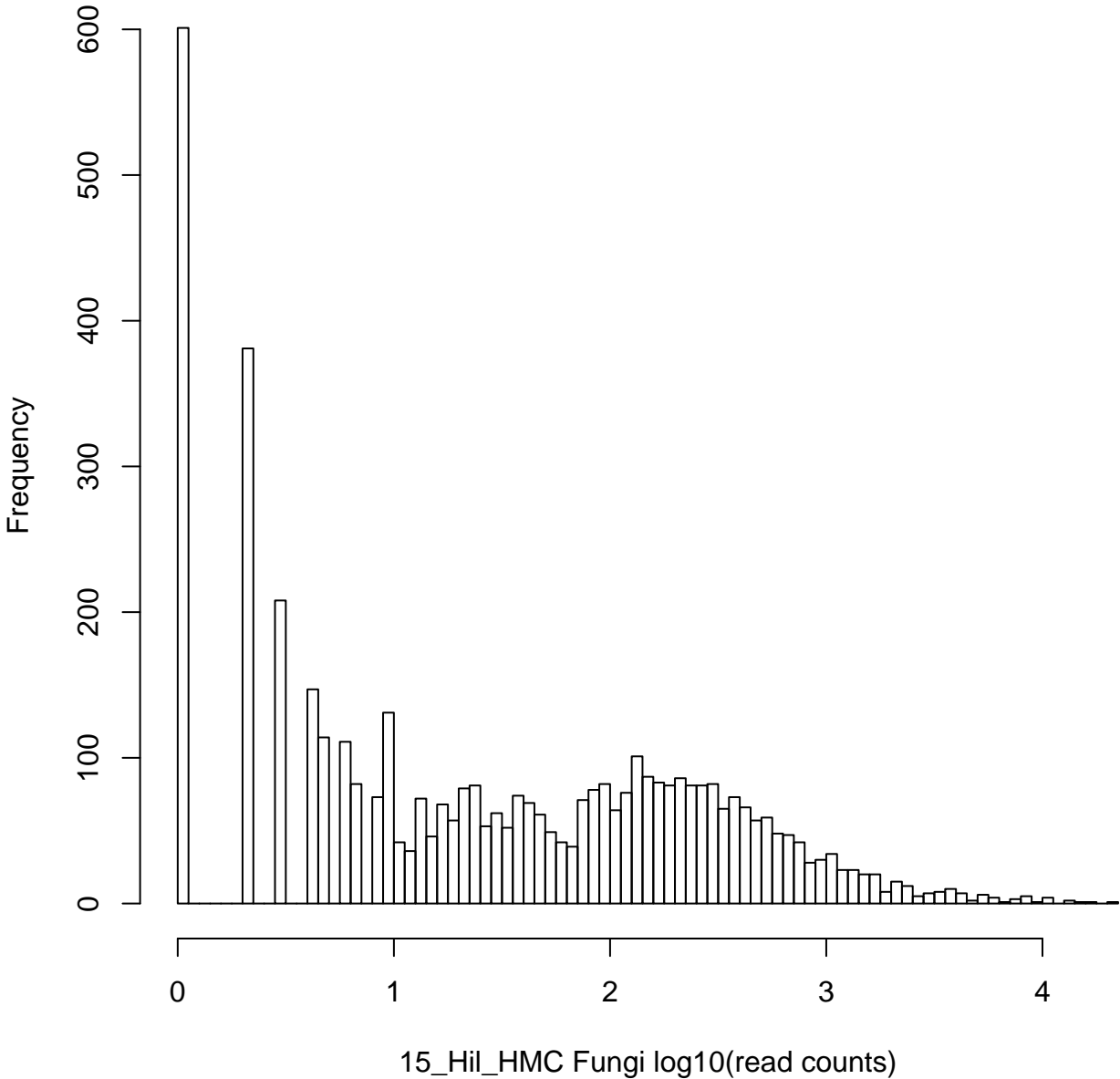
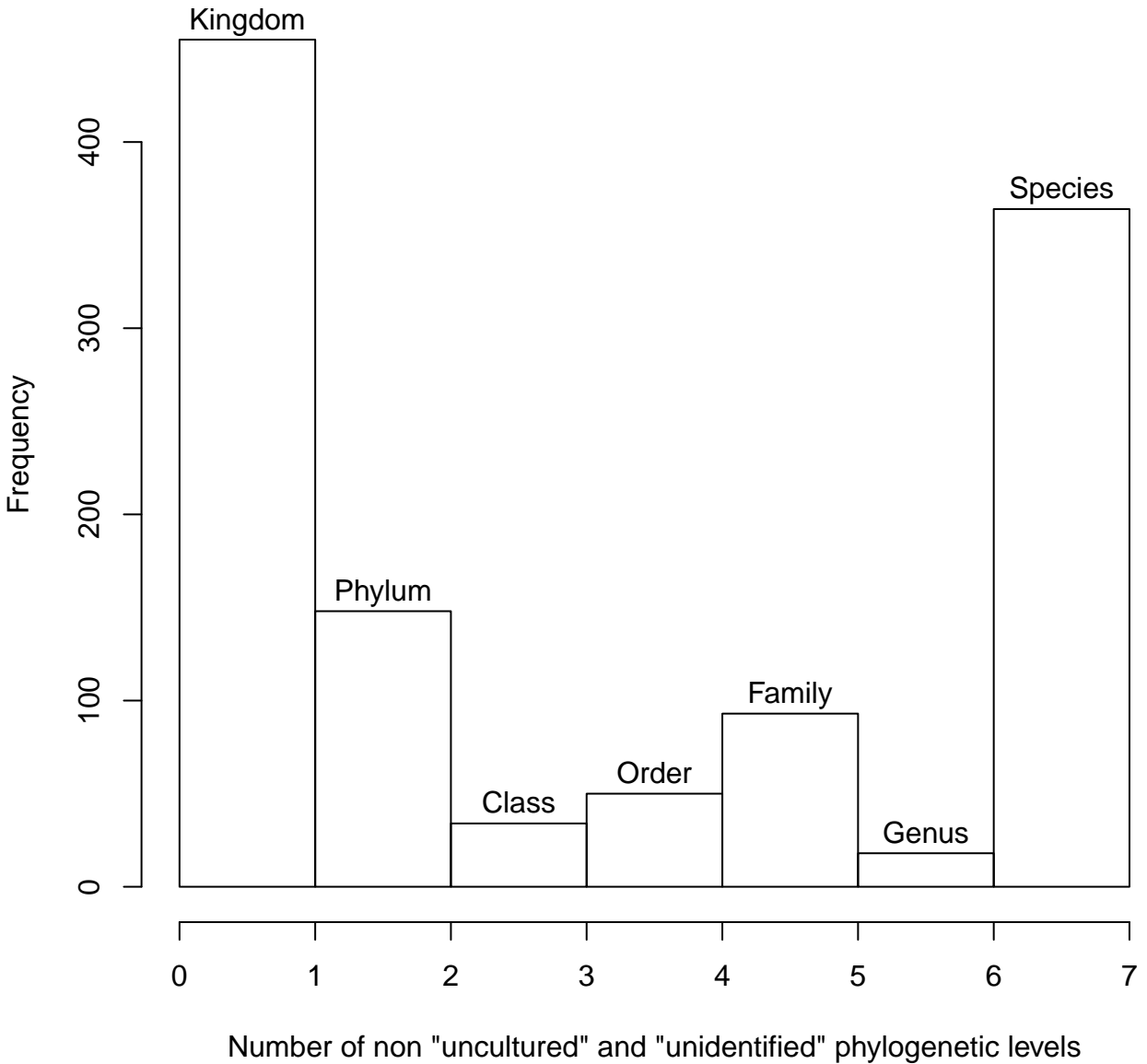


**15\_Hil\_HMC Fungi**  
**Histogram of reads per sample/OTU**

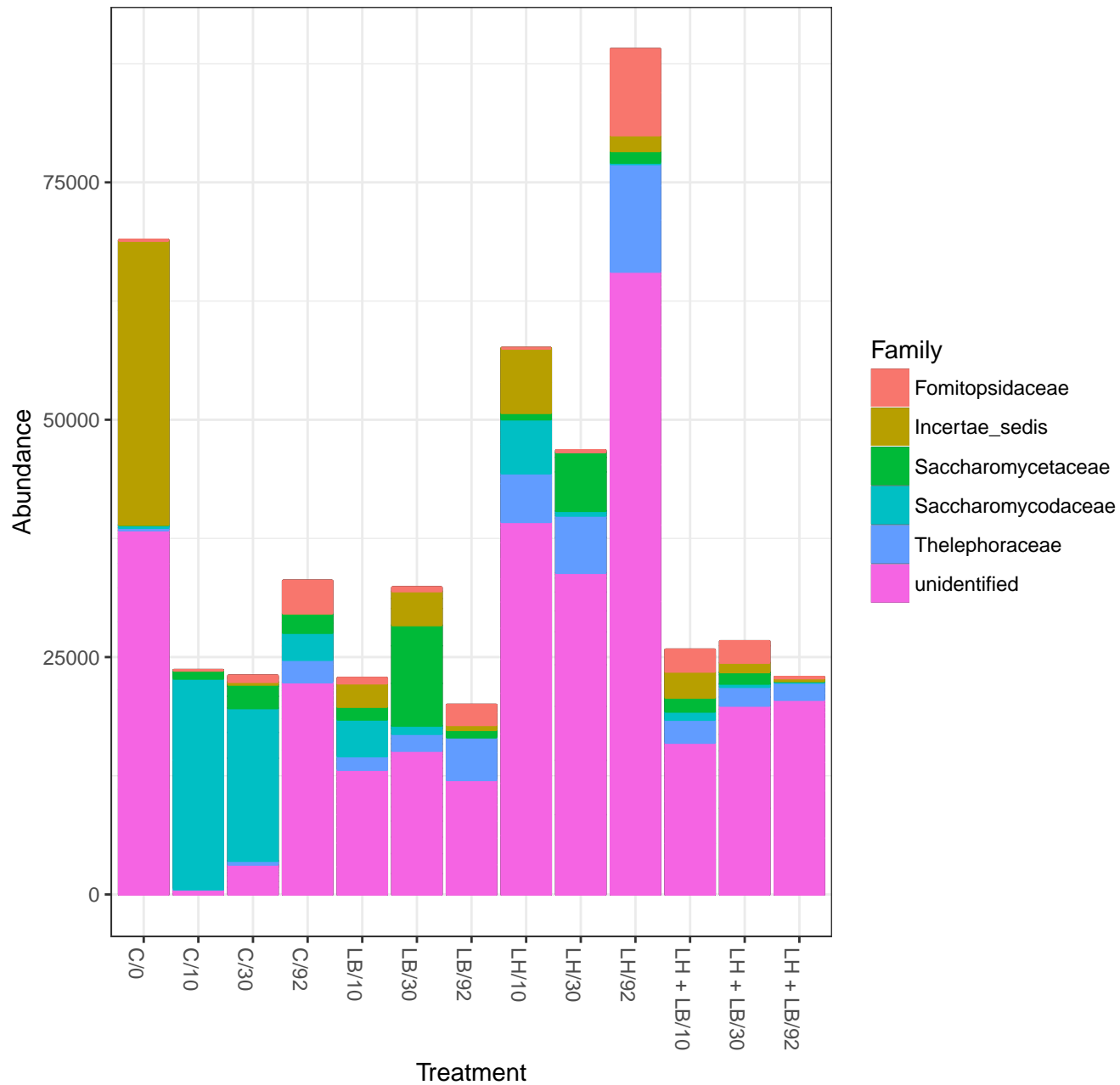


# 15\_Hil\_HMC Fungi

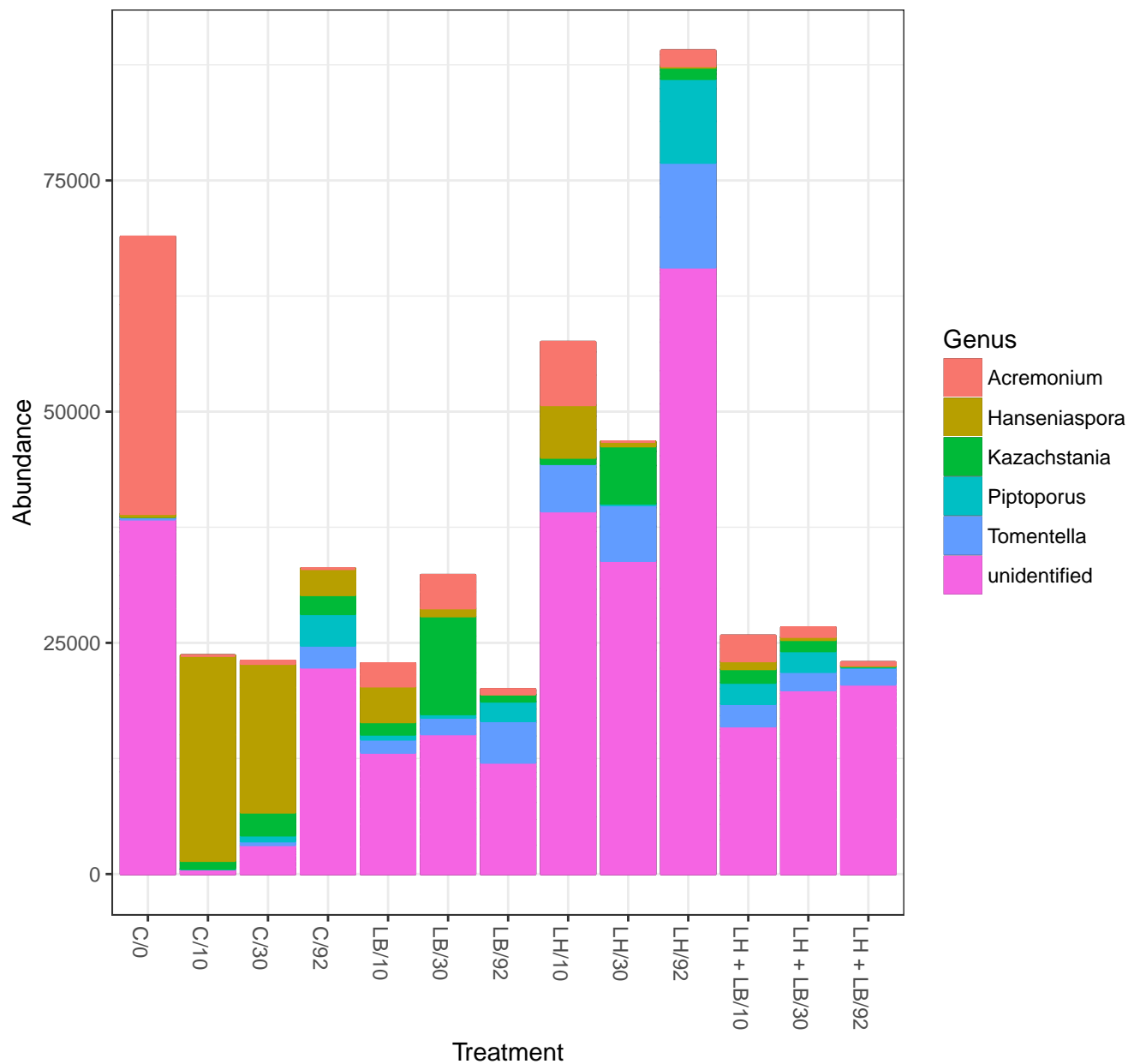
## Histogram of INFERRED most specific rank



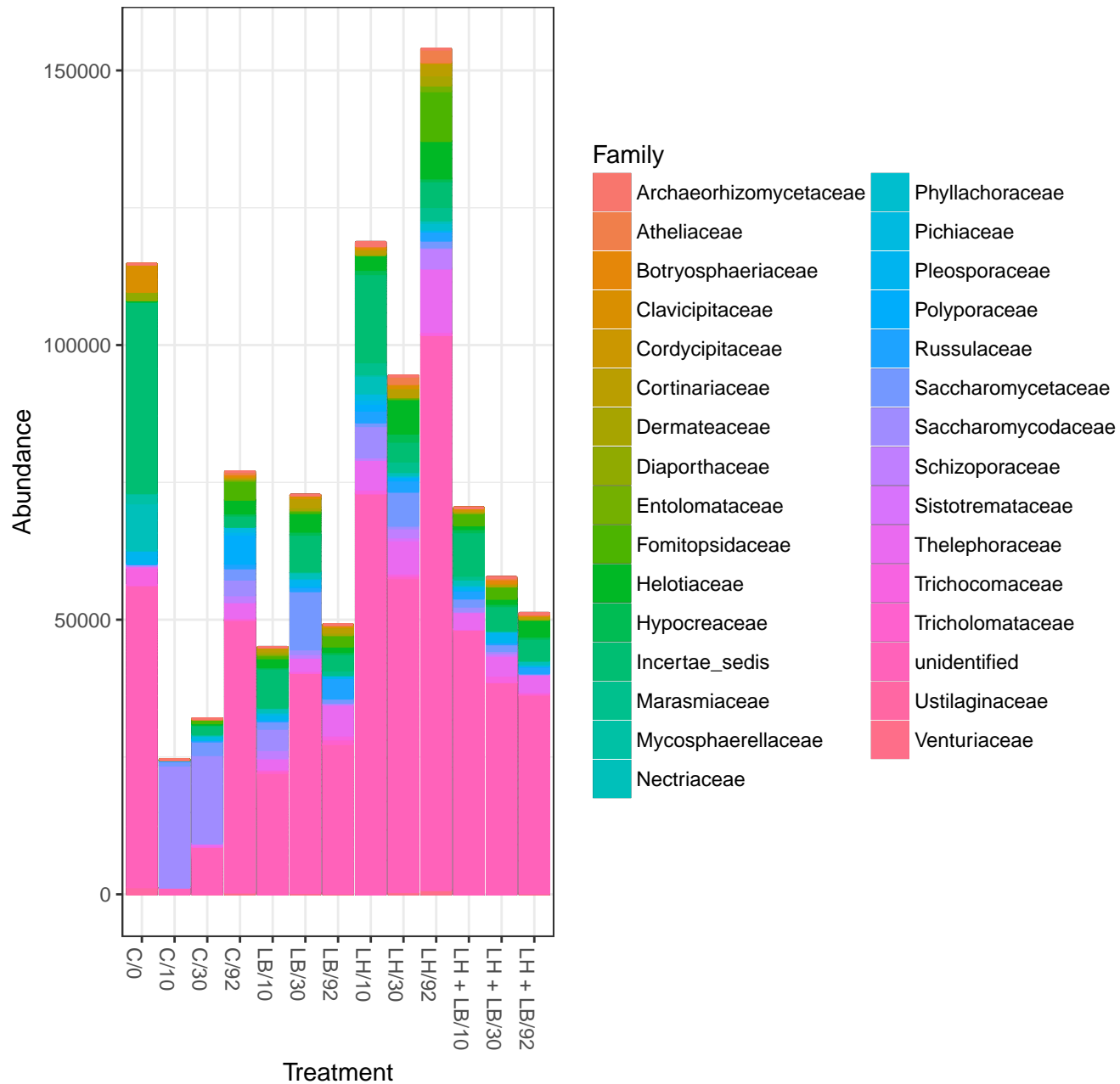
15\_Hil\_HMC Fungi, top 10 OTUs



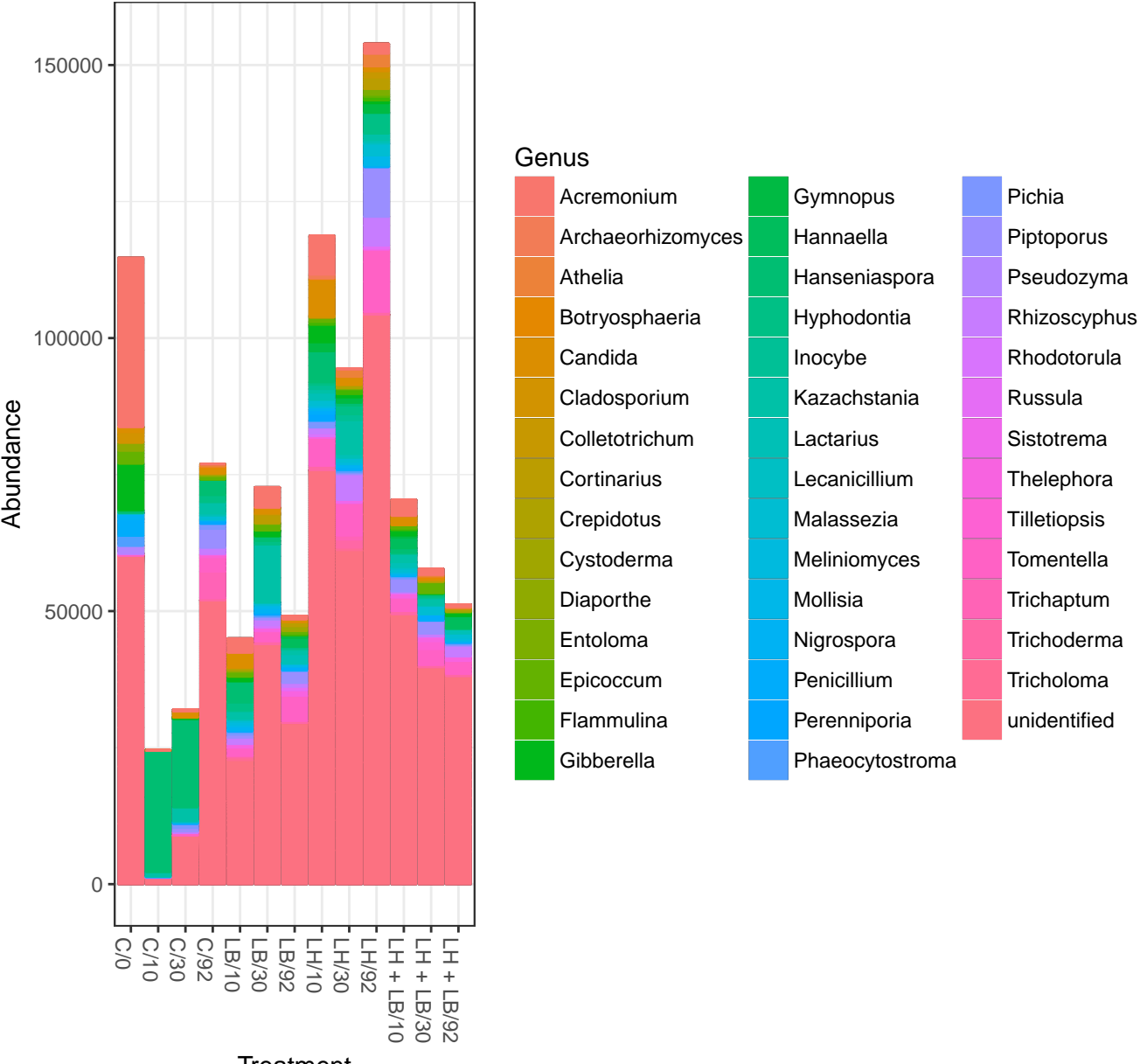
# 15\_Hil\_HMC Fungi, OTUs with at least 1/100 of max reads/sample/OTU in at least one treatment group



# 15\_Hil\_HMC Fungi, top 10 OTUs



15\_Hil\_HMC Fungi, OTUs with at least 1/100 of max reads/sample/OTU  
in at least one treatment group



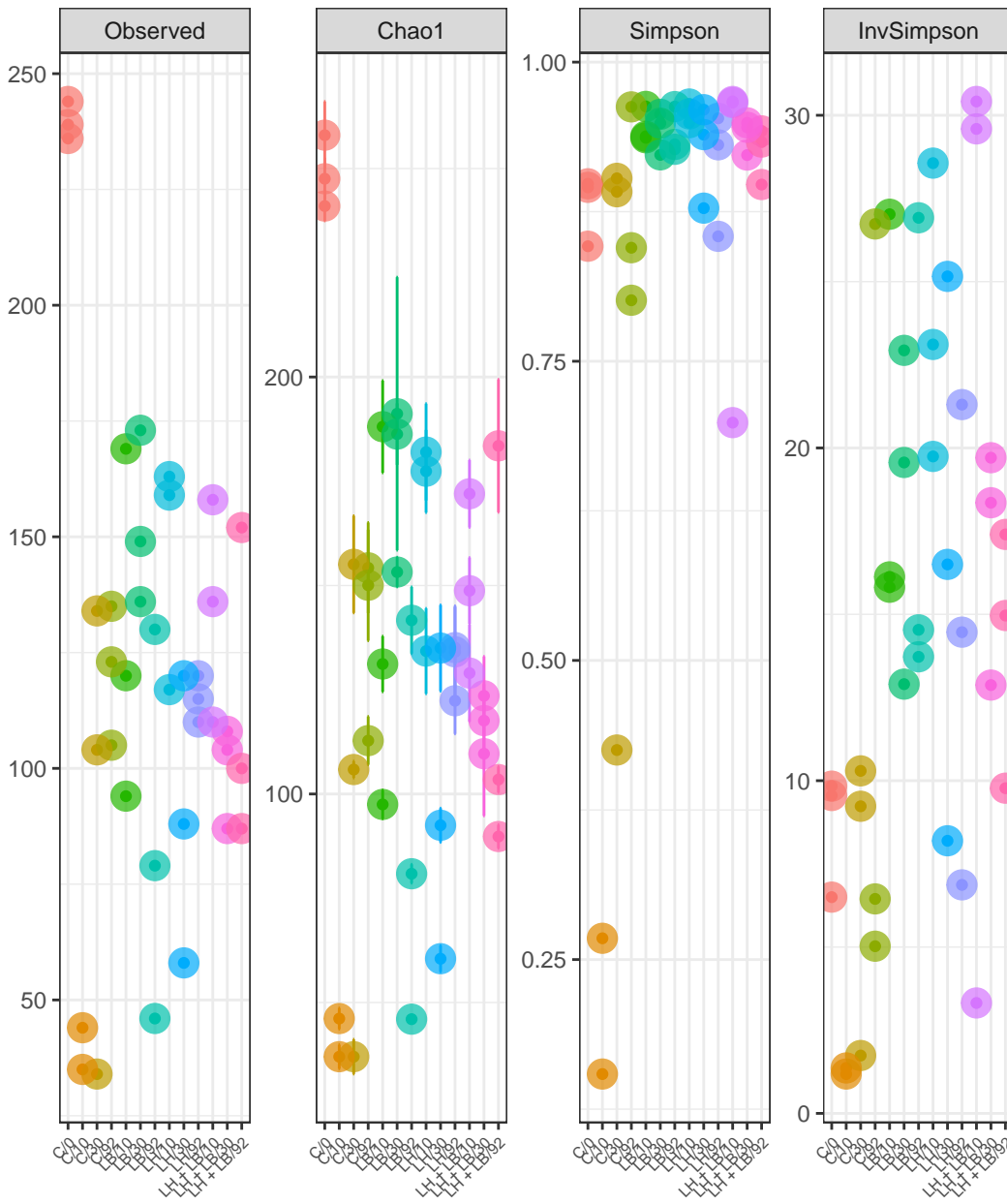
# 15\_Hil\_HMC Fungi

Alpha Diversity Measure

Treatment

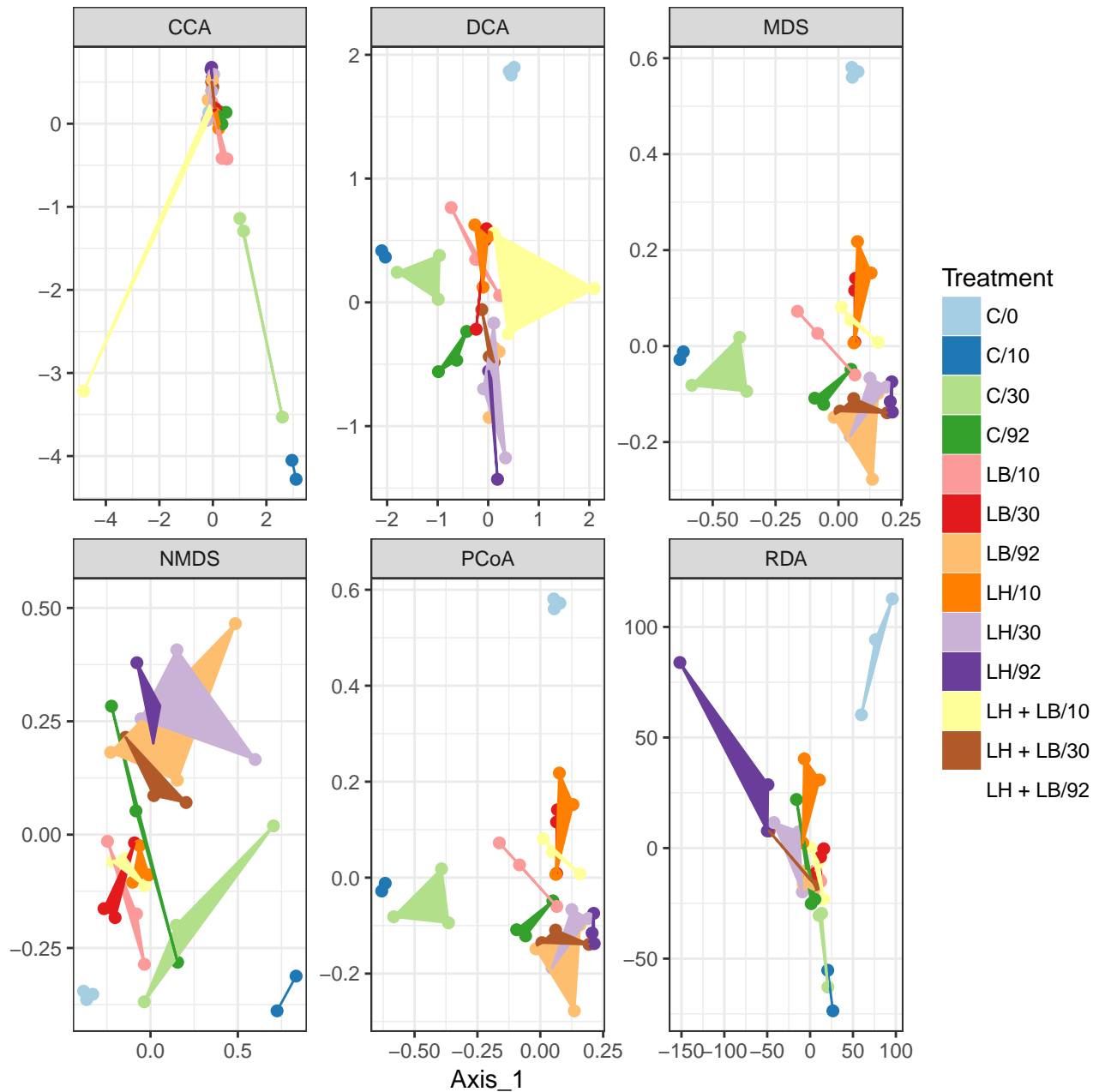
Treatment

- C/0
- C/10
- C/30
- C/92
- LB/10
- LB/30
- LB/92
- LH/10
- LH/30
- LH/92
- LH + LB/10
- LH + LB/30
- LH + LB/92



# 15\_Hil\_HMC Fungi

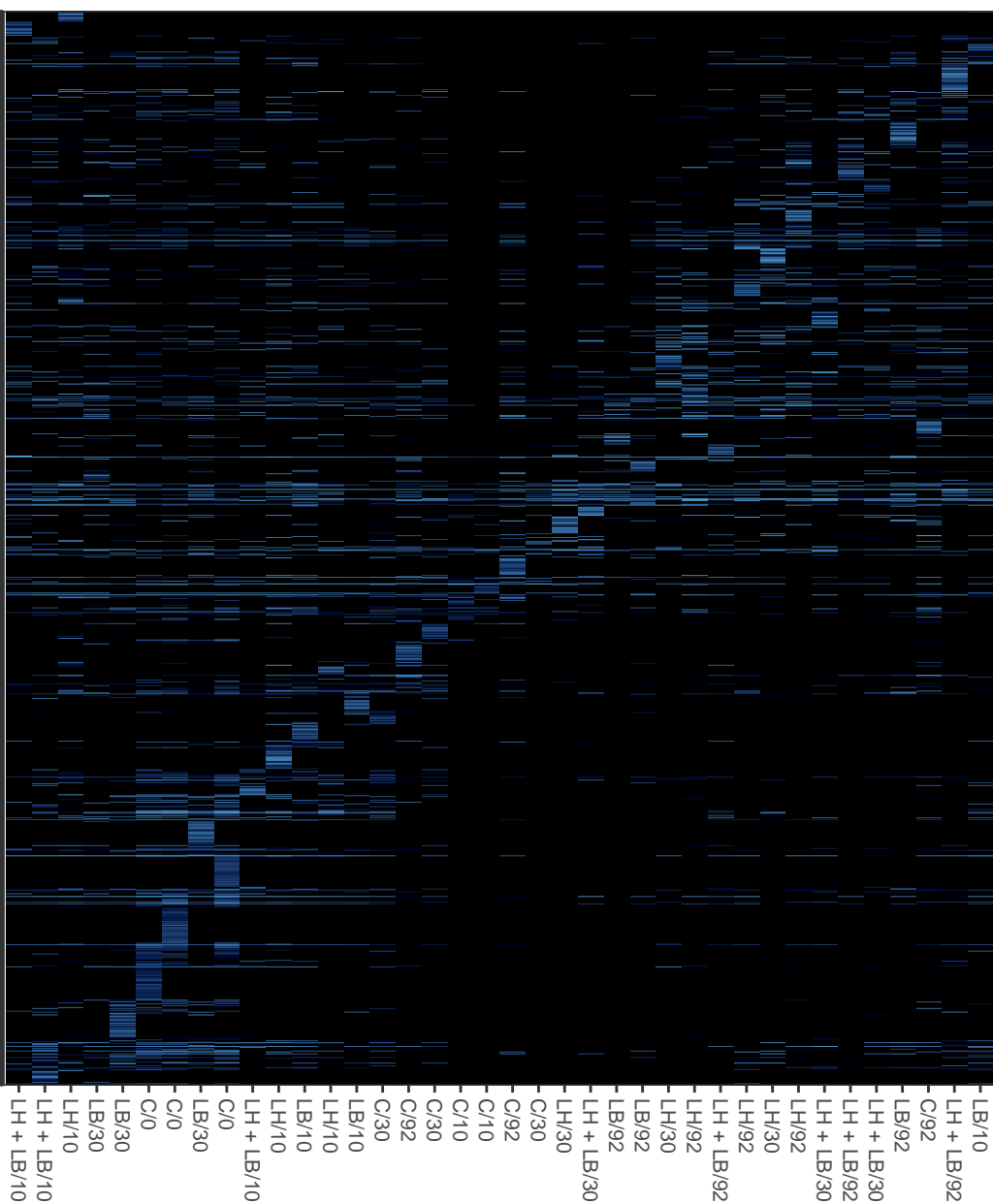
Axis\_2





# 15\_Hil\_HMC, Fungi

OTU



Abundance

4096

256

16

1

Treatment

# 15\_Hil\_HMC Fungi: Differential abundance ratio, C/0 vs. LH/30 @ 0.05

