

Figure S1. Relative abundance of 12 ASVs present in all samples in different tissues of leafhopper *R. dorsalis* at different developmental stages. Circle area corresponds to relative abundance of bacteria in each sample. N1, N2, N3, N4, N5, A refer to 1st-instar nymphs, 2nd-instar nymphs, 3rd-instar nymphs, 4th-instar nymphs, 5th-instar nymphs and adults, respectively. Mg, midgut; Mt, Malpighian tubule; Sg, salivary gland.

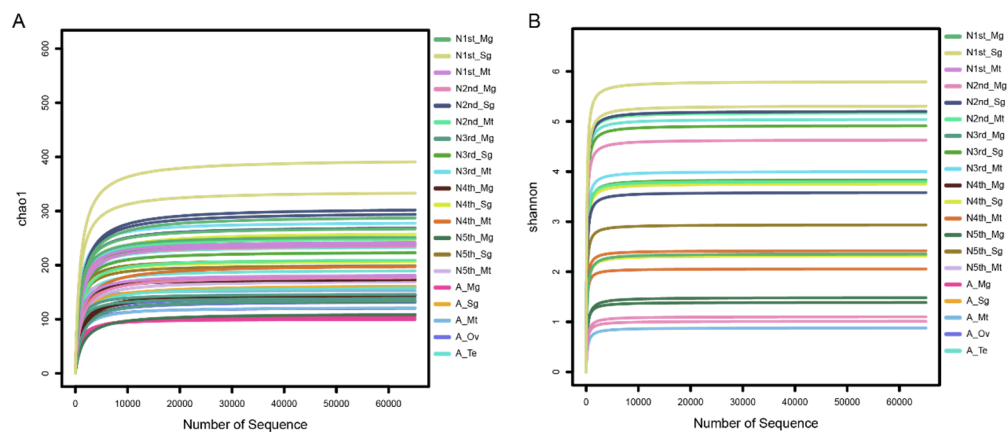


Figure S2. Rarefaction curves showing bacterial richness based on the Chao1 index (A) and bacterial richness and evenness on the Shannon index (B).

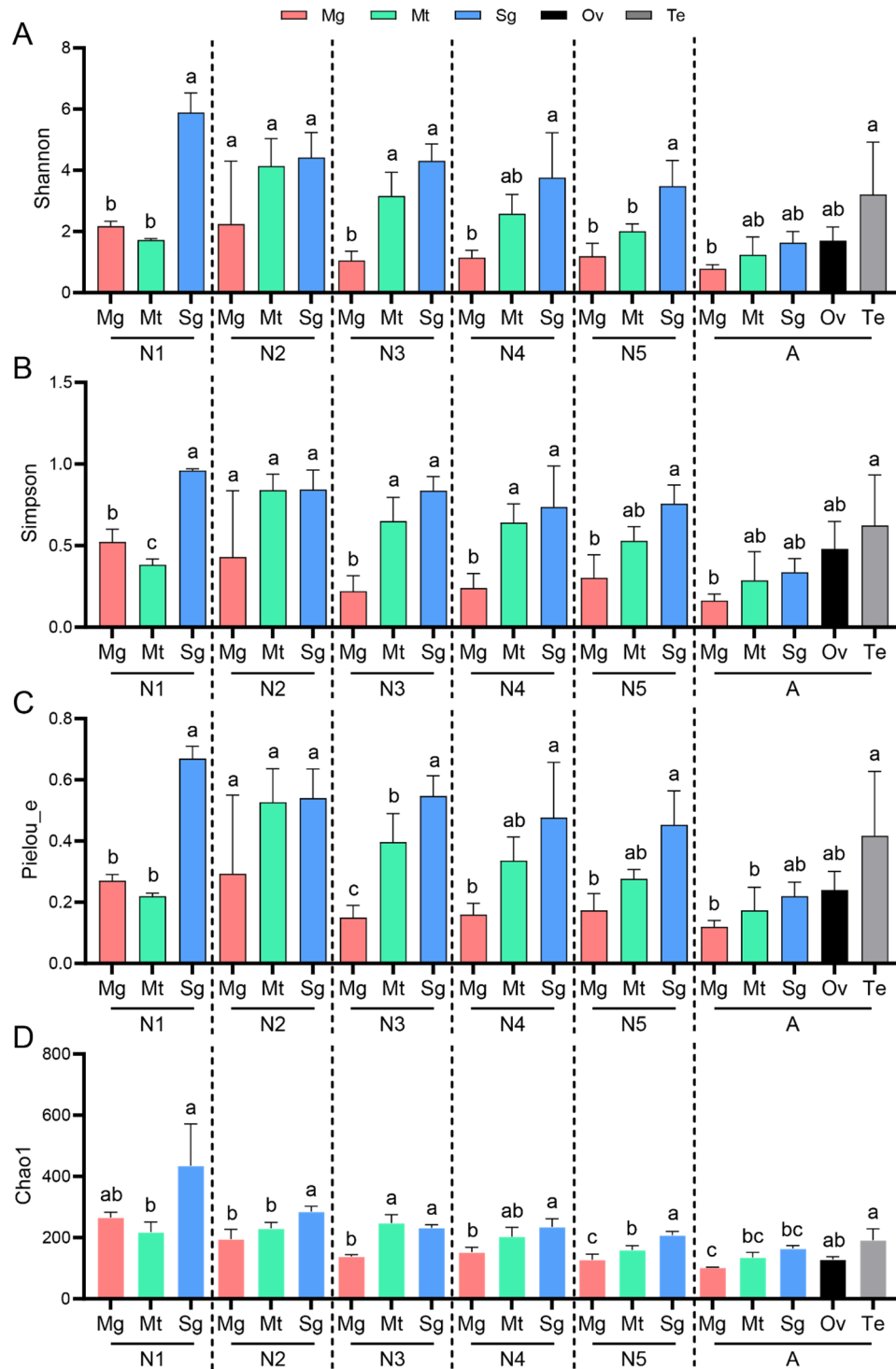


Figure S3. Analysis of alpha diversity among different tissues of leafhoppers at the same developmental stage. Shannon (A) and Simpson (B) indexes represent the species diversity, Pielou_e (C) index represents species evenness, Chao1 (D) species indexes represent species richness. Each bar represents the mean \pm standard error of three replicates. Different lowercase letters indicate significant differences (ANOVA, $p < 0.05$). N1, N2, N3, N4, N5, A refer to 1st-instar nymphs, 2nd-instar nymphs, 3rd-instar nymphs, 4th-instar nymphs, 5th-instar nymphs and adults, respectively.