

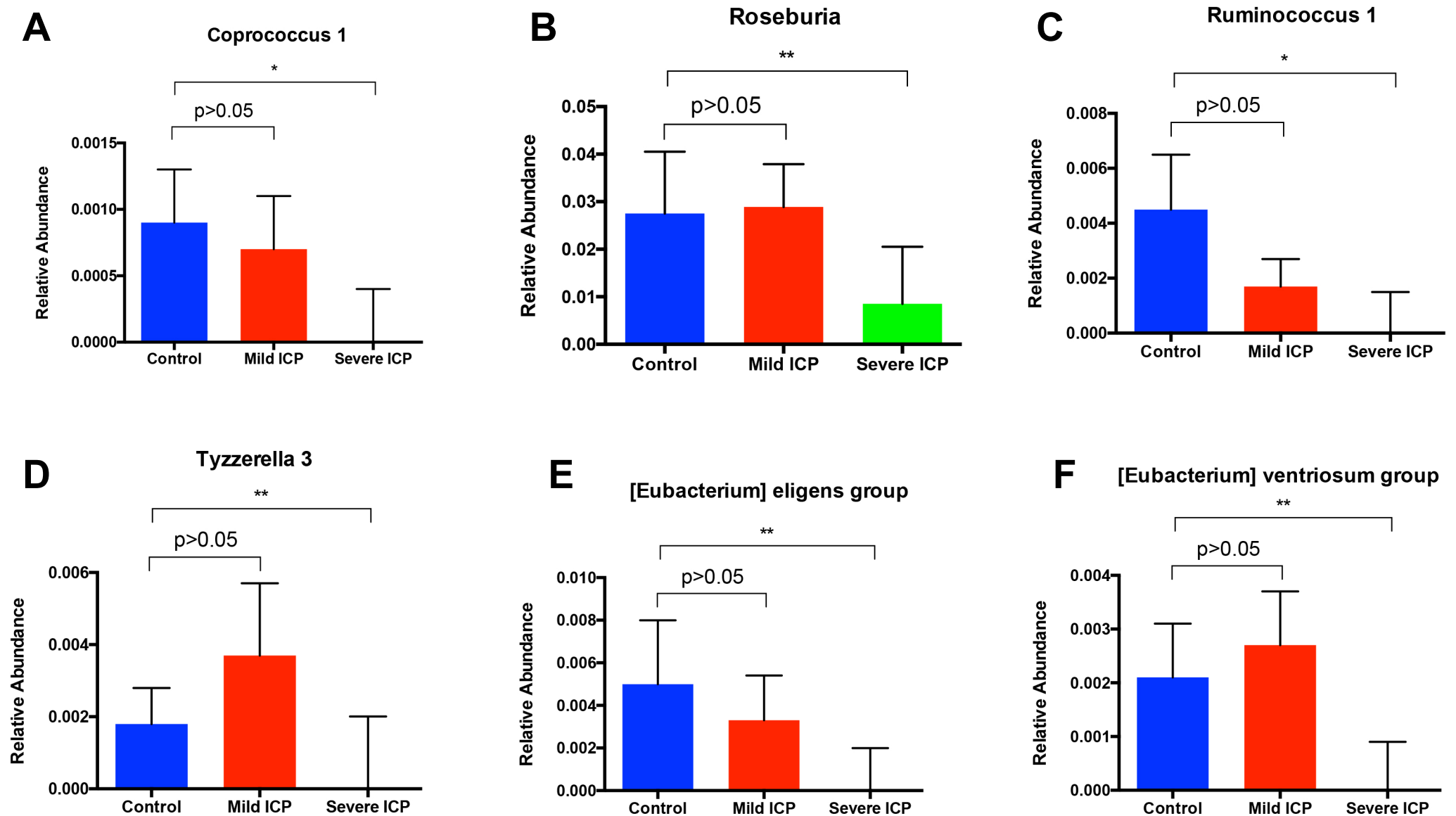


Figure S2. Microbiome composition in different groups at the genus level

(A-F) Abundance of representative differentially abundant genera in the different groups: Coprococcus 1, Roseburia, Ruminococcus 1, Tyzzerella 3, [Eubacterium] eligens group, and [Eubacterium] ventriosum group.