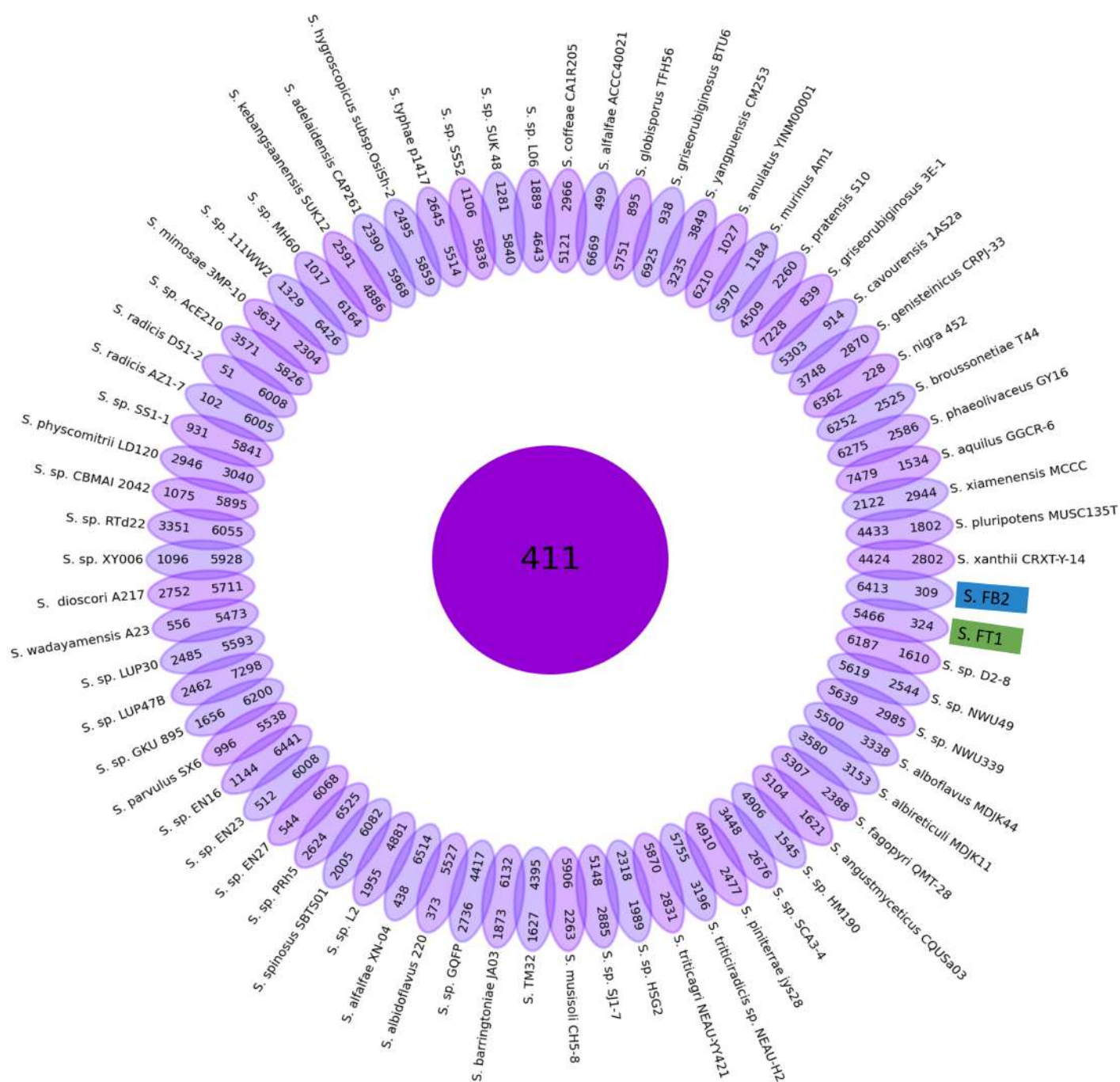




Supplementary Figure 1. Flower diagrams represent each genus's core, unique, and accessory genes. The numbers underneath the names represent the number of accessory genes. The pan-genome is open. C, D. Circular plots of FT1 and FB2 strains. The three innermost rings indicate the genome size, GC content, and GC skew of FT1 and FB2 strains, respectively. The outer rings illustrate the BLAST genome comparisons against the genome reference.