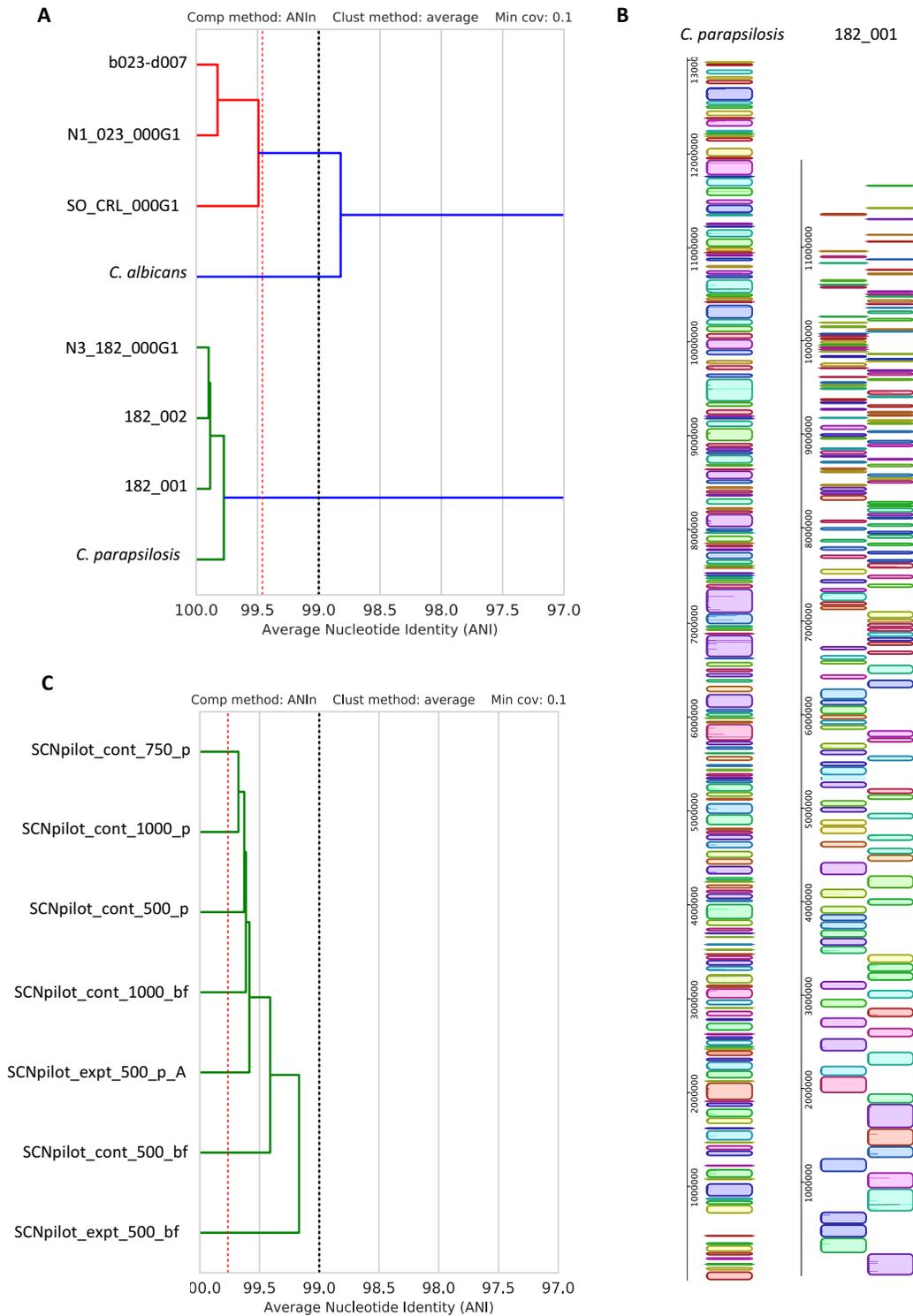


Supplemental Figure S5



**Supplemental Figure S5 Dendrograms of infant-derived fecal sample and thiocyanate reactor sample genomes based on Average Nucleotide Identity (ANI).** The compare\_wf workflow of dRep (Olm et al. 2017) with default parameters was used to cluster highly similar genomes and calculate ANI. *C. albicans* and *C. parapsilosis* genomes were obtained from the Candida Genome Database (Skrzypek et al.). (A) Fungal genomes binned from infant fecal-derived samples. (B) Rotifer genomes binned from thiocyanate reactor samples.