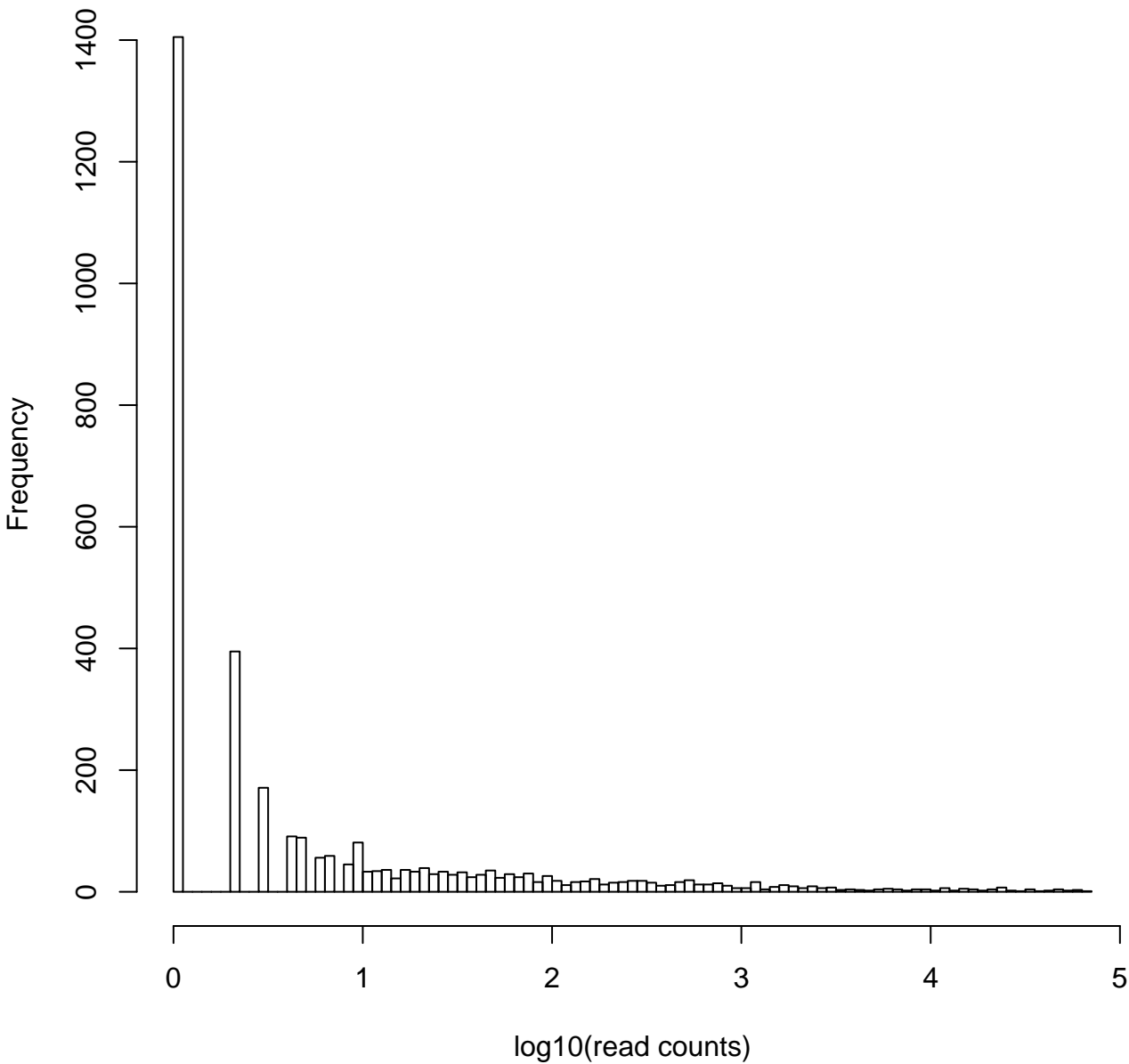
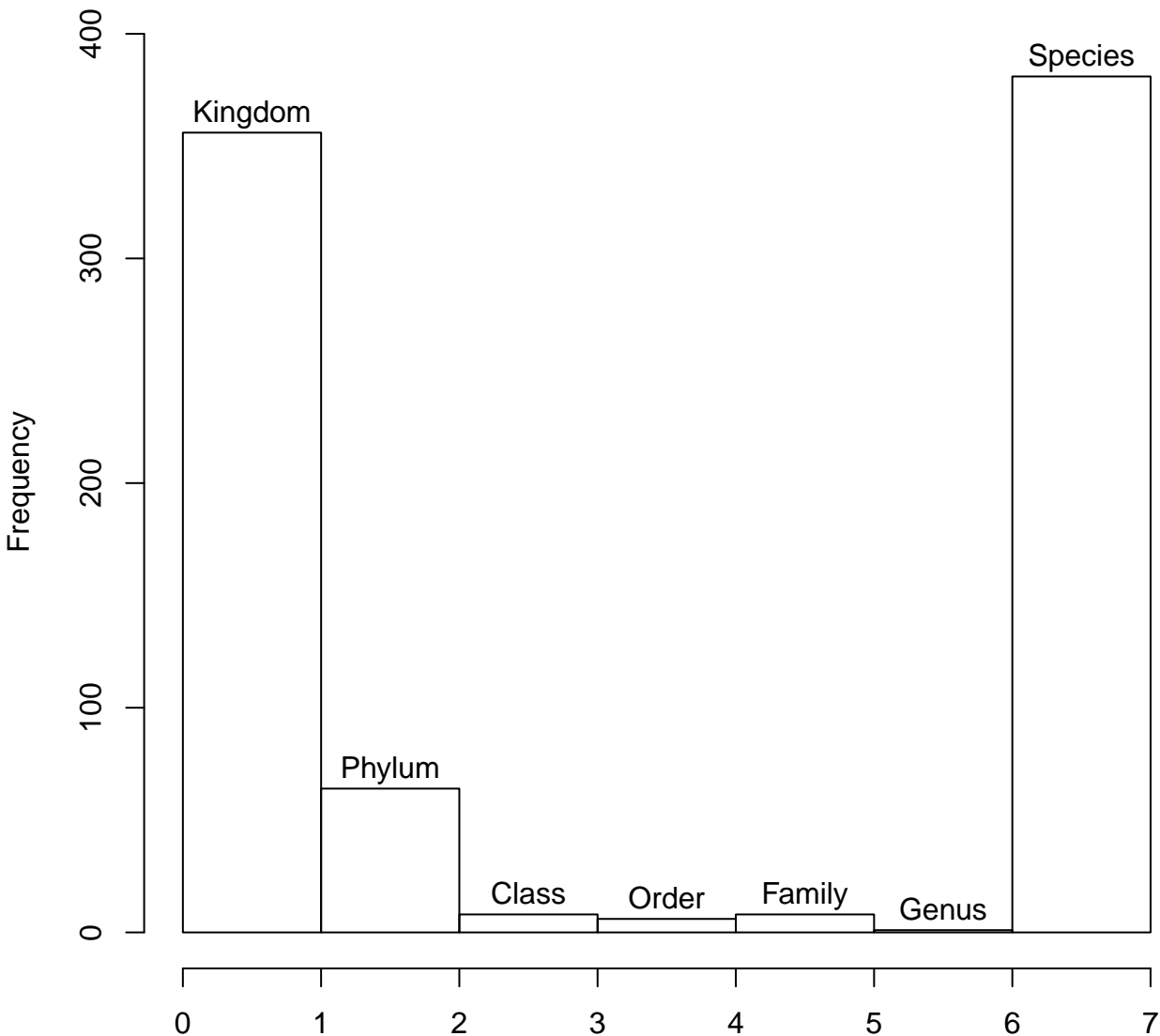


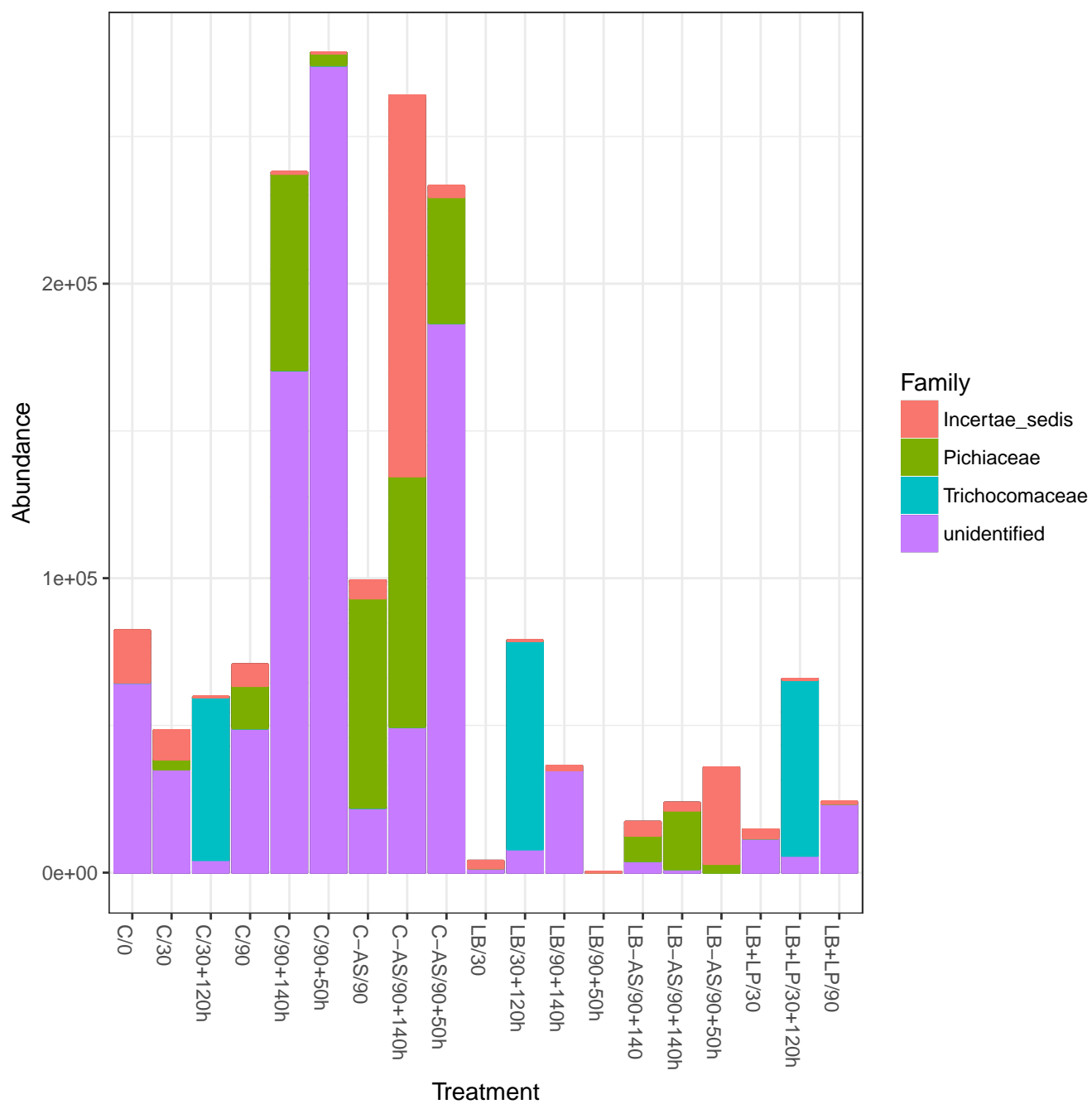
Histogram of reads per sample/OTU

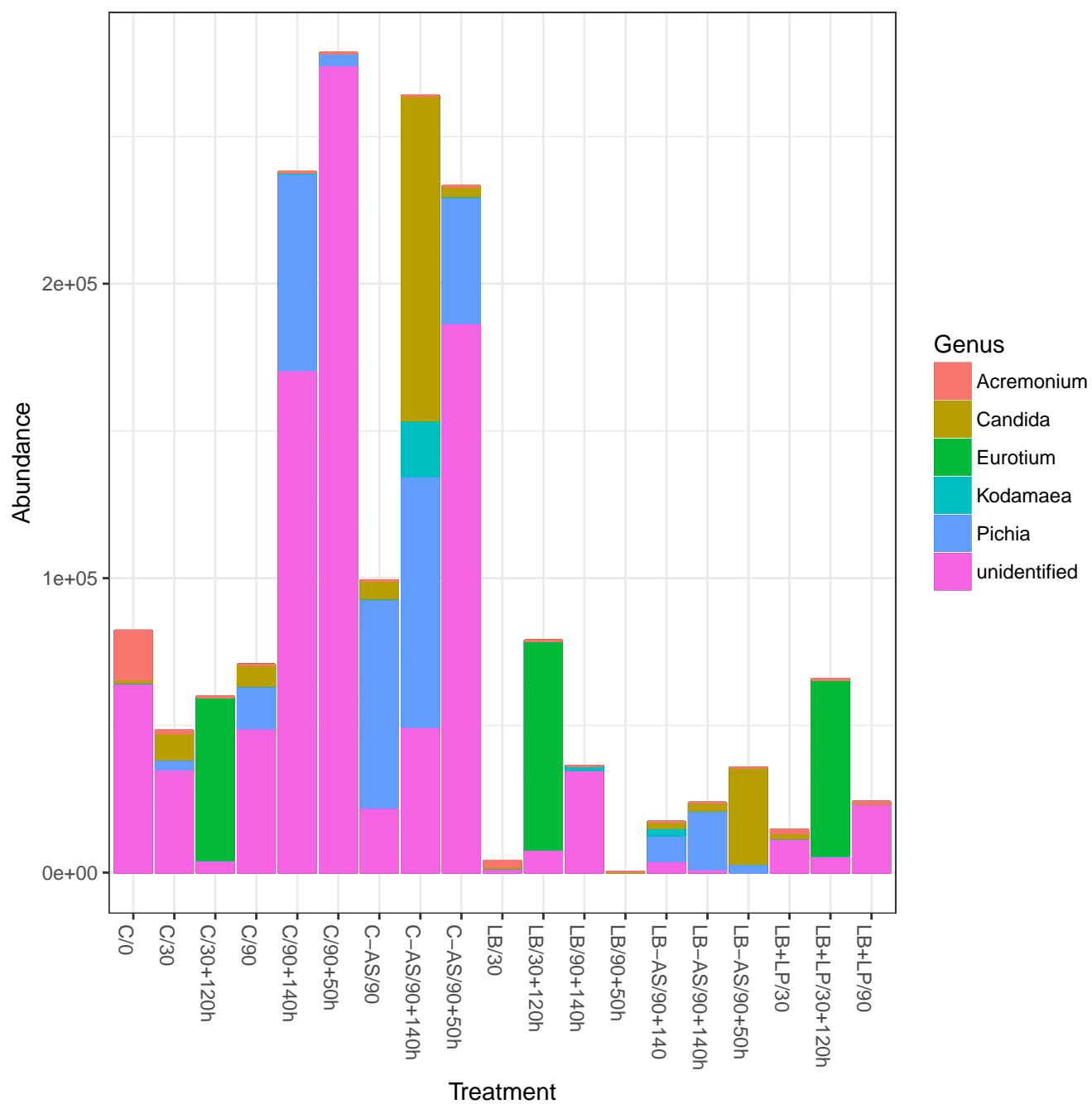


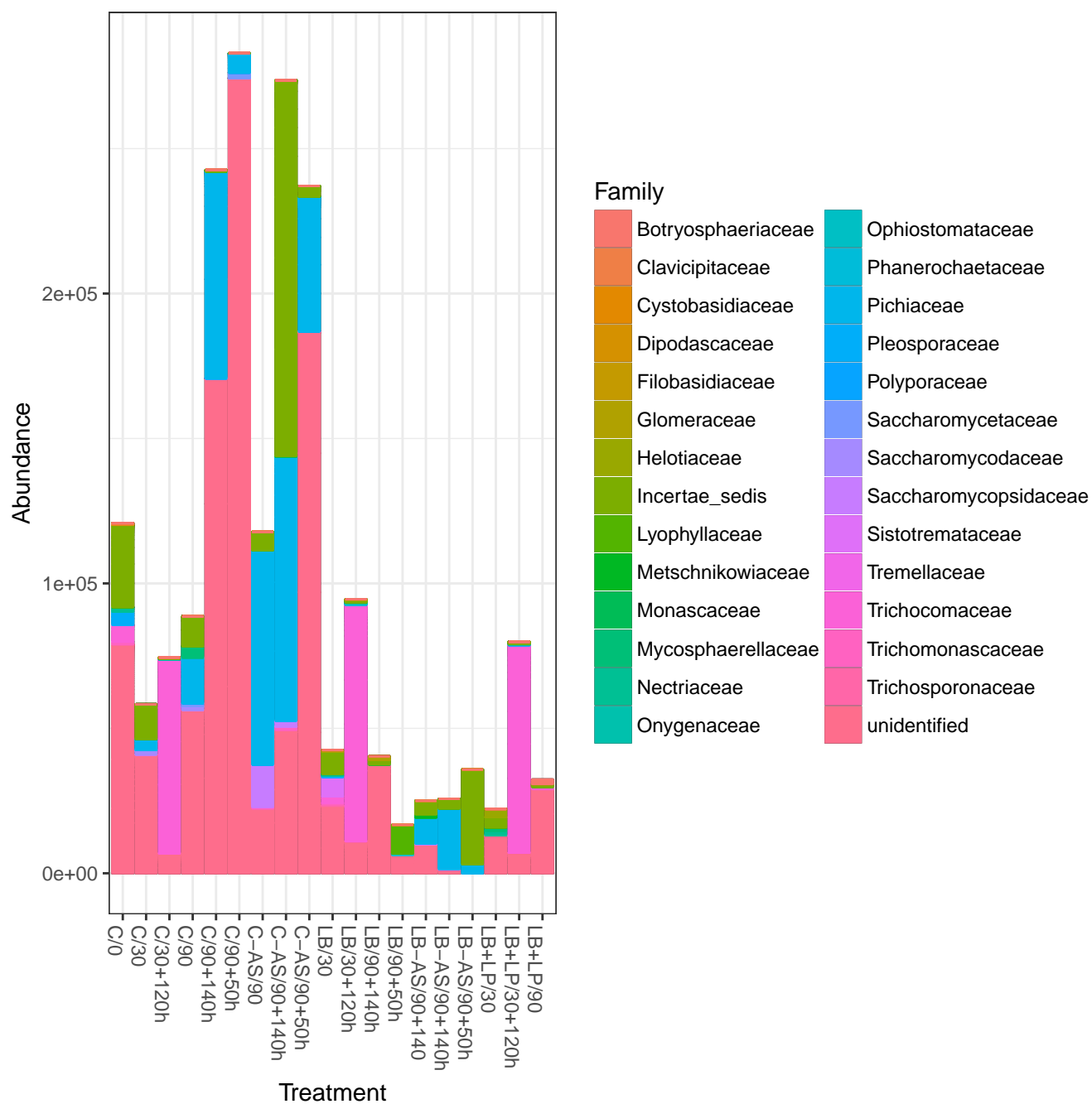
Histogram of INFERRED most specific rank

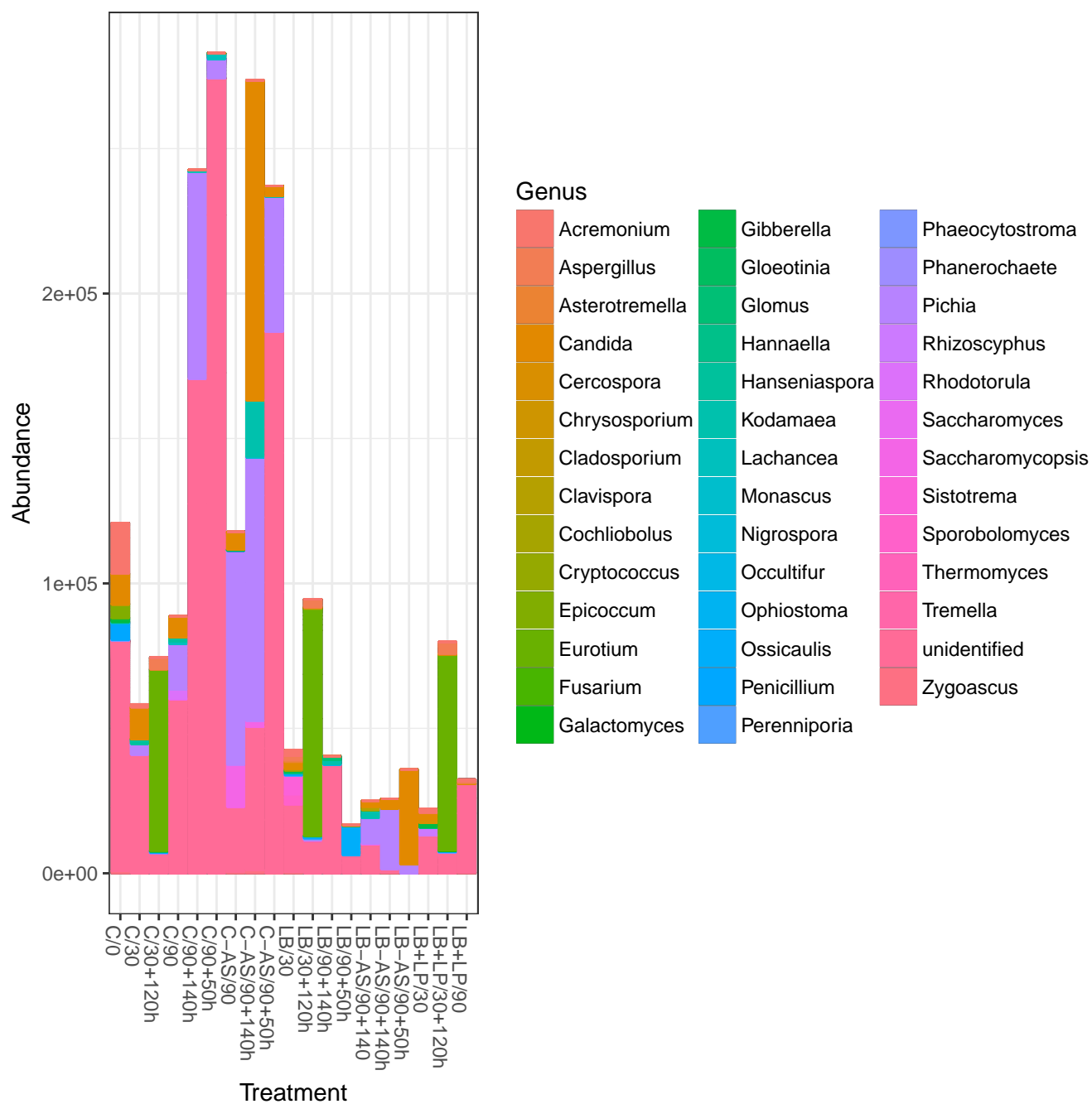


Number of non "uncultured" and "unidentified" phylogenetic levels

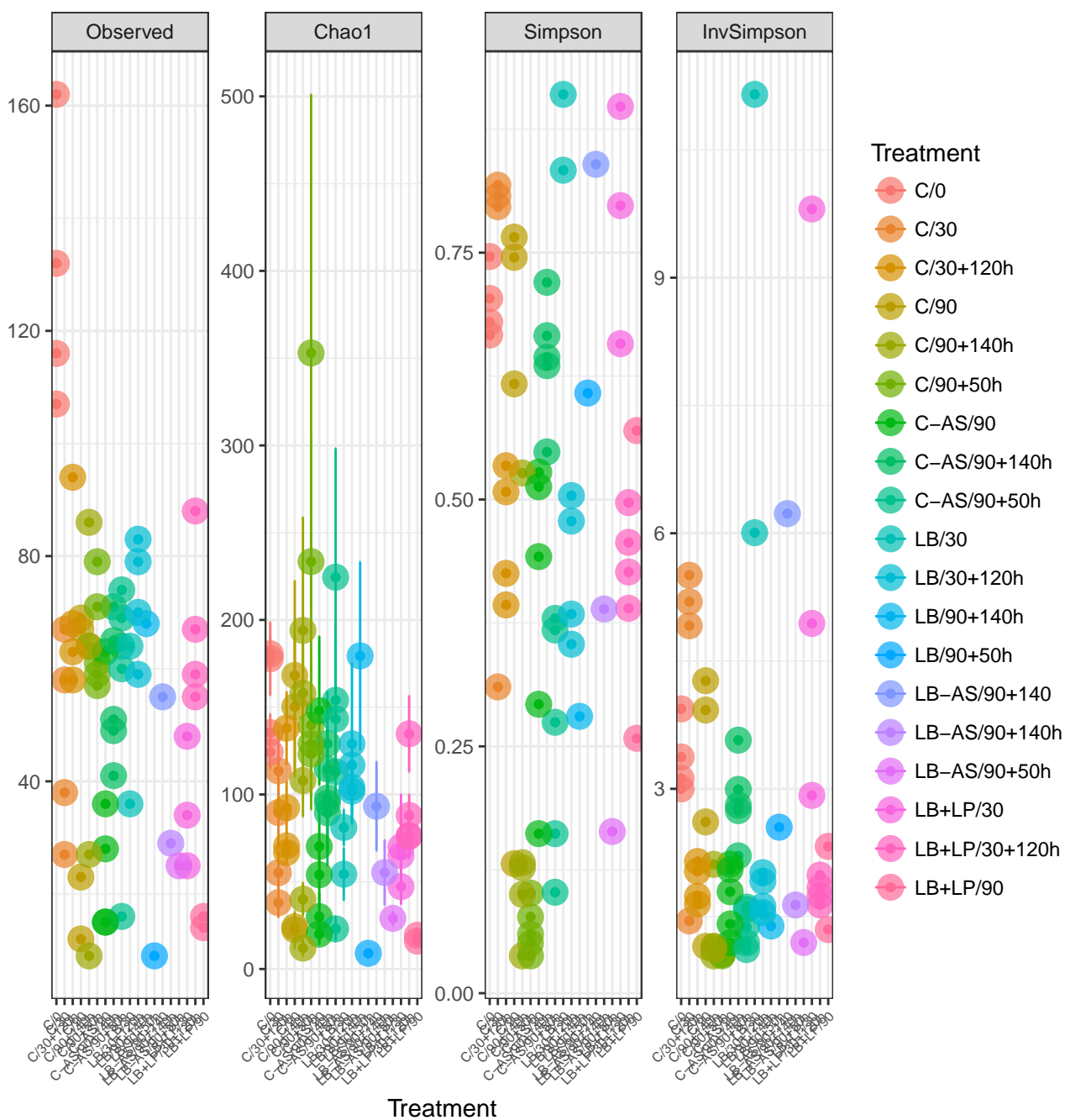




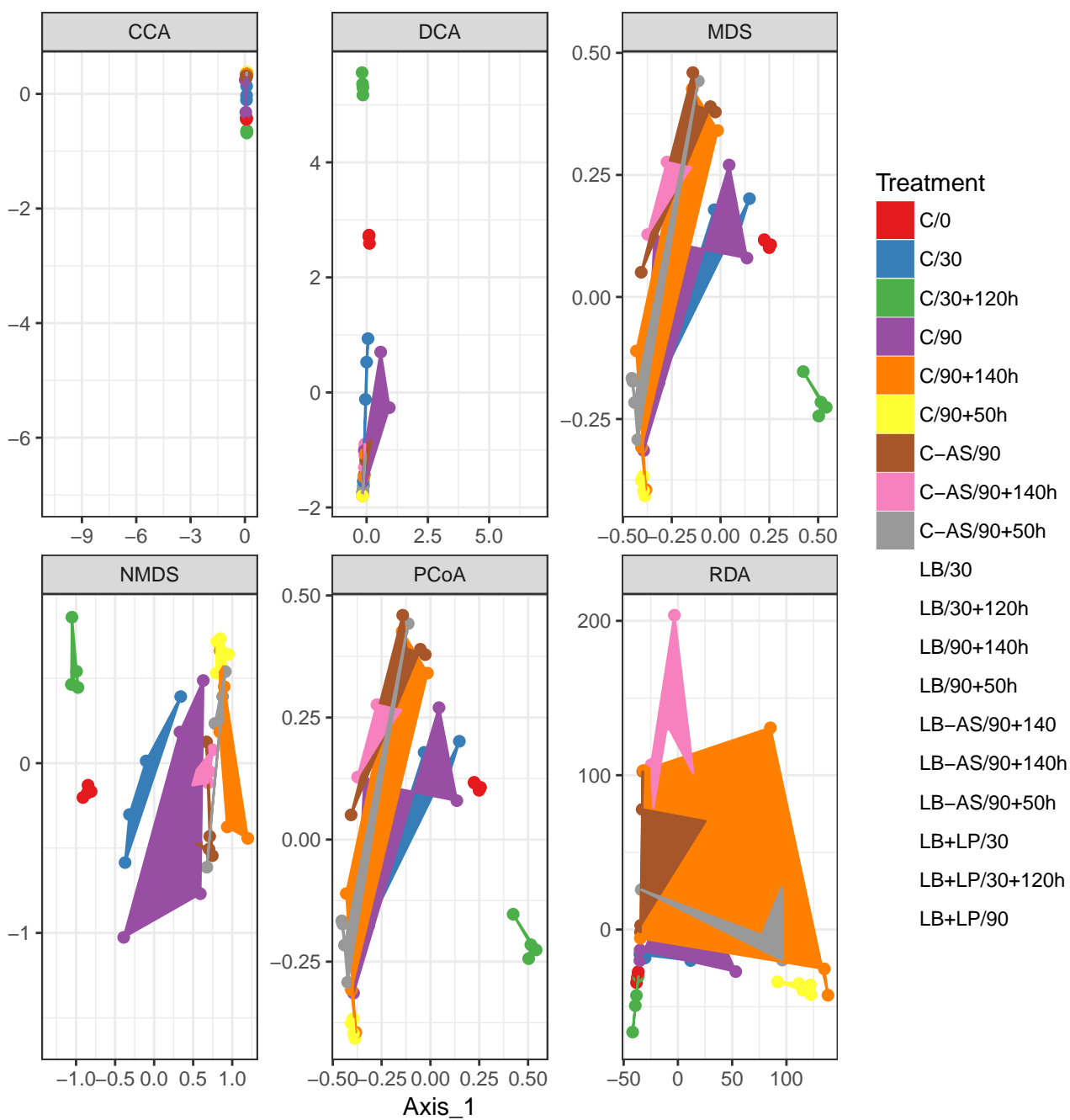




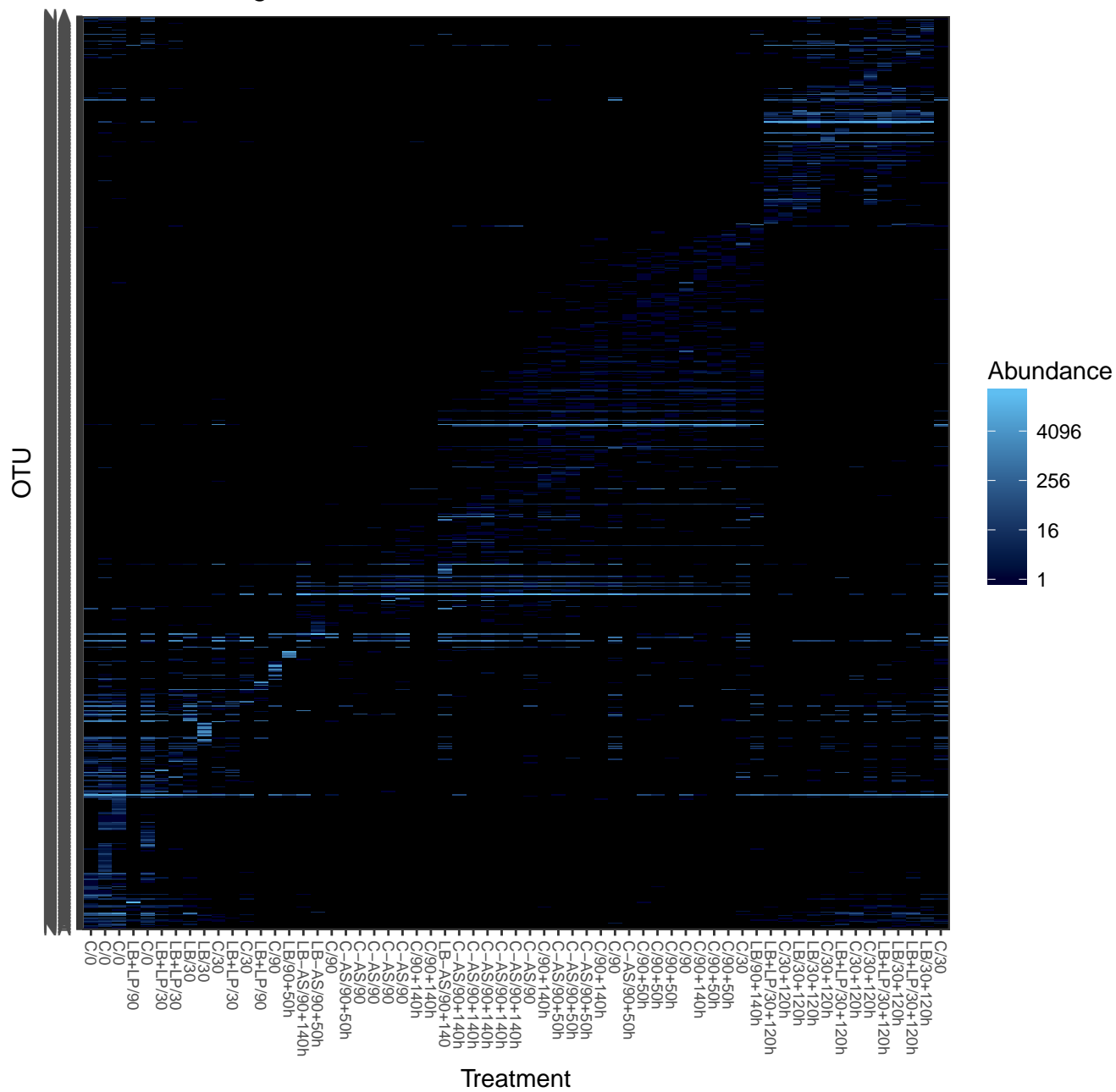
Alpha Diversity Measure



Axis_2



V-HMC, Fungi



Differential abundance ratio, C/0 vs. LB-AS/90+50h

