105, Gm^r	(5)
pECP61.5	Contains an *rhlA-lacZ* translation fusion and IPTG-inducible *rhlR* , Ap^r	(6)
pPROBE-GT	Broad-host-range pVS1/p15a GFP reporter, Gm^r	(7)
pJF01	pPROBE-GT with -1 through-501 5' region of *rhlA* inserted with HindIII and BamHI, Gm^r	This work
pBS351	pPROBE-GT with -1 through-501 5' region of *lasI* inserted with HindIII and BamHI, Gm^r	This work

Primers	
lasR.cloning.EcoR1 F	5' – TTTTTGAATTCTAGCGCTATGGCCTTGGTT – 3'
lasR.cloning.Xba1 R	5' – AAAAATCTAGAGCAAGATCAGAGAGTAATAAGACCC – 3'
rhlA.GFPcloning.HindIII F	5' – TTTTTAAGCTTGCATGCGAGGCCTGCGAA – 3'
rhlA.GFPcloning.BamHI R	5' – AAAAAAGGATCCTCTAGAAACCGATACCAACAGACTTTCGC – 3'
GFP.QC R	5' – TTCTTTCCTGTACATAACCTTCGGGCA – 3'

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