

Supplemental Table 1. Strains used in this study

Species	Strain ID	MLST (*if high risk clone)	T3SS effector	Virulence in animal model of bloodstream infection (Allen <i>et al.</i> (2020))	Other relevant Characteristics	Reference
<i>P. aeruginosa</i>	PABL004	ST1734	ExoS	medium	Human bacteremia isolate, GenBank assembly: GCA_003411985.2	Scheetz <i>et al.</i> (2009)
<i>P. aeruginosa</i>	PABL006	ST235*	ExoU	high	Human bacteremia isolate, GenBank assembly: GCA_003412035.2	Scheetz <i>et al.</i> (2009)
<i>P. aeruginosa</i>	PABL012	ST708	ExoS	high	Human bacteremia isolate, GenBank assembly: GCA_003429185.1	Scheetz <i>et al.</i> (2009)
<i>P. aeruginosa</i>	PABL048	ST298*	ExoU	medium	Human bacteremia isolate, GenBank assembly: GCA_003411785.2	Scheetz <i>et al.</i> (2009)
<i>P. aeruginosa</i>	PABL049	ST244*	ExoS	high	Human bacteremia isolate, GenBank assembly: GCA_003411745.2	Scheetz <i>et al.</i> (2009)
<i>P. aeruginosa</i>	PABL054	ST111*	ExoS	medium	Human bacteremia isolate, GenBank assembly: GCA_003411595.2	Scheetz <i>et al.</i> (2009)