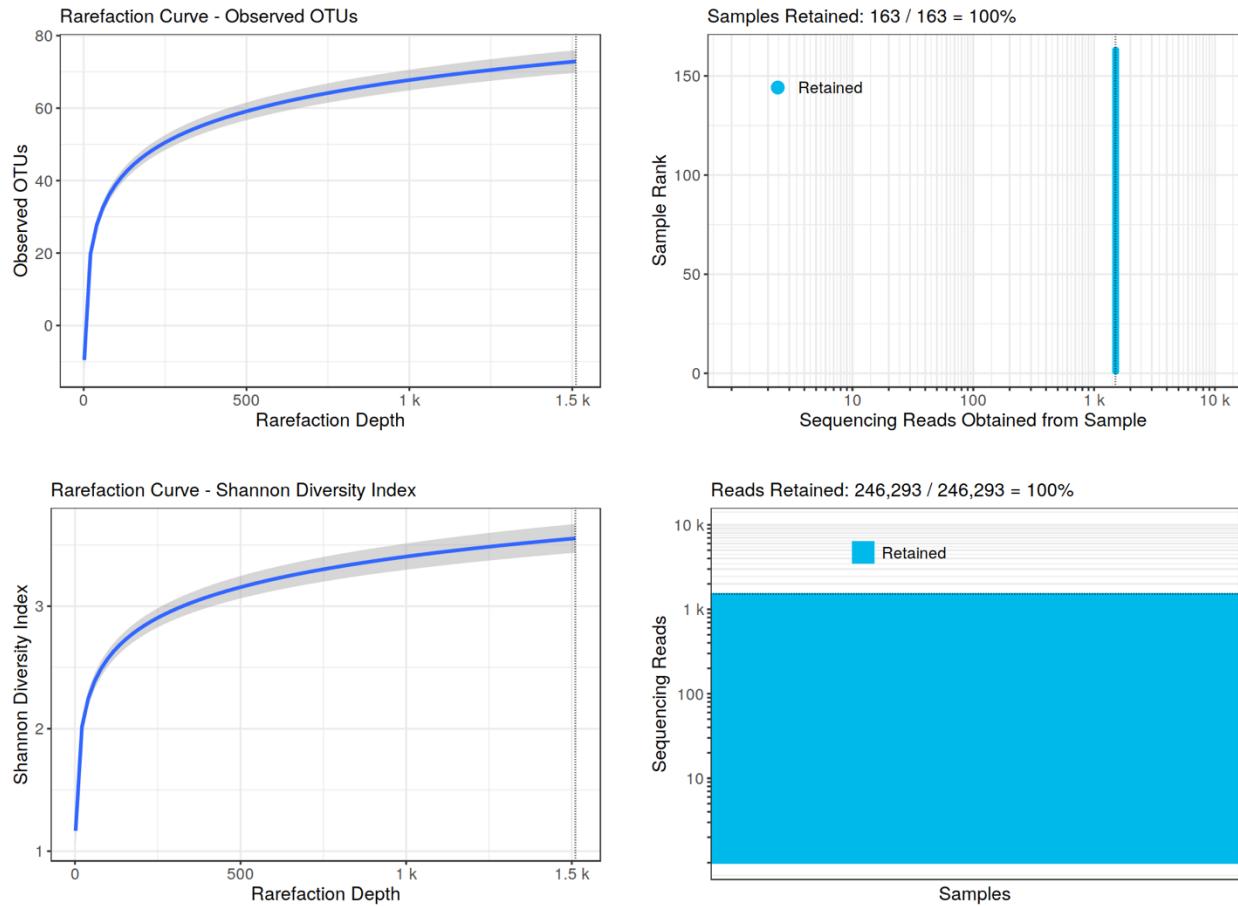
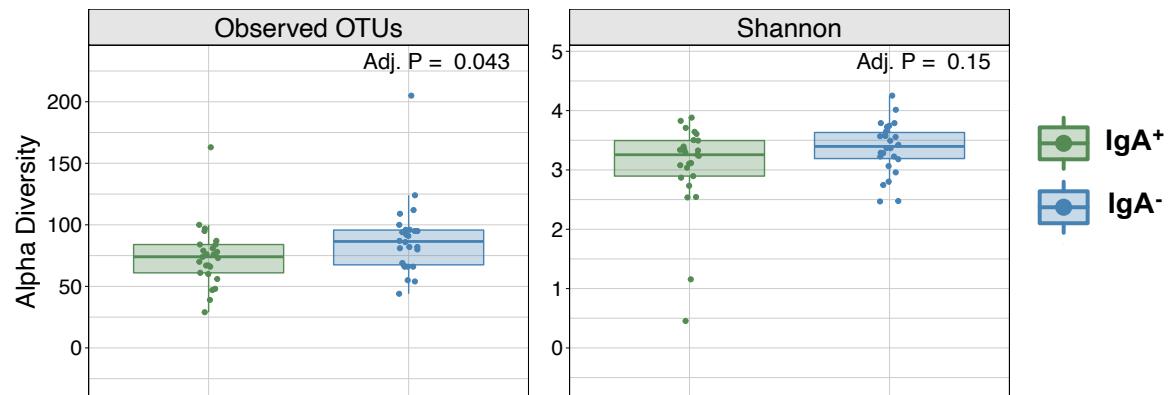


# Supplementary Material

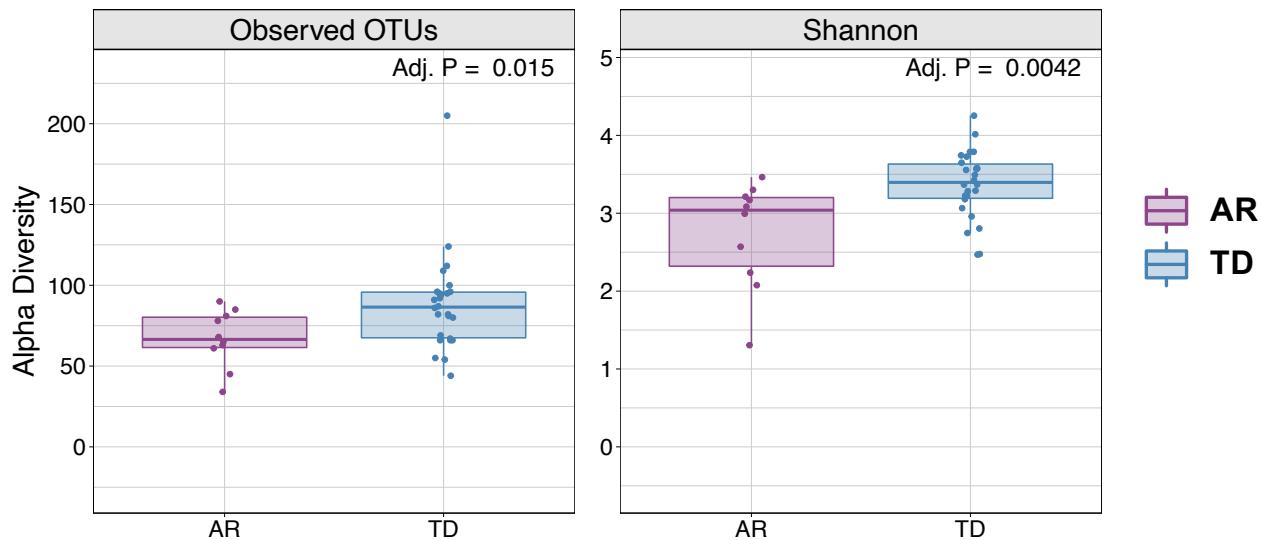
## IgA-Biome Profiles Correlate with Clinical Parkinson's Disease Subtypes



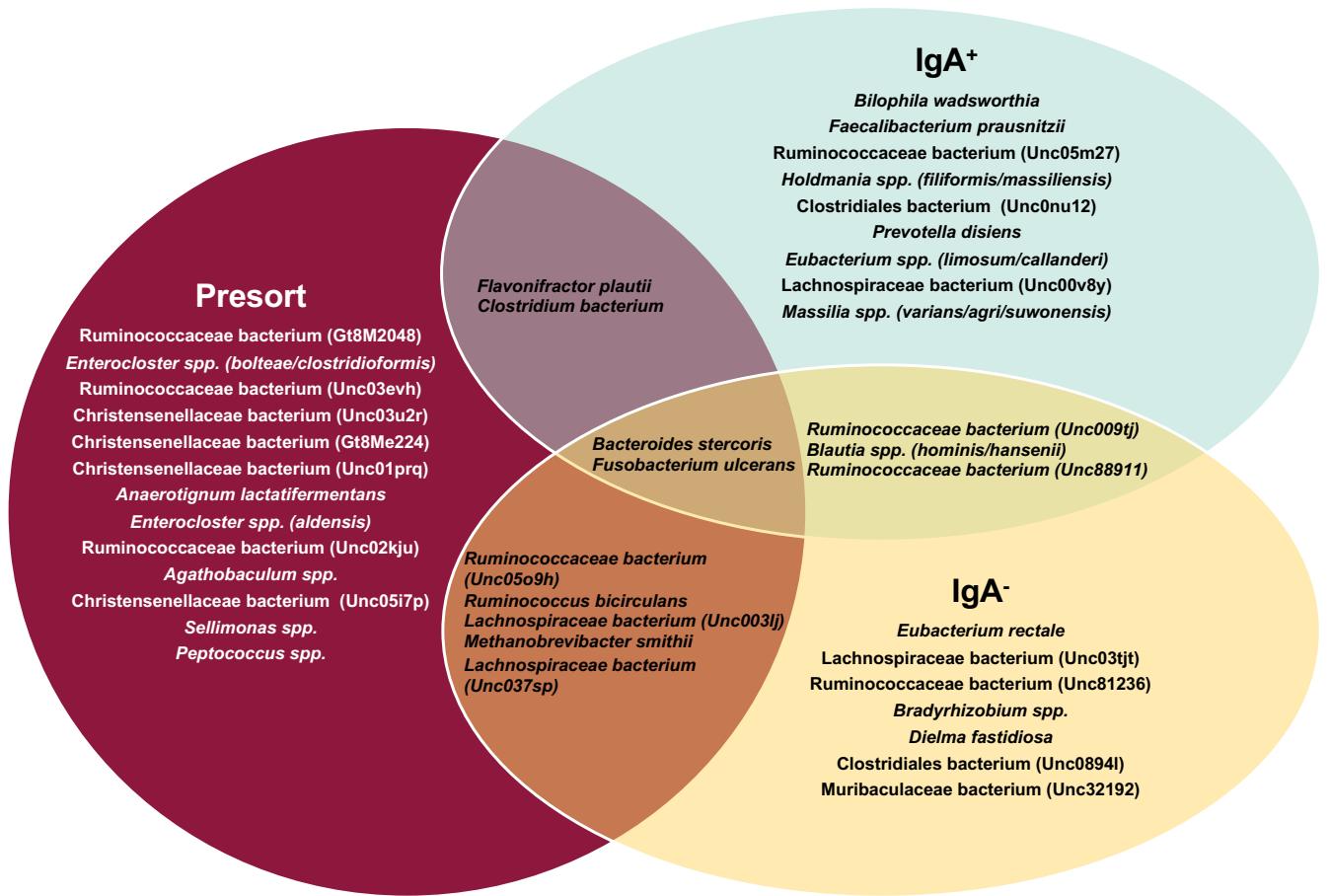
**Supplementary Figure 1.** Rarefaction curves of IgA-Biome samples.



**Supplementary Figure 2.** IgA-Biome alpha diversity of patients classified to the TD group. Box plots depict sample richness (Observed OTU) and evenness (Shannon Diversity Index) among sorted samples according to IgA-bound status. \*p=0.0427 FDR adjusted Mann-Whitley t test.



**Supplementary Figure 3.** IgA<sup>-</sup> compartment comparison between patients classified as either AR or TD. Box plots depict sample richness (Observed OTU) and evenness (Shannon Diversity Index). p=0.0148 and p=0.00424 FDR adjusted Mann-Whitney.



**Supplementary Figure 4.** Overlap in LEfSe-identified differential OTUs.