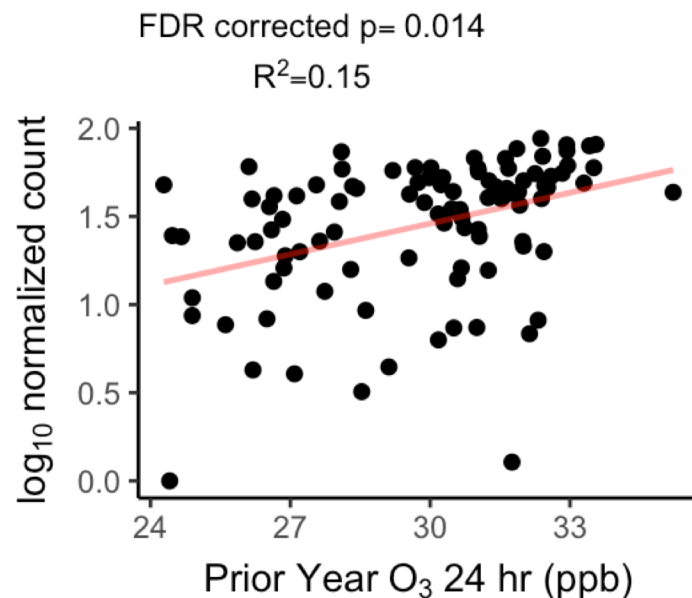


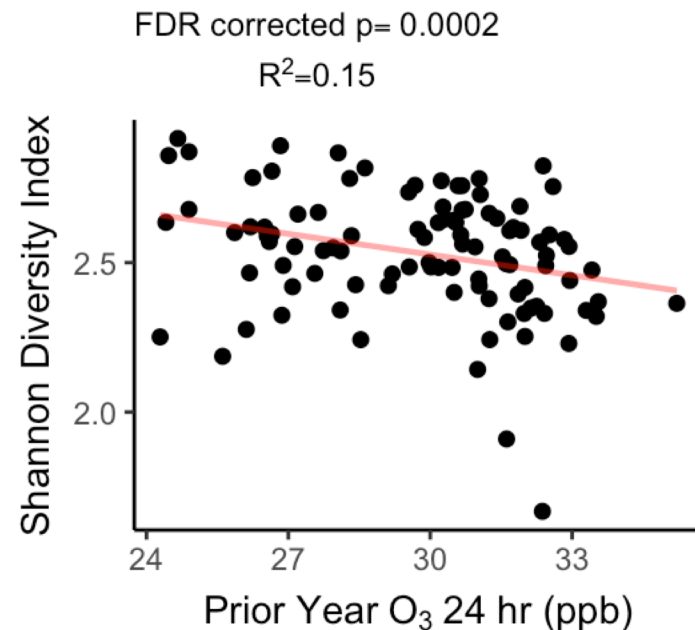
Supp Figure 6: Exposure to O₃ was Significantly Associated with a Higher Relative Abundance of *Bacteroides* and a Lower Diversity and Evenness Using Whole Genome Sequencing (WGS) Data that were Taxonomically Classified by MetaPhlan2

A

Bacteroides



B



C

