

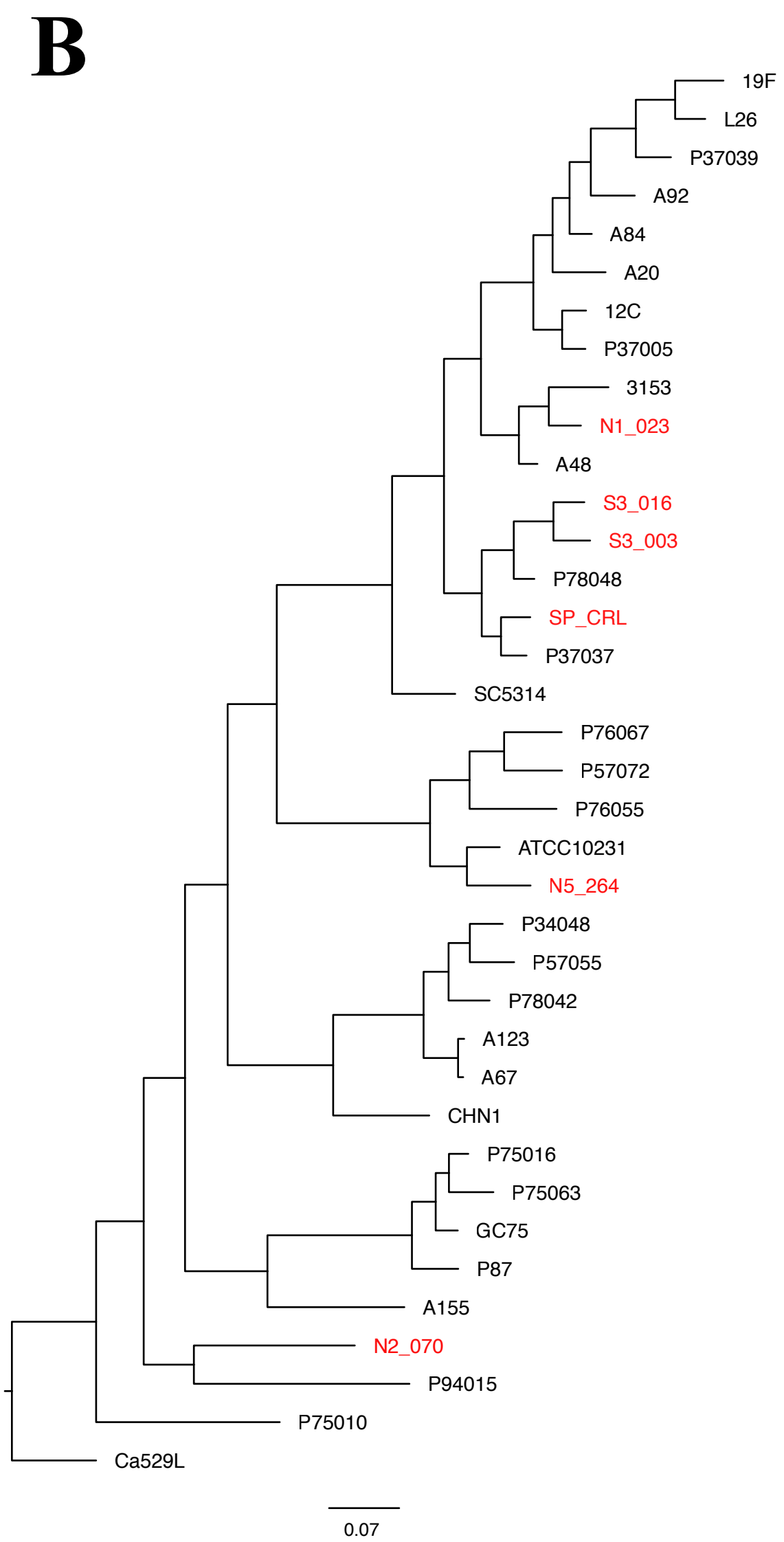
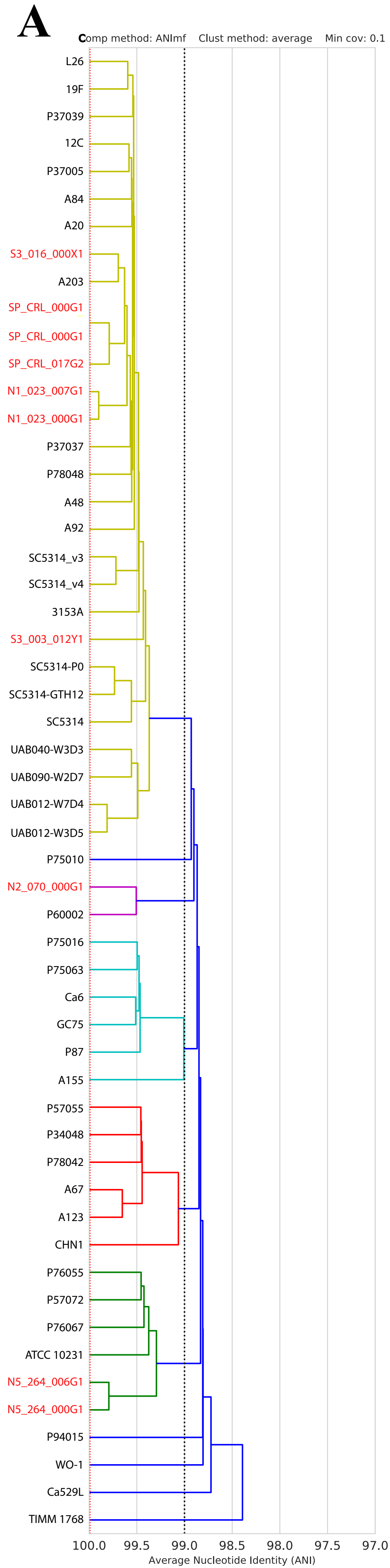


Figure S2: Phylogenetic and ANI comparisons of *C. albicans* genomes show no clear hospital associated population. (A) ANI comparisons for each *C. albicans* strain. All publically available *C. albicans* genomes from NCBI were included. Genomes assembled in this study are highlighted in red. If multiple genomes were assembled for the same strain from different time points, all assembled genomes were included in this analysis. **(B)** A phylogenetic tree of *C. albicans* strains constructed from concatenated SNVs. Strains with genomes assembled in this study are highlighted in red. All publically available *C. albicans* genome read sets on NCBI are included in this analysis.