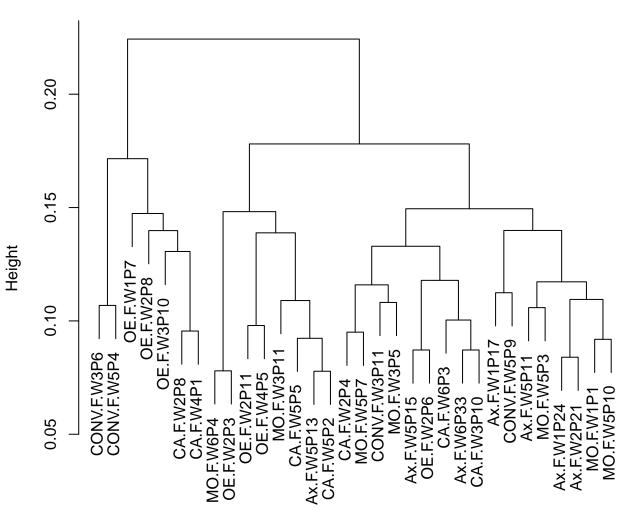
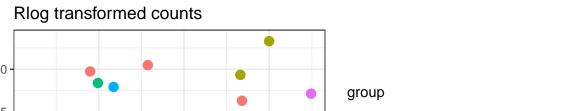
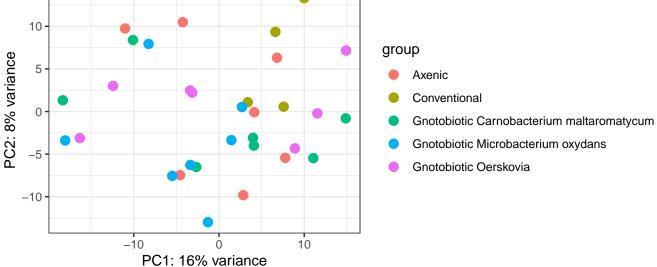
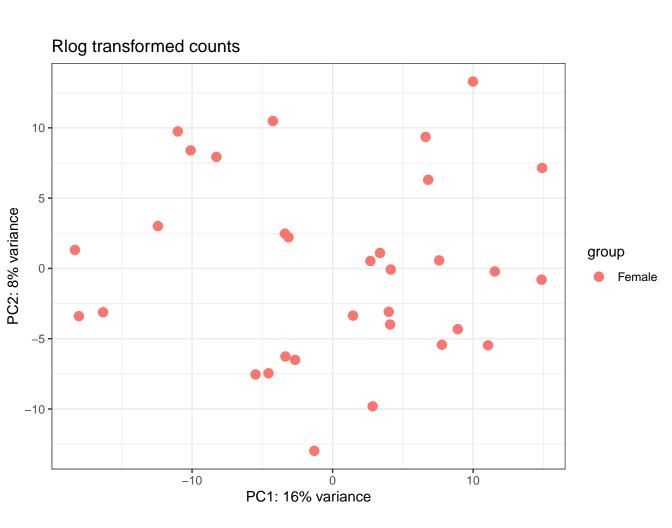
## rlog transformed read counts distance: Pearson correlation

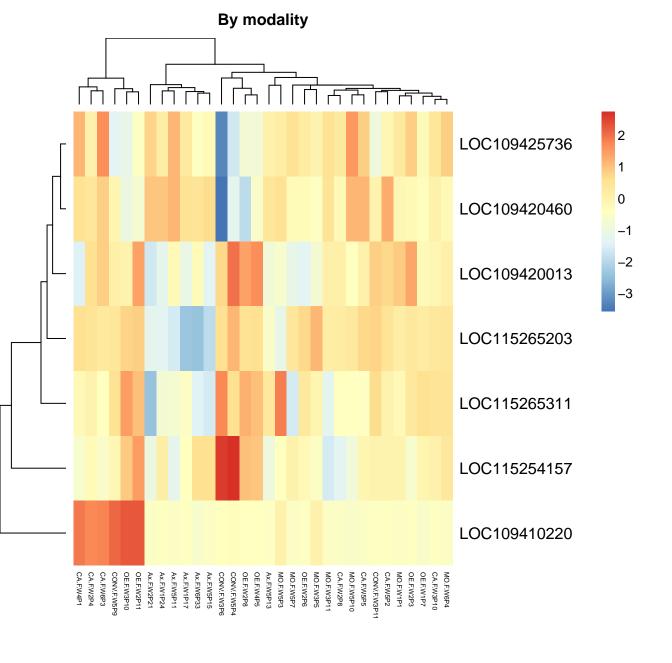


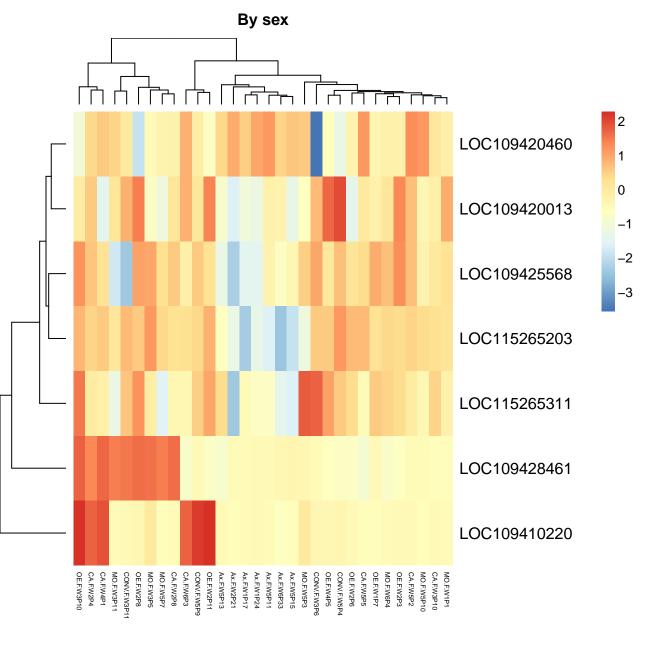
distance.m\_rlog hclust (\*, "complete")

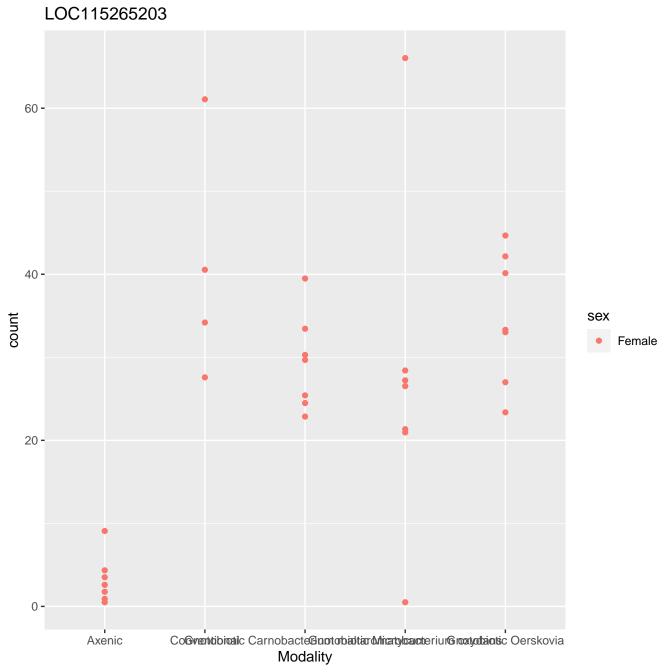












## MA plot: group1 vs group2 (padj < 0.05)

