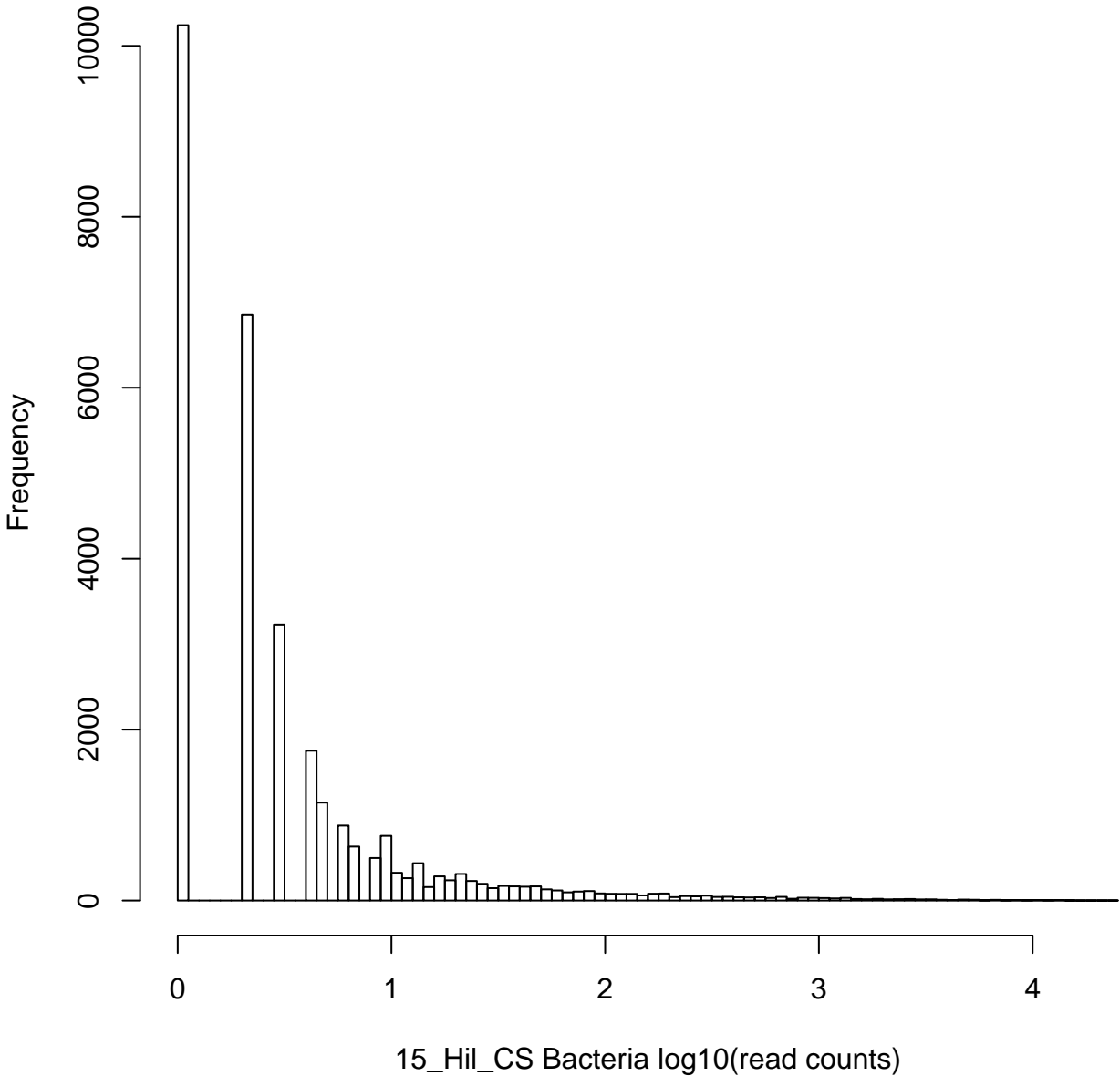
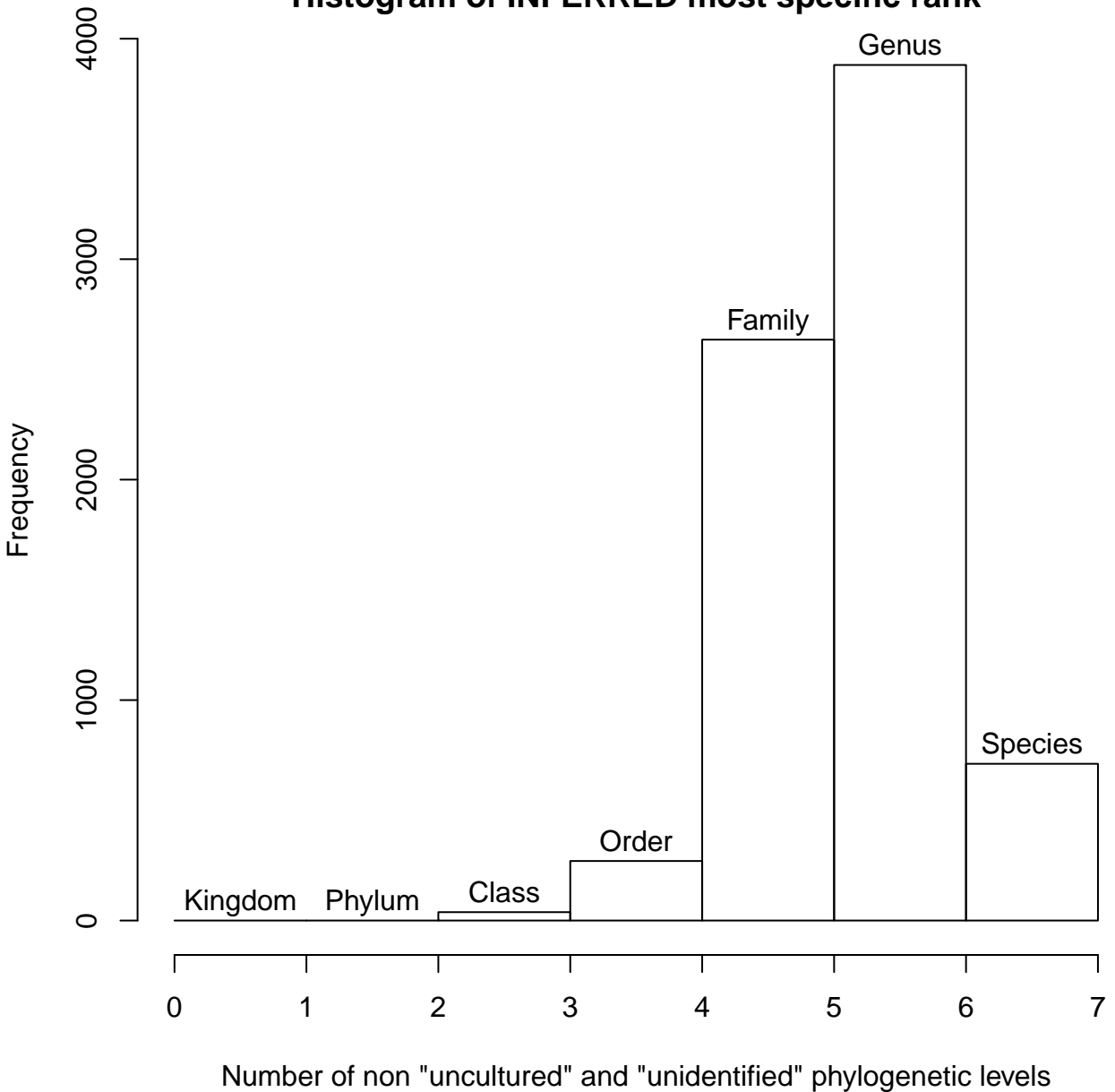


15_Hil_CS Bacteria
Histogram of reads per sample/OTU

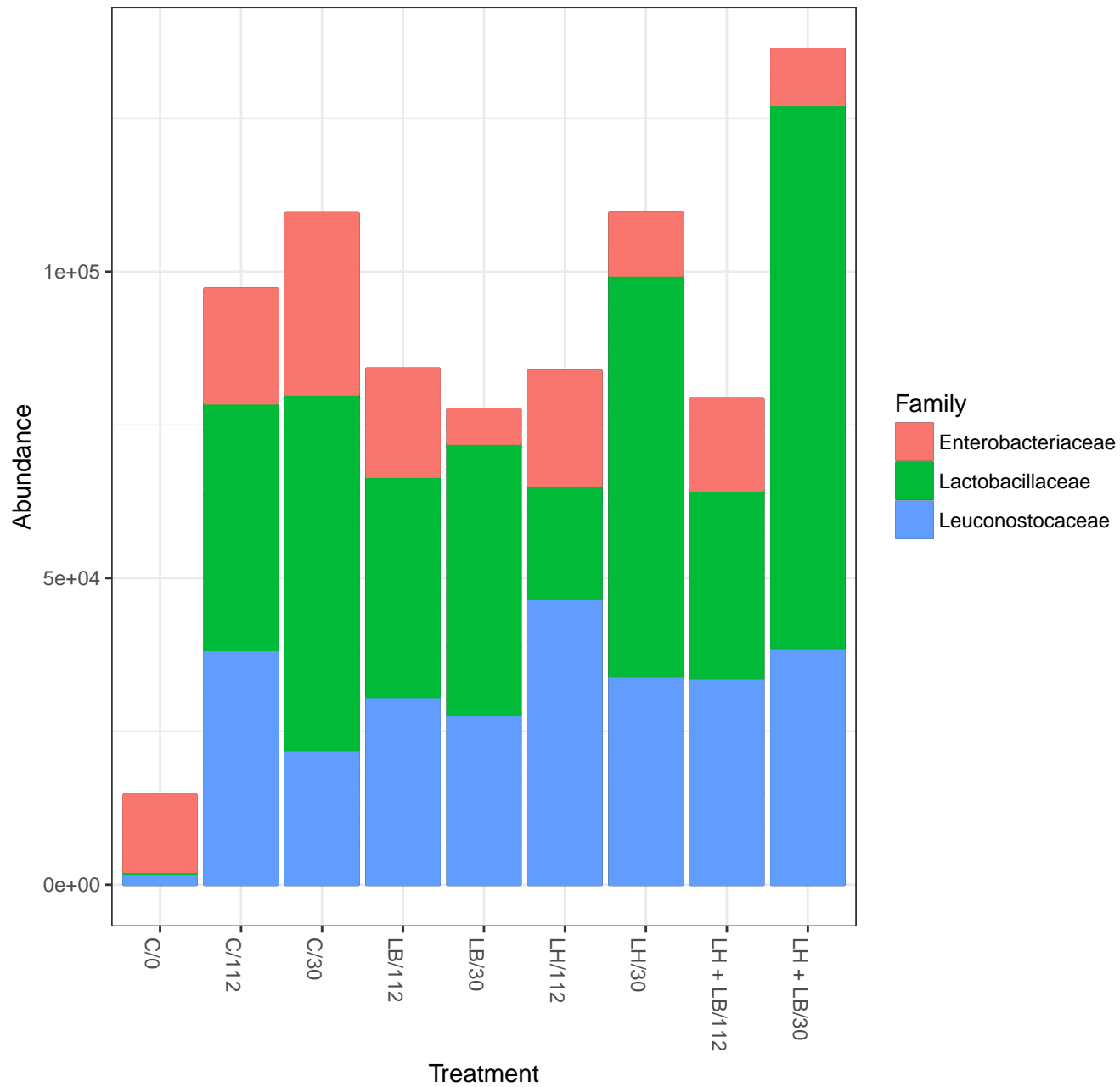


15_Hil_CS Bacteria

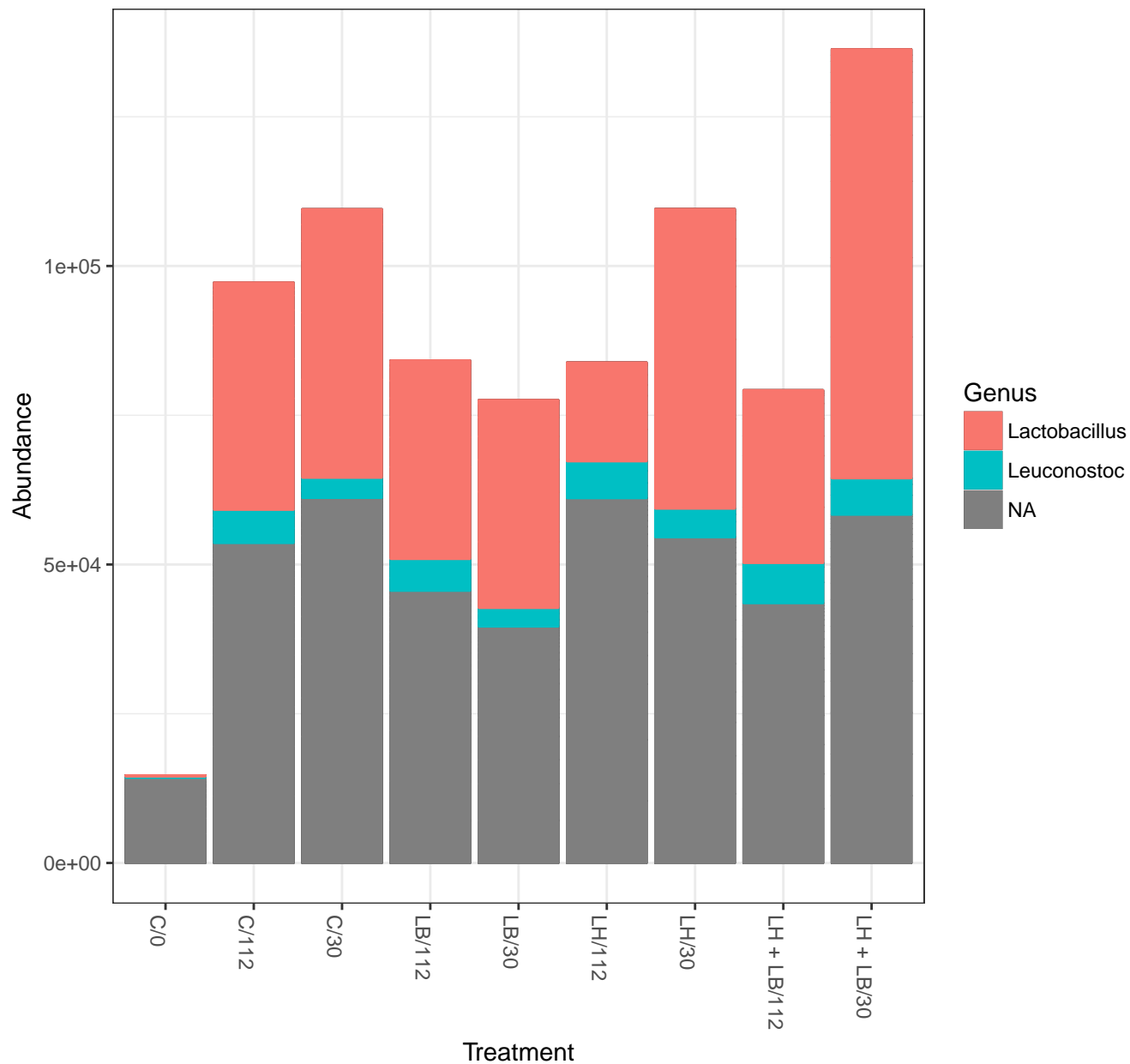
Histogram of INFERRED most specific rank



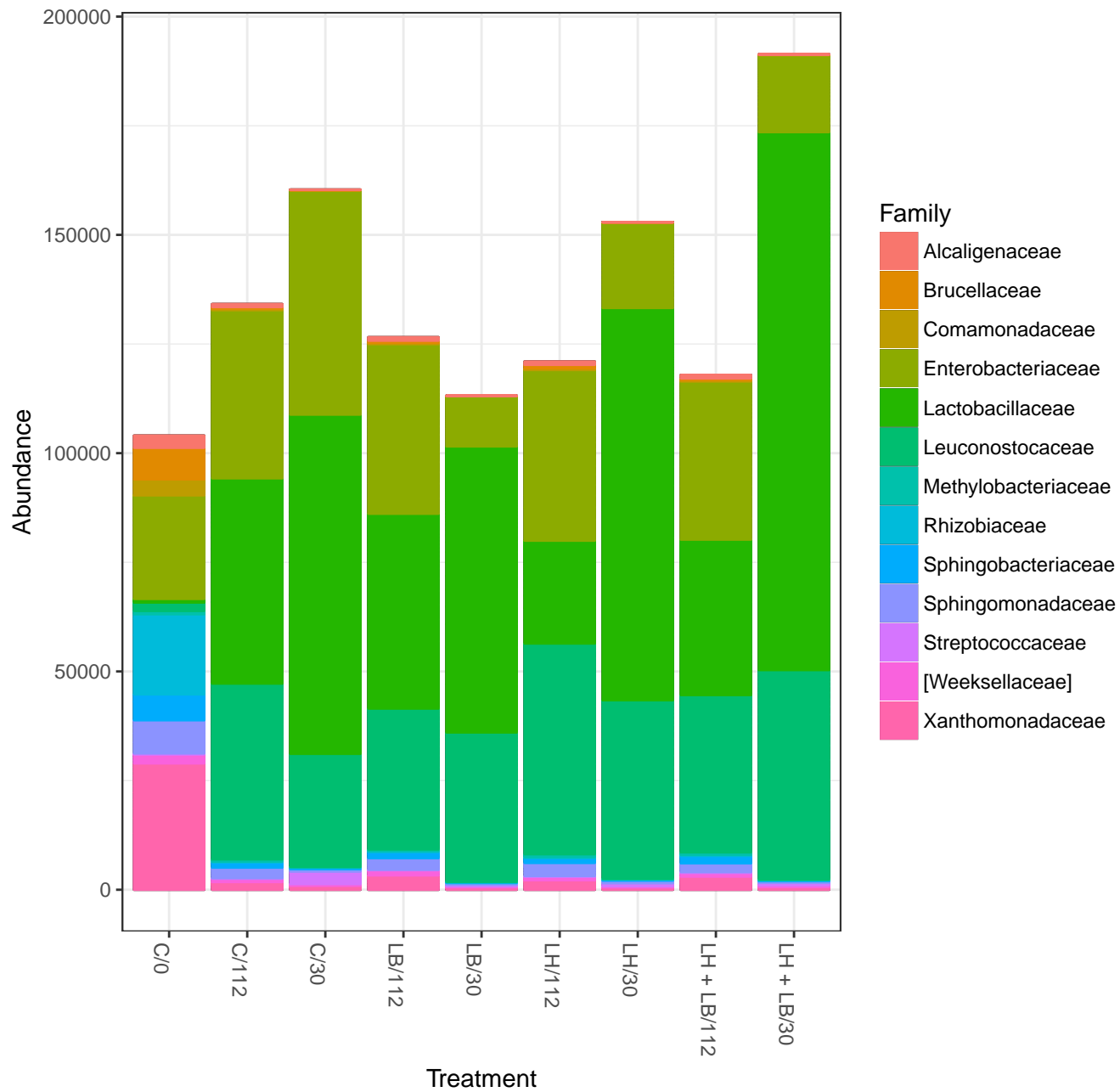
15_Hil_CS Bacteria, top 10 OTUs



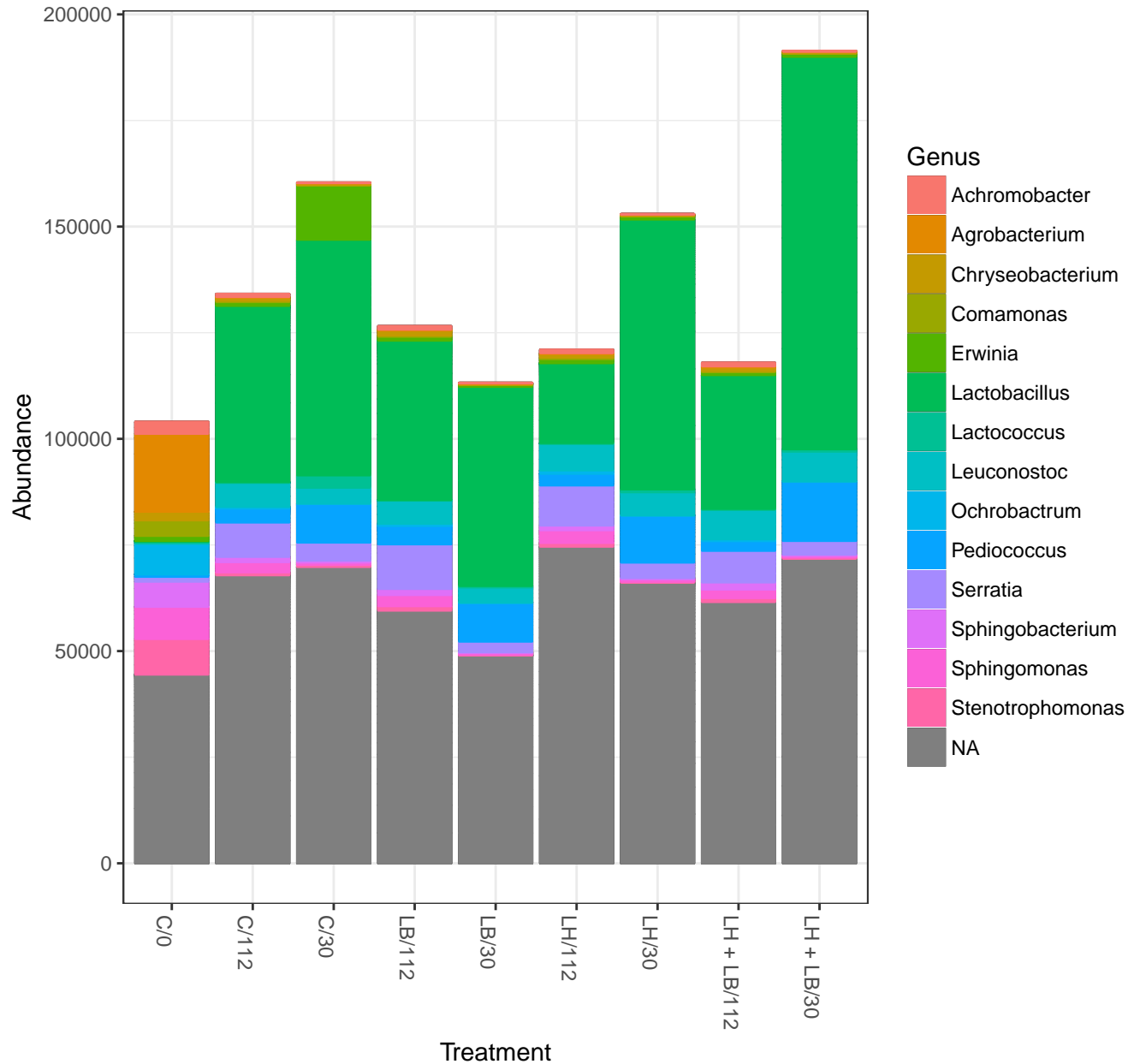
15_Hil_CS Bacteria, OTUs with at least 1/100 of max reads/sample/OTU
in at least one treatment group



15_Hil_CS Bacteria, top 10 OTUs

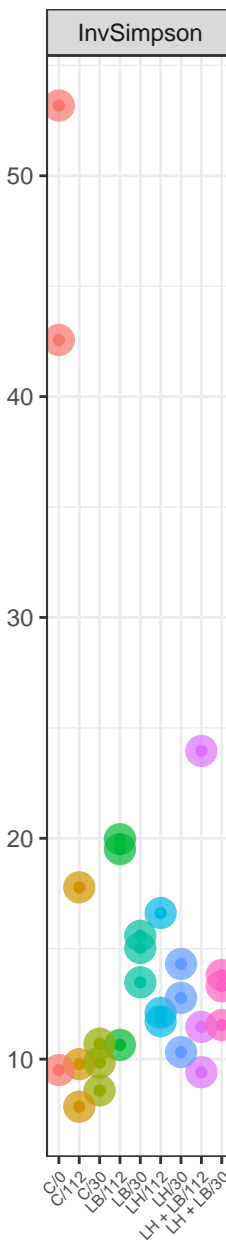
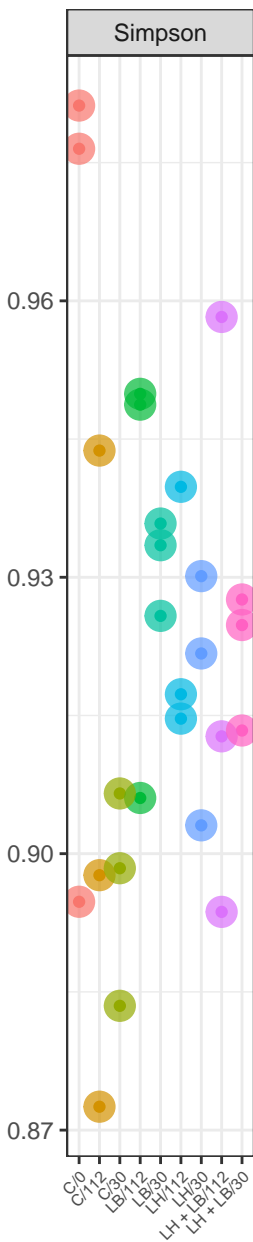
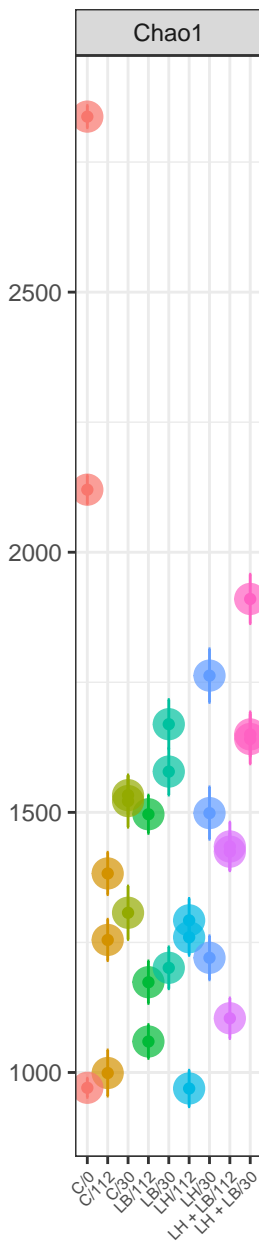
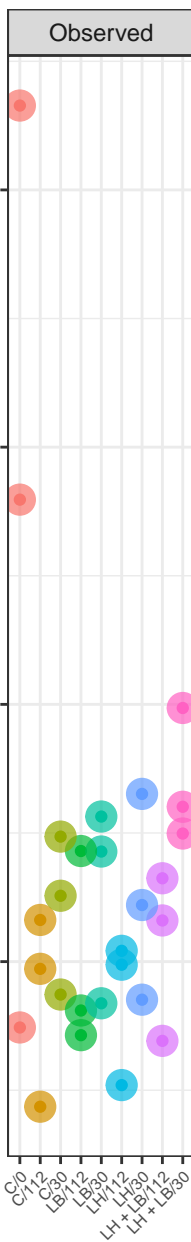


15_Hil_CS Bacteria, OTUs with at least 1/100 of max reads/sample/OTU
in at least one treatment group



15_Hil_CS Bacteria

Alpha Diversity Measure



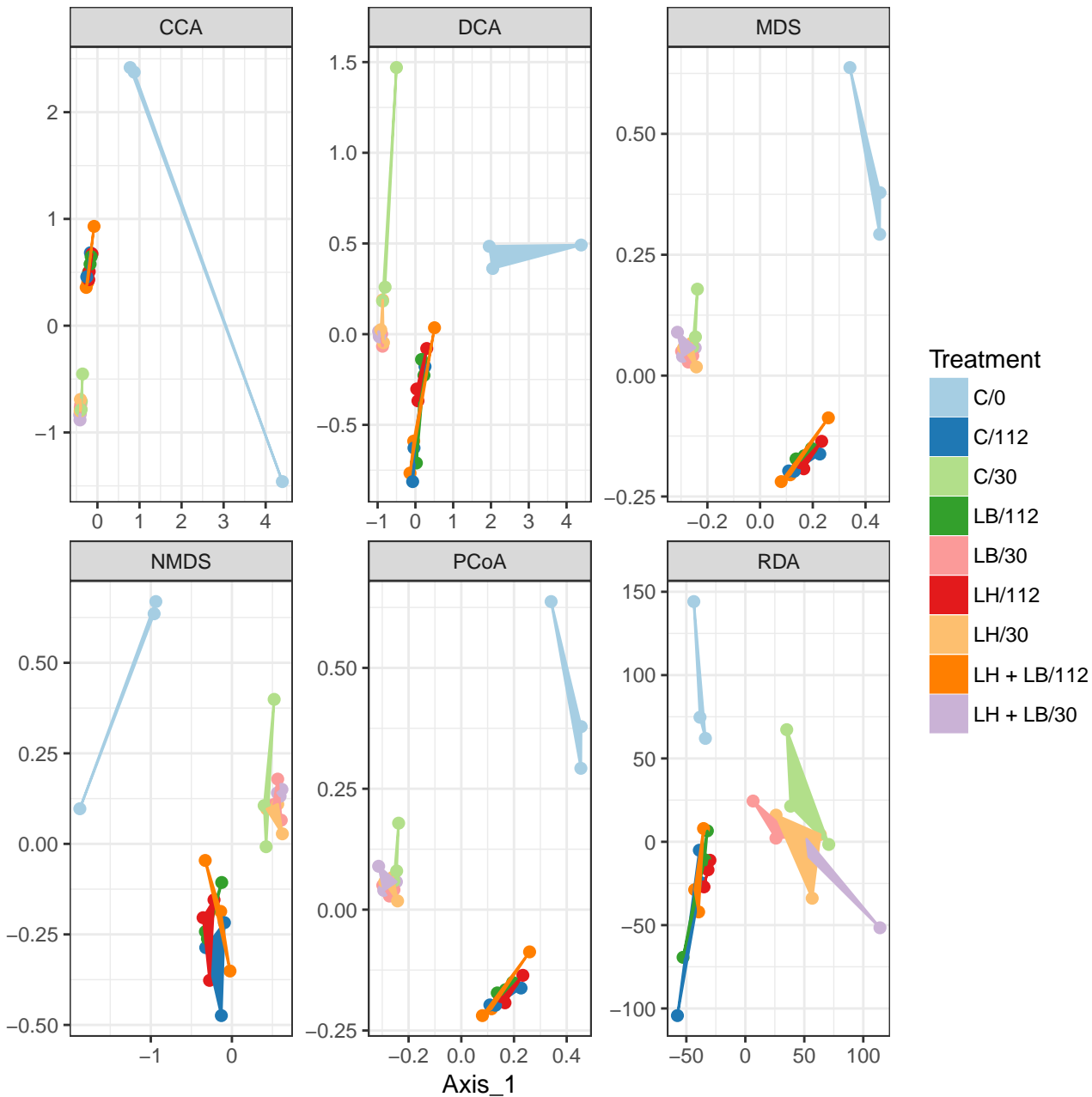
Treatment

- C/0
- C/12
- C/30
- LB/12
- LB/30
- LH/12
- LH/30
- LH + LB/12
- LH + LB/30

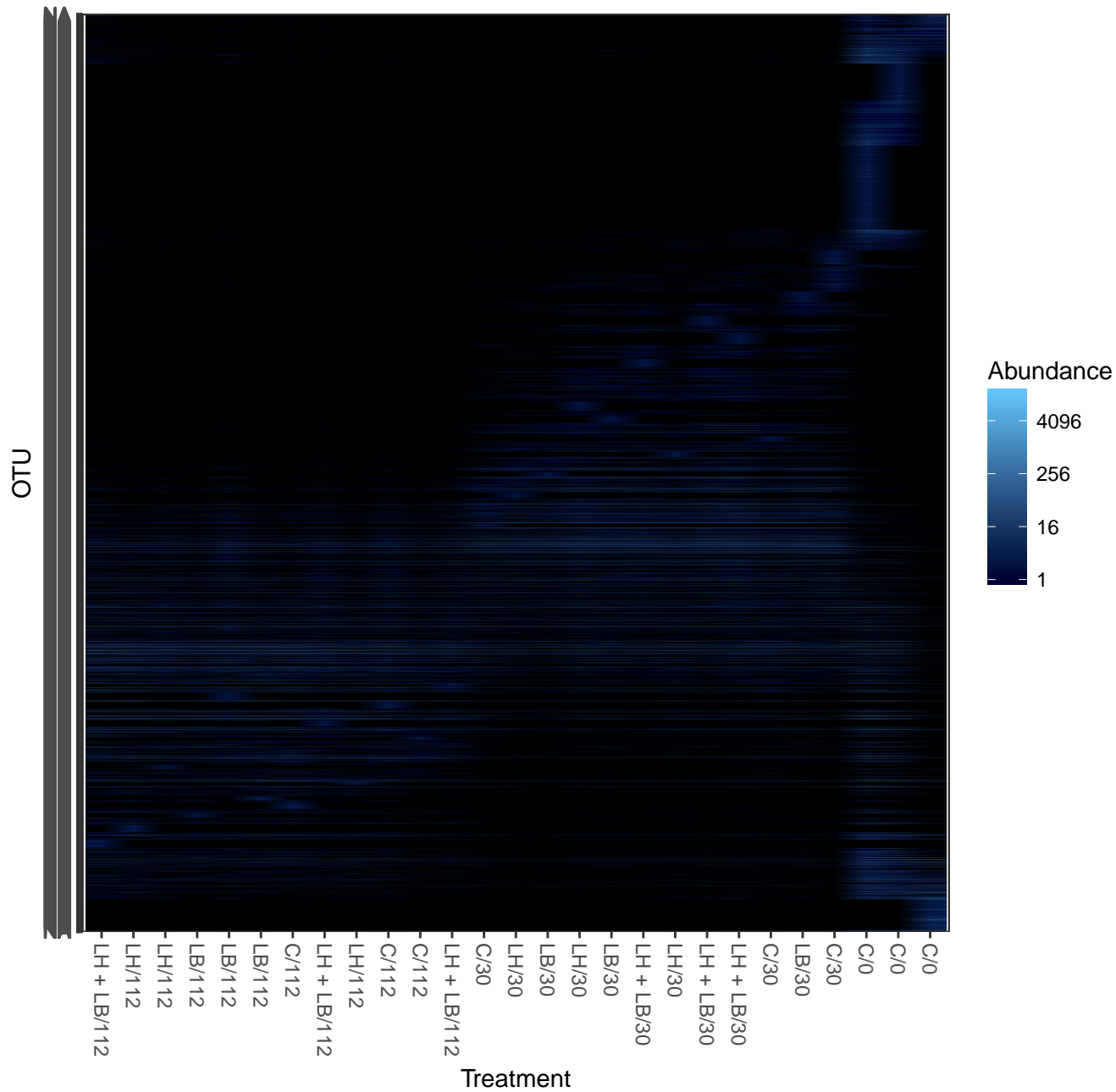
Treatment

15_Hil_CS Bacteria

Axis_2



15_Hil_CS, Bacteria



15_Hil_CS Bacteria: Differential abundance ratio, C/0 vs. LH/30 @ 0.05

