

S2 Table. Bacterial Strains used in this study.

Species	Genotype	Source
<i>E. coli</i>	MG1655	ATCC 47076
	<i>fpr1</i> *	This study
	<i>fpr2</i> *	This study
	<i>mprA</i> *	This study
	<i>ompC</i> *	This study
	<i>ompC</i> * <i>fpr2</i> *	This study
	Δfpr	This study
	$\Delta mprA$	This study
	$\Delta ompC$	This study
<i>P. aeruginosa</i>	PA14	[1]
	$\Delta pvdJ$	This study
	$\Delta pchE$	This study
	$\Delta pqsA$	This study
	$\Delta pqsE$	This study
	$\Delta phzA1-G1 \Delta phzA2-G2$	This study
	$\Delta pvdJ \Delta pchE$	This study
	$\Delta pvdJ \Delta pchE \Delta pqsA$	This study
	$\Delta pvdJ \Delta pchE \Delta pqsE$	This study
	$\Delta pvdJ \Delta pchE \Delta phzA1-G1 \Delta phzA2-G2$	This study

1. Rahme LG, Stevens EJ, Wolfort SF, Shao J, Tompkins RG, Ausubel FM. Common virulence factors for bacterial pathogenicity in plants and animals. *Science*. 1995;268(5219):1899-902. doi: 10.1126/science.7604262.