



Figure S2: Input OTUs used in the network analyses (A), number of network nodes using three different models (B), and number of links in co-occurrence networks (C) of rhizosphere microbiota from samples collected from week 1 and week 3 across the given levels of dilution treatment; Spearman correlations between input OTUs and dilutions were listed under panel; Spearman correlations between nodes (or links) and dilutions in each network were listed in Table S1; Asterisk means a statistically significant ($*P < 0.05$).