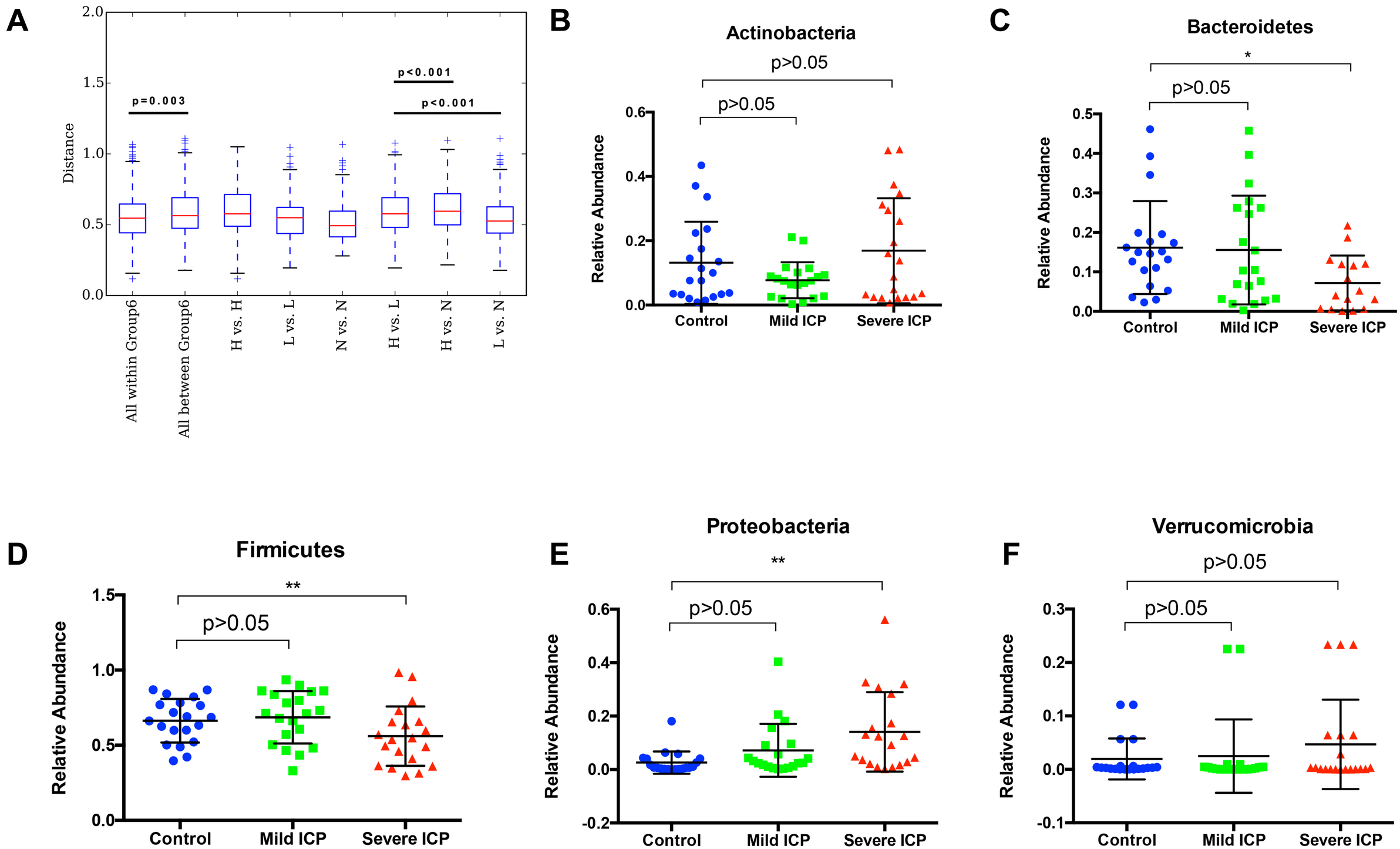

Figure S1. Diversity and microbial features at the phylum level of the gut microbiome of pregnant women with ICP

(A) Unweighted UniFrac-based distance among groups; N indicates the control group, L indicates the mild ICP group, and H indicates the severe ICP group. (B-F) Abundance of representative differentially abundant phyla in different groups: Actinobacteria, Bacteroidetes, Firmicutes, Proteobacteria and Verrucomicrobia.