



Figure S1. goeBURST of the 3321 sequence types listed in the *Pseudomonas aeruginosa* PubMLST database (August 2019). Blue points represent sequence types (STs); lines connect single-locus variants (SLVs). Red boxes within the circles are the STs found in this study, light green boxes represent the clonal complex group founder, while dark green boxes the sub-group founder. ST1189 and ST3277 are singletons.