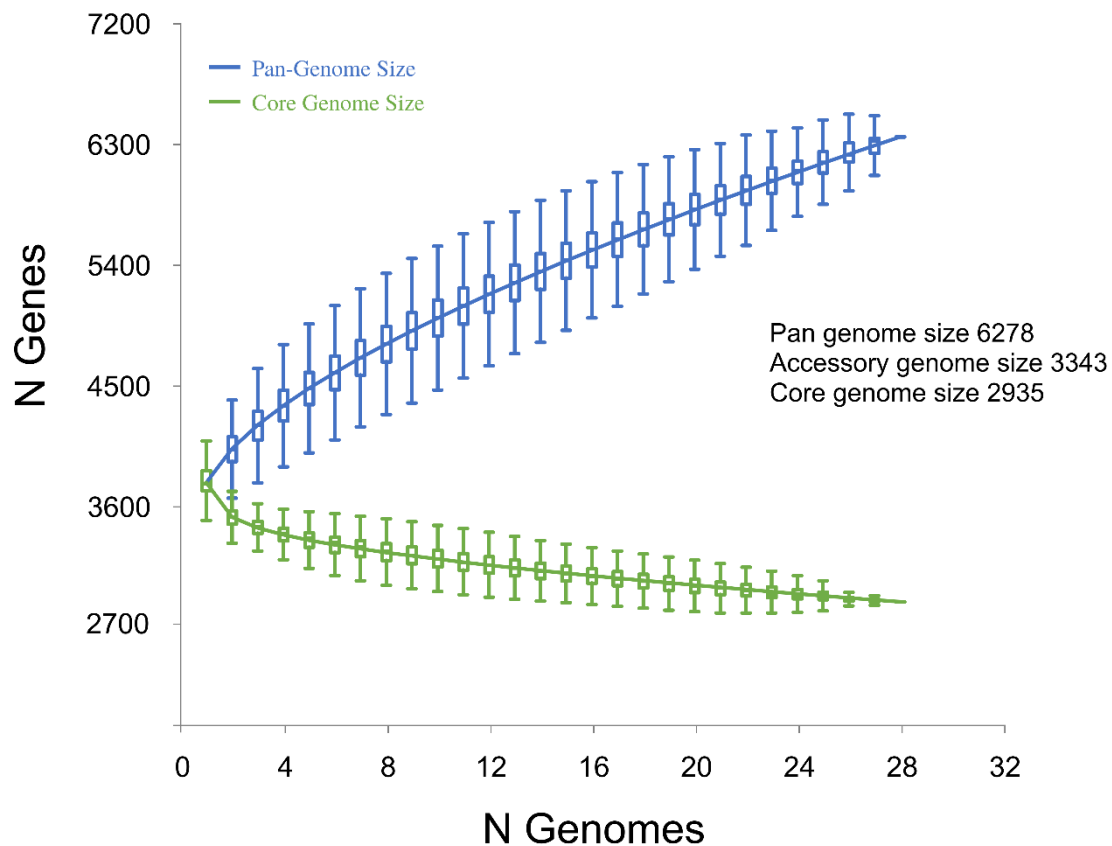


## Supplementary Material

**Knight DR\*, Squire MM, Collins DA and Riley TV (2016).** Genome analysis of *Clostridium difficile* PCR ribotype 014 lineage in Australian pigs and humans reveals a diverse genetic repertoire and signatures of long-range interspecies transmission. *Front. Microbiol.* **7**:2138. doi: 10.3389/fmicb.2016.02138

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### Supplementary Image 3 | Pan-genome analysis of *C. difficile* RT014 from humans

The total number of genes in the pan (blue) and core (green) genomes from humans ( $n=28$ ) are plotted as a function of the number of genomes sequentially added (see Methods and manuscript Figure 6 for details).