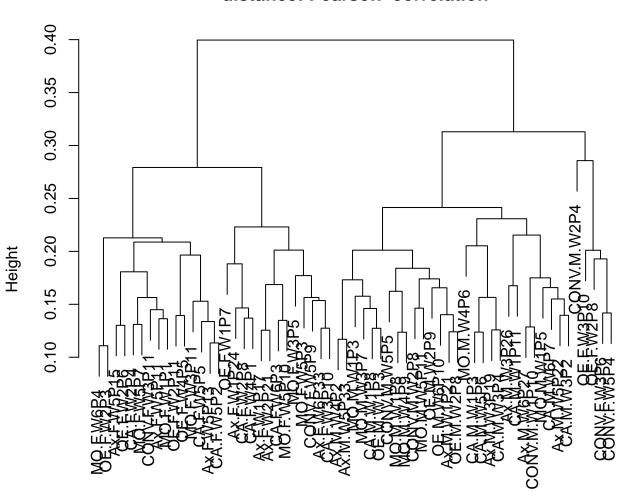
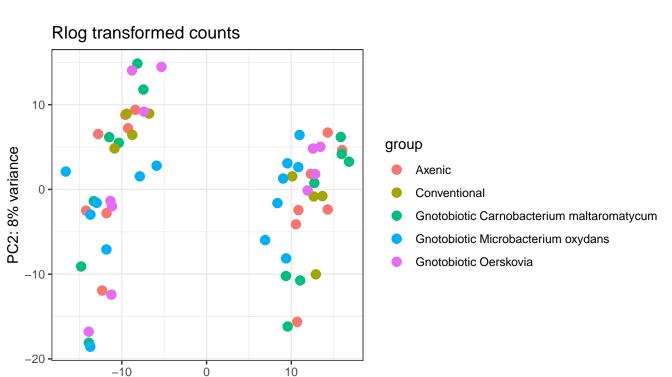
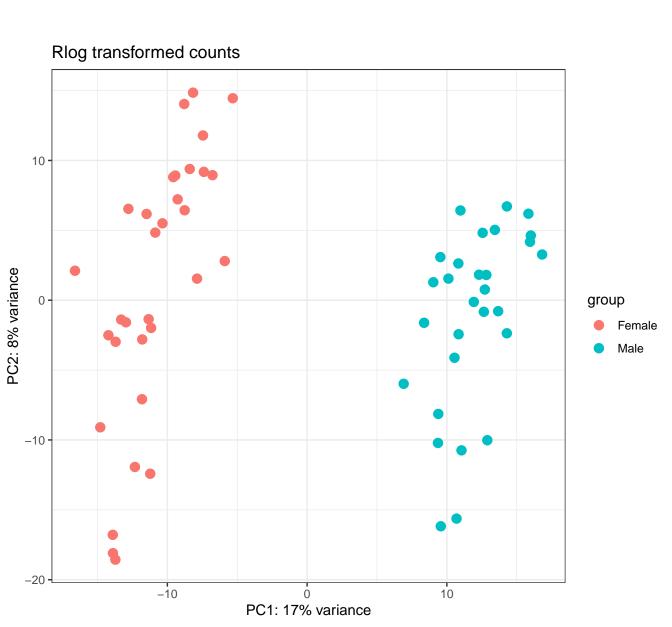
rlog transformed read counts distance: Pearson correlation

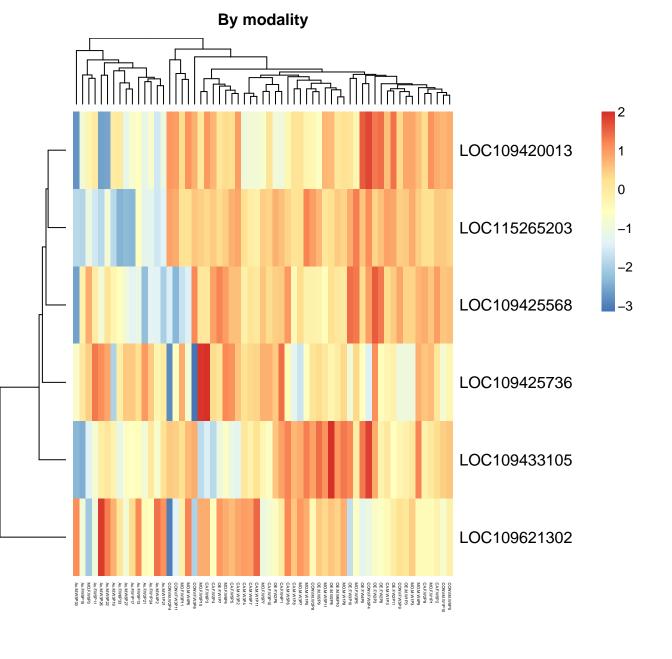


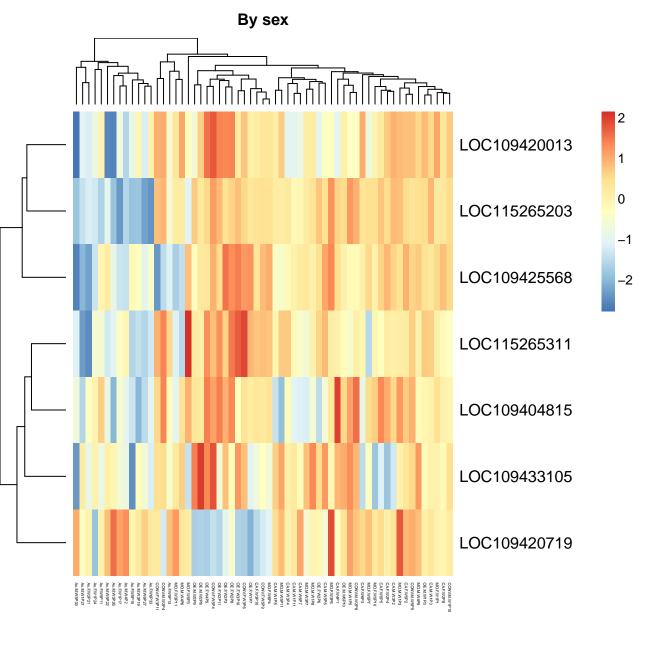
distance.m_rlog
hclust (*, "complete")

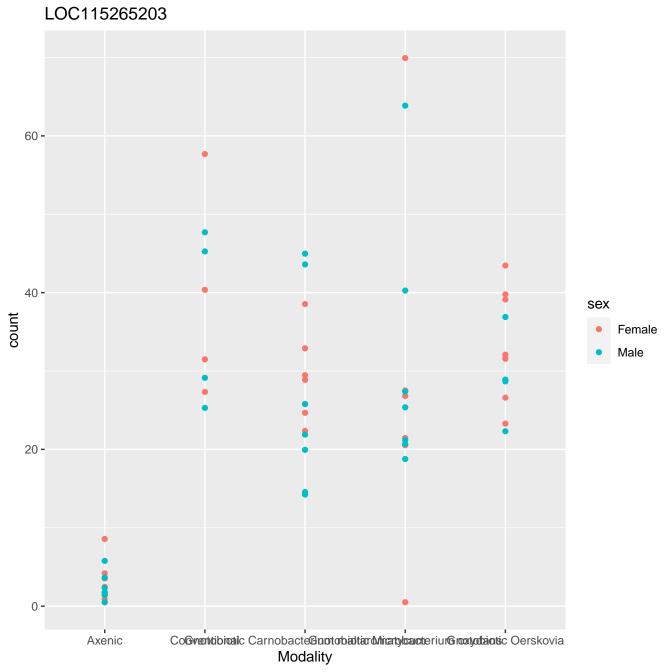


PC1: 17% variance









MA plot: group1 vs group2 (padj < 0.05)

