

SUPPLEMENTAL FIGURES

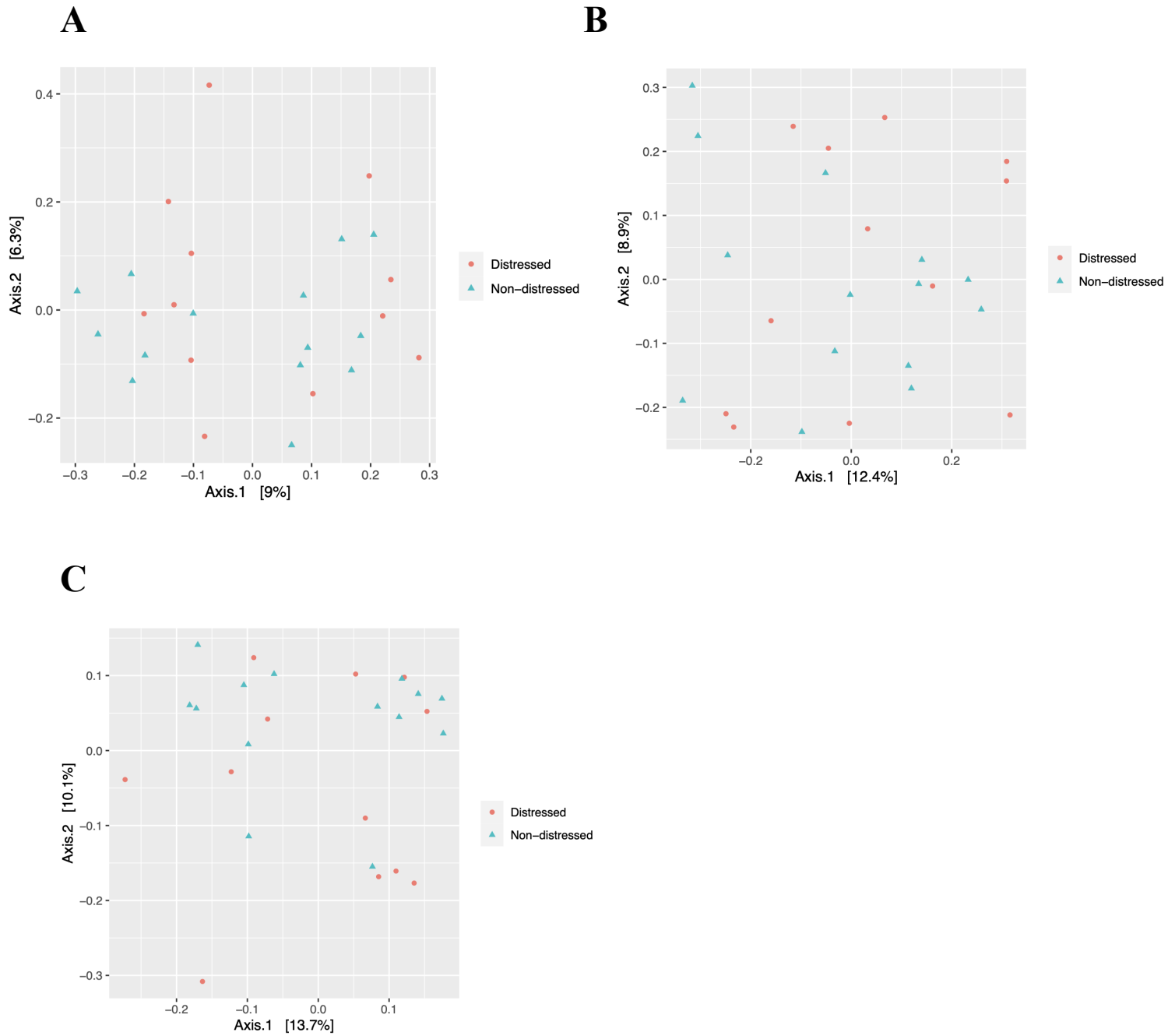


Figure S1. Patient status does not impact microbial community composition. The gut microbiome composition of distressed and non-distressed PD patients was analyzed with (A) Jaccard, (B) Bray-Curtis, and (C) Unweighted UniFrac beta diversity metrics.