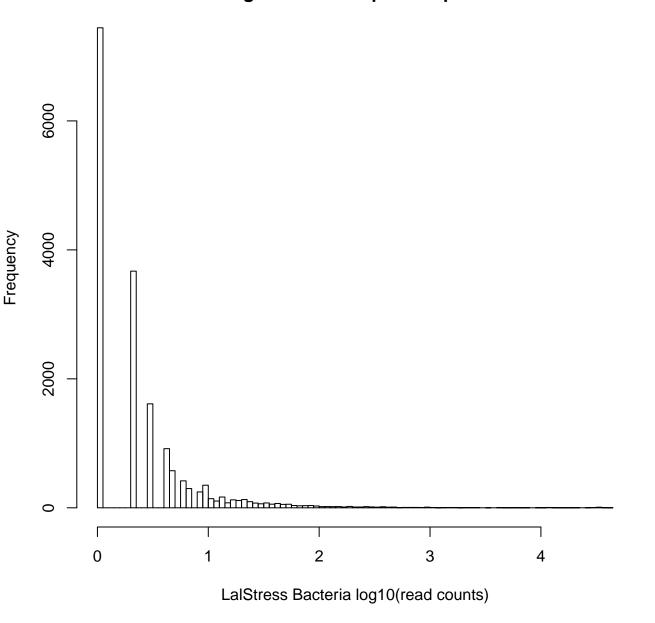
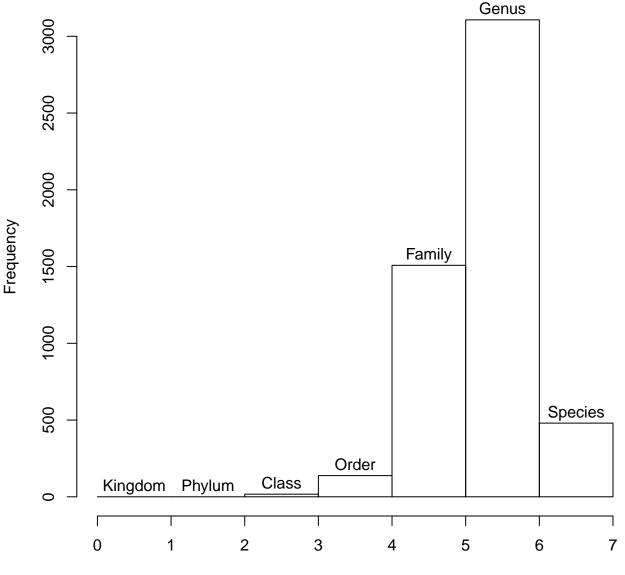
LalStress Bacteria
Histogram of reads per sample/OTU



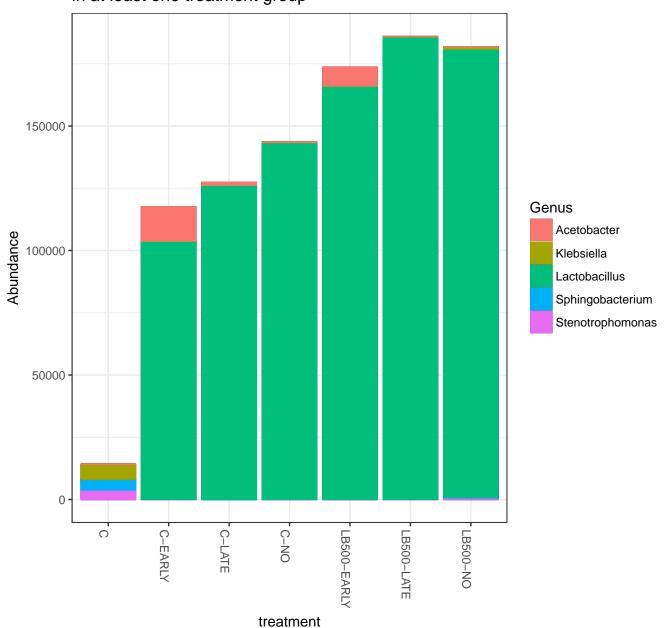
## LalStress Bacteria Histogram of INFERRED most specific rank

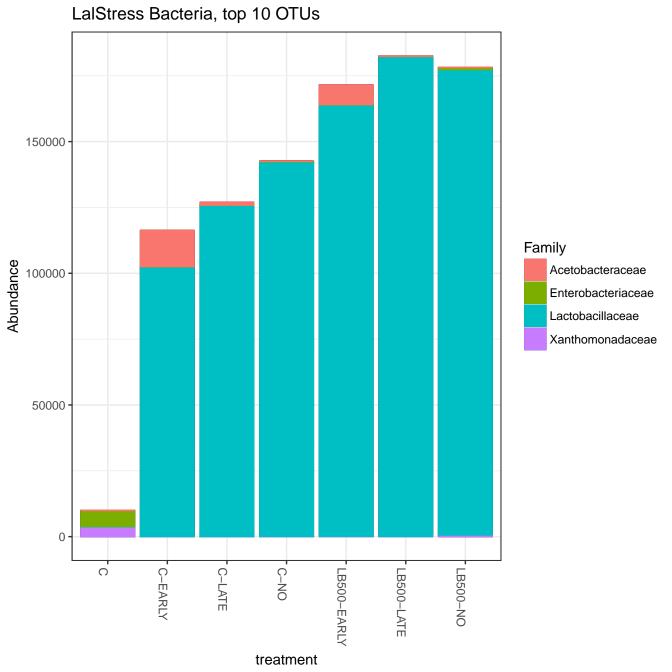


Number of non "uncultured" and "unidentified" phylogenetic levels

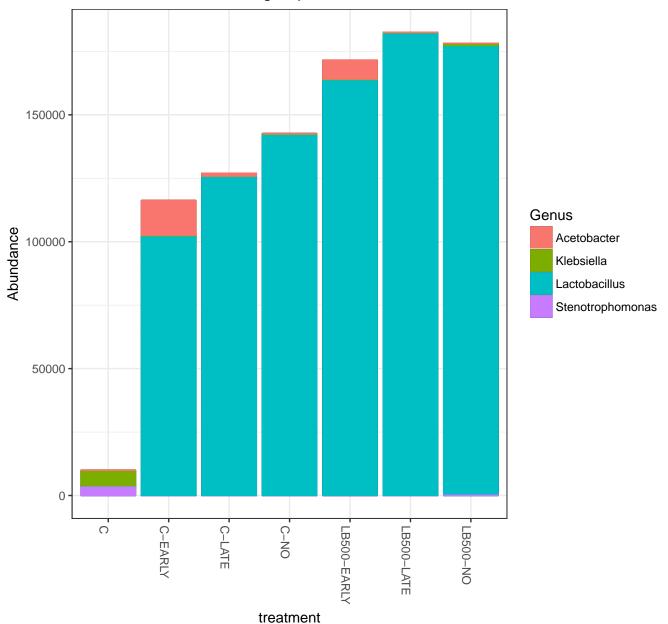
LalStress Bacteria, top 10 OTUs 150000 -Family Acetobacteraceae Abundance 100000 -Enterobacteriaceae Lactobacillaceae Sphingobacteriaceae Xanthomonadaceae 50000 -0 C C-EARLY LB500-EARLY LB500-LATE treatment

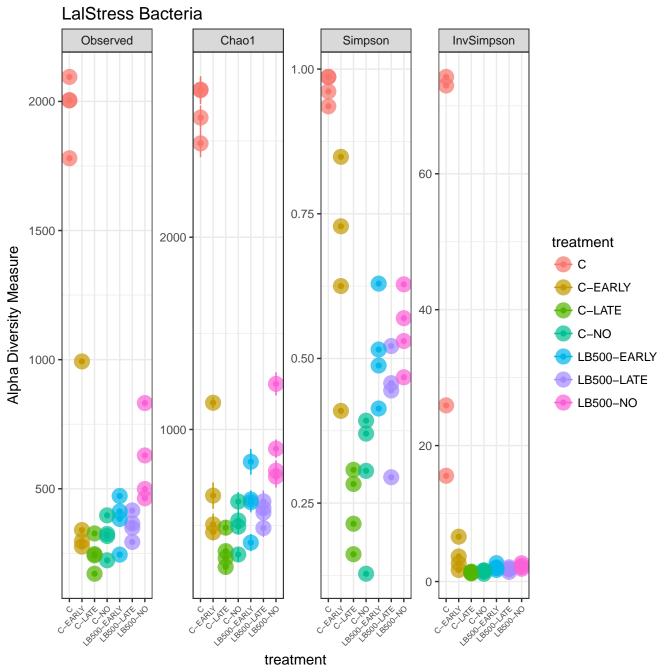
LalStress Bacteria, OTUs with at least 1/100 of max reads/sample/OTU in at least one treatment group

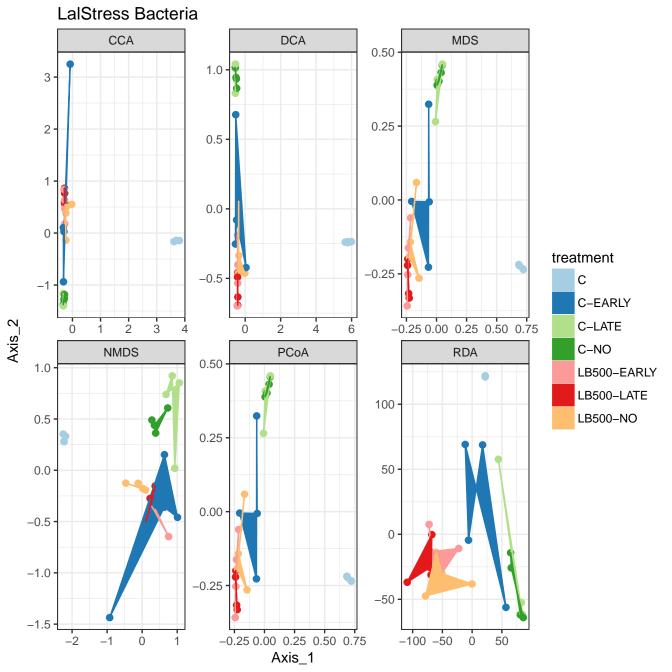


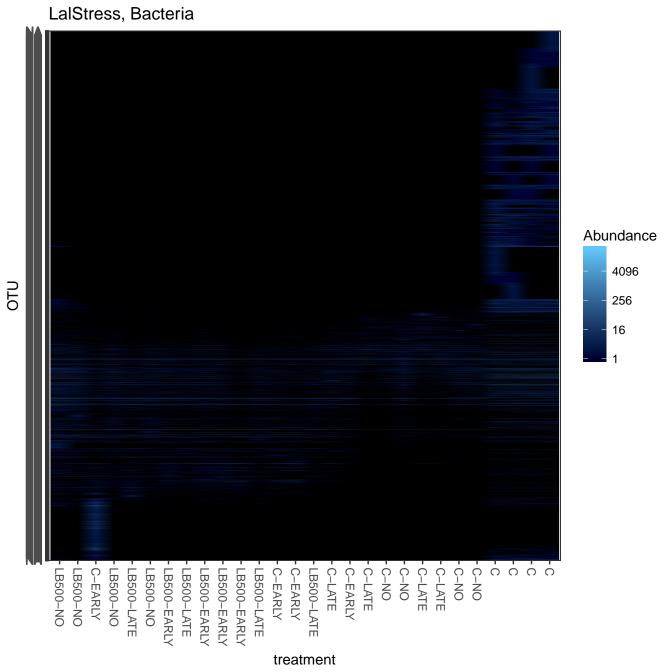


LalStress Bacteria, OTUs with at least 1/100 of max reads/sample/OTU in at least one treatment group









LalStress Bacteria: Differential abundance ratio, C vs. LB500-EARLY @ 0.05 5 0 Phylum log2FoldChange Actinobacteria Proteobacteria **Bacteroidetes** Firmicutes -5· -10 Bacilli Actinobacteria Betaproteobacteria Sphingobacteriia Gammaproteobacteria Alphaproteobacteria Class