

Small Farm Holder Cropping Systems Influence Microbial Profiles in an Equatorial Rainforest Agroecosystem

Supplementary Material

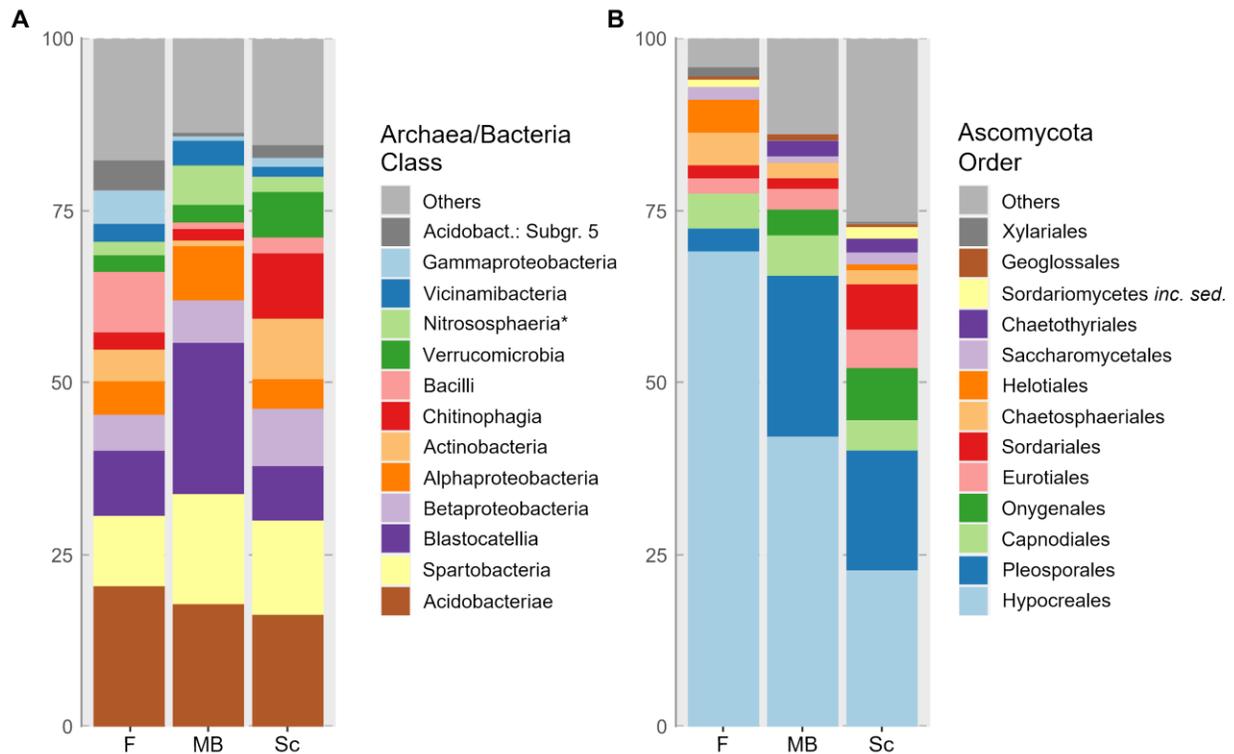


Figure. S1. Relative abundance of amplified sequence variants grouped by (A) prokaryote class (16S RNA) and (B) Ascomycota order (28S RNA) for each cropping system. Asterisk denotes Archaea. Cropping system include: F – forest, MB – maize-bean, Sc – sugarcane.

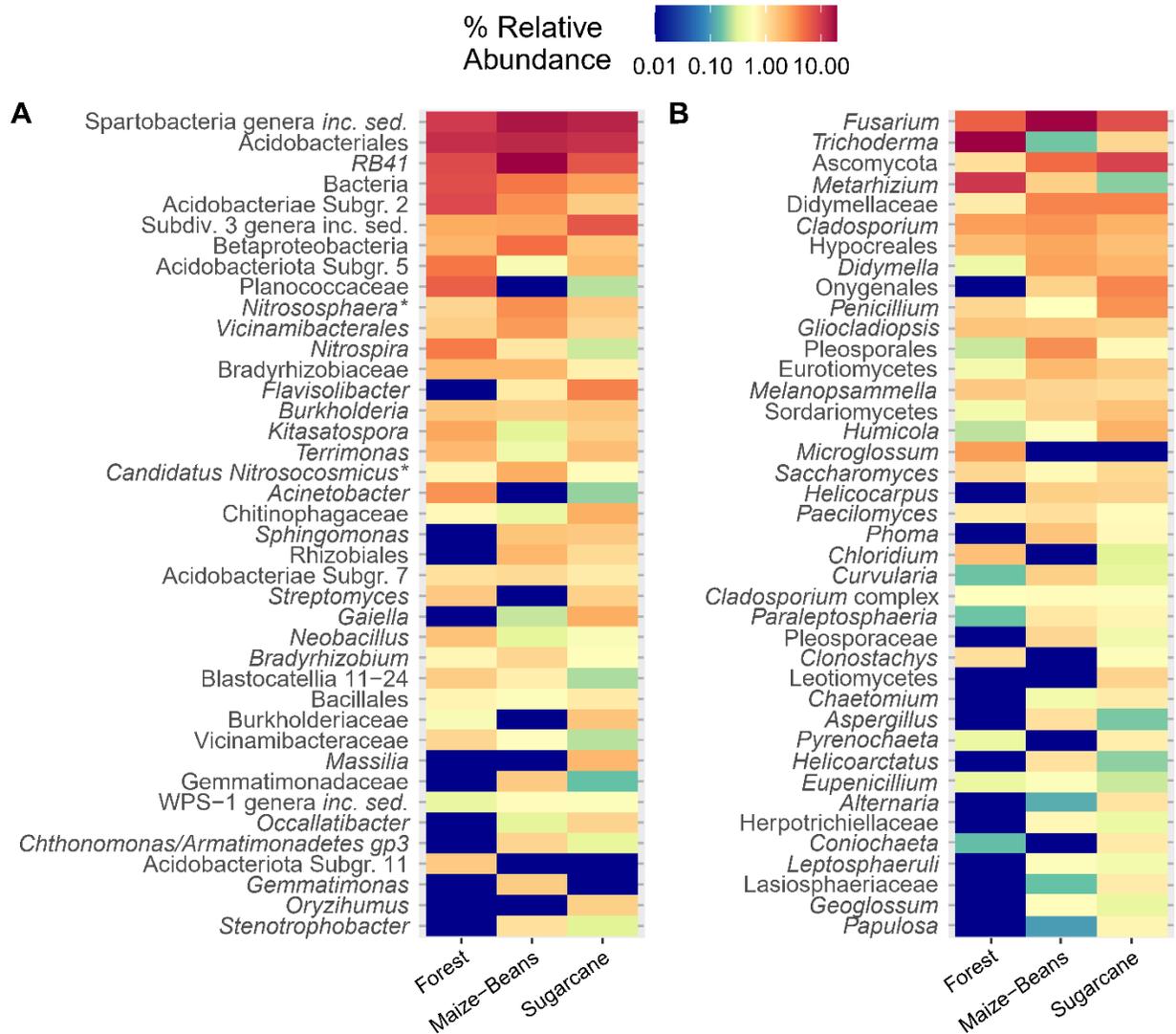


Figure. S2. Log relative abundance heatmap of the top 40 ASVs within (A) prokaryote and (B) Ascomycota samples. Asterisks identify groups of Archaea.