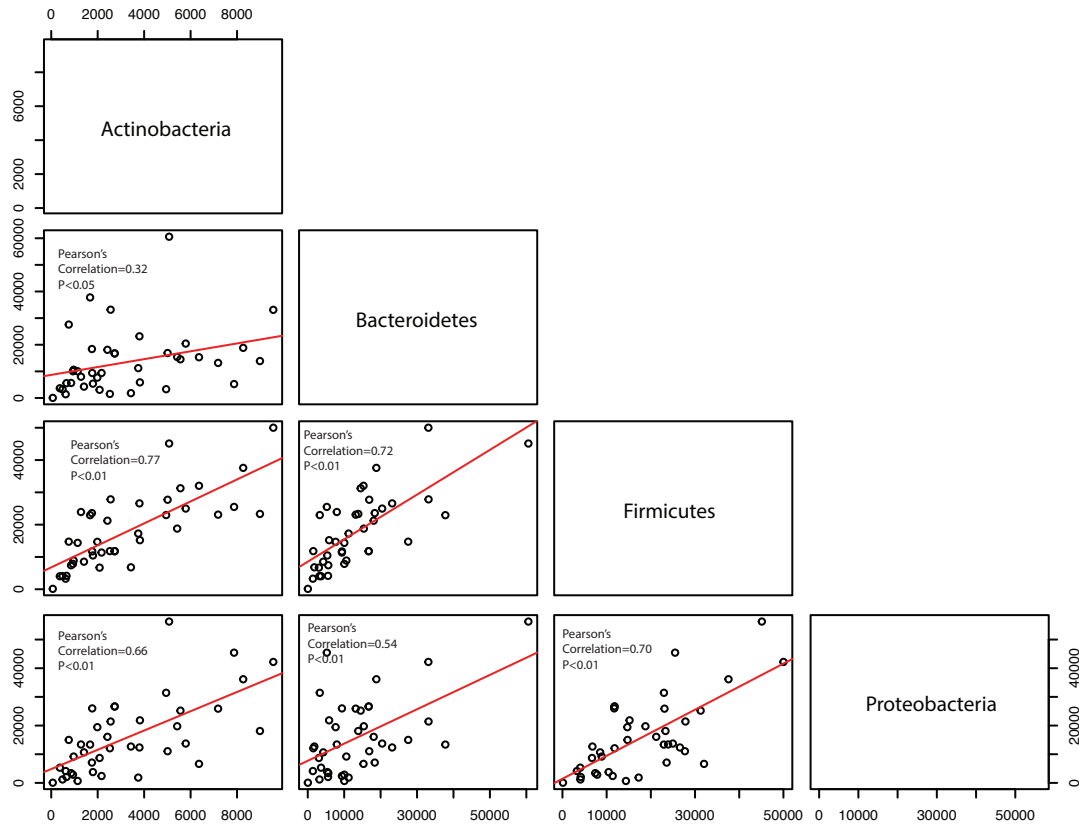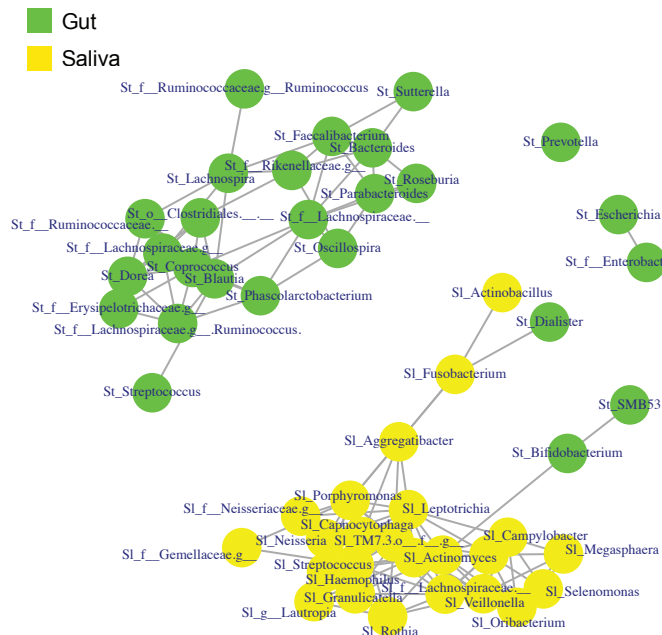# A. Pearson's correlation of salivary Actinobacteria, Bacteroidetes, Firmicutes and Proteobacteria co-occurrence cluster

B. Gut and oral microbiome genus level SparCC co-occurrence network, Controls (Corr>0.5)

C. Gut and oral microbiome genus level SparCC co-occurrence network, ASD (Corr>0.5)