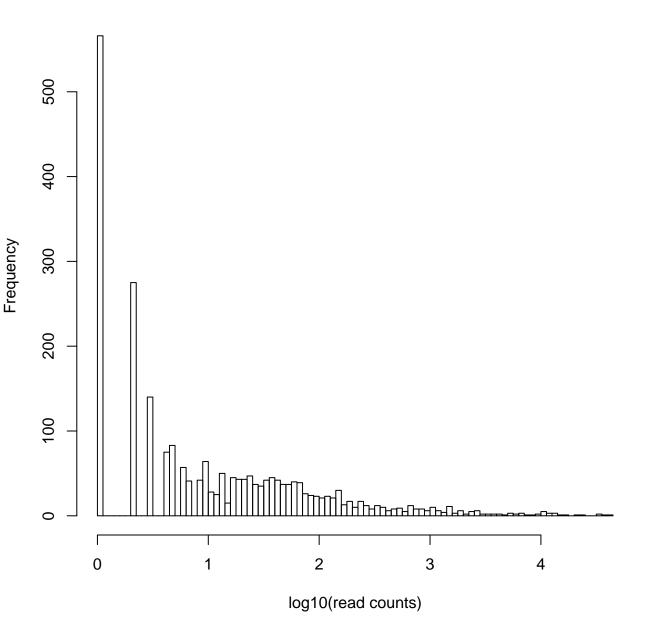
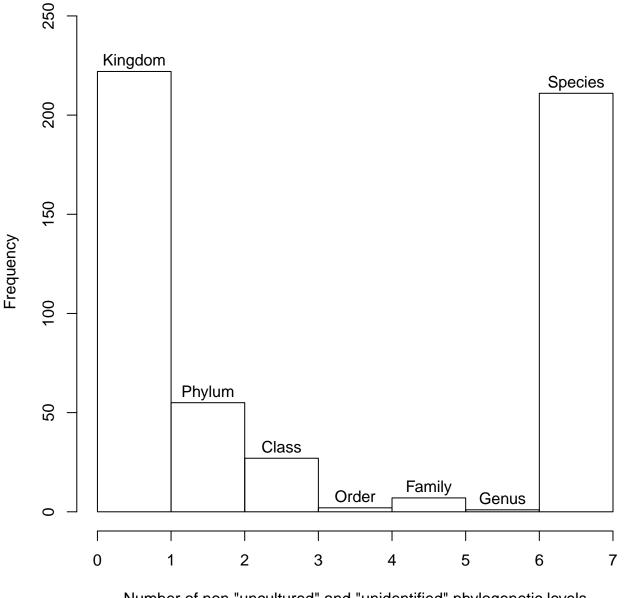
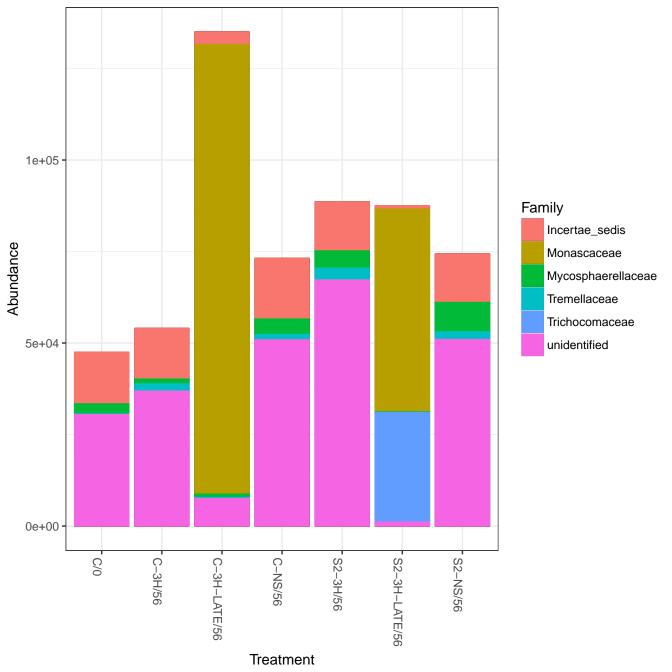
Histogram of reads per sample/OTU

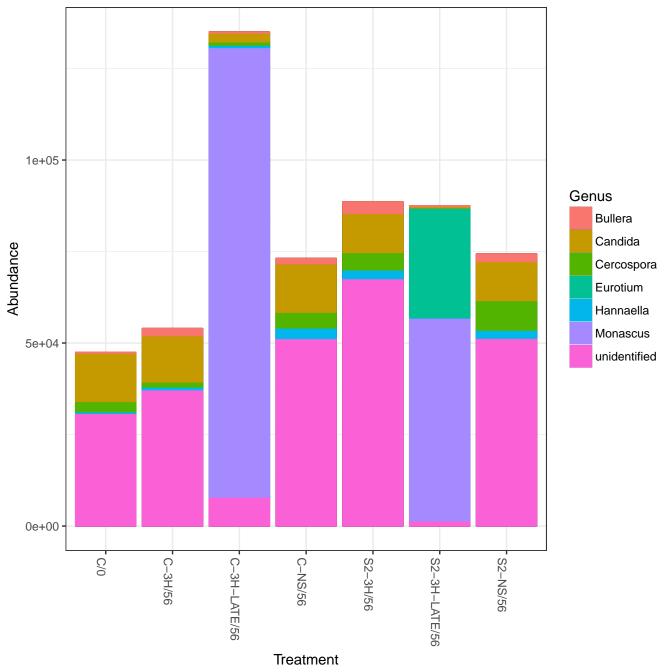


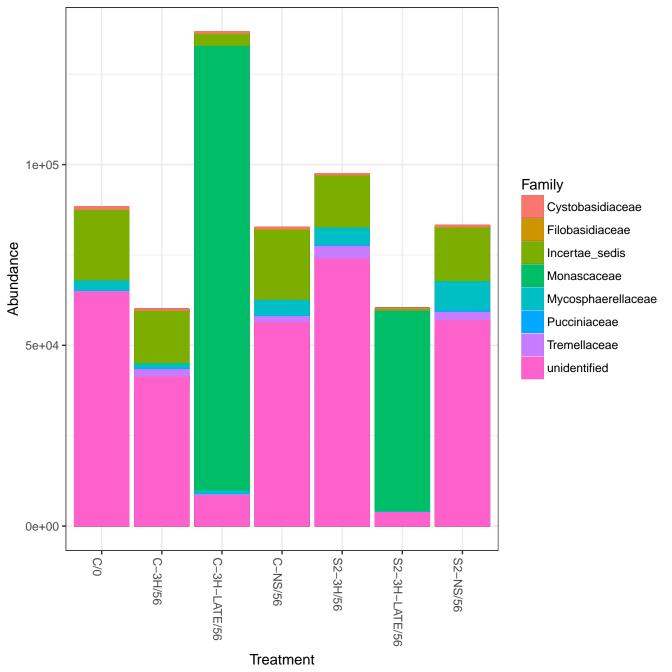
Histogram of INFERRED most specific rank

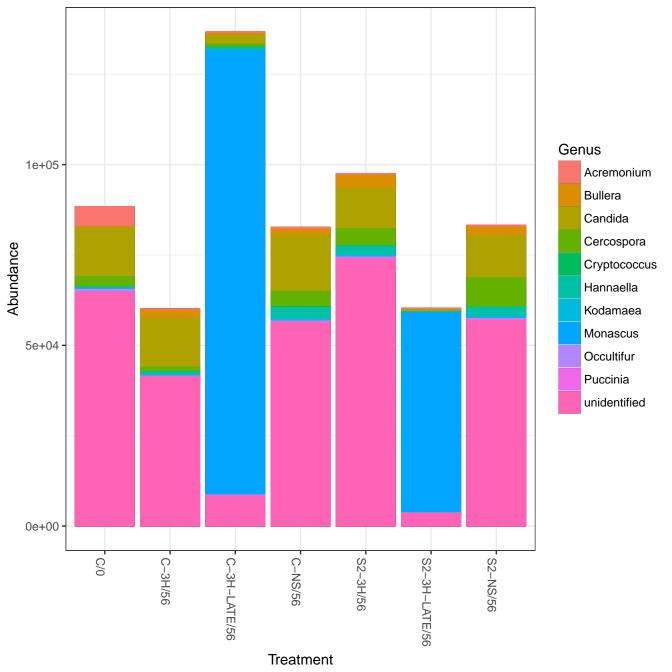


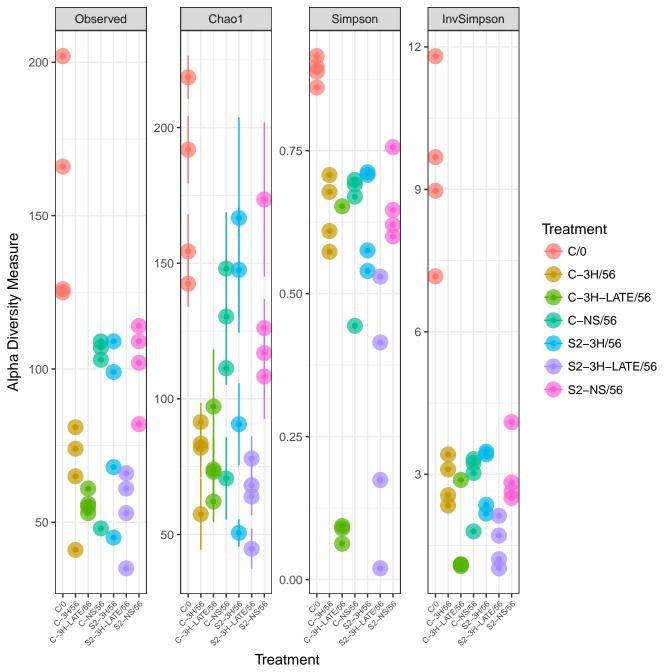
Number of non "uncultured" and "unidentified" phylogenetic levels

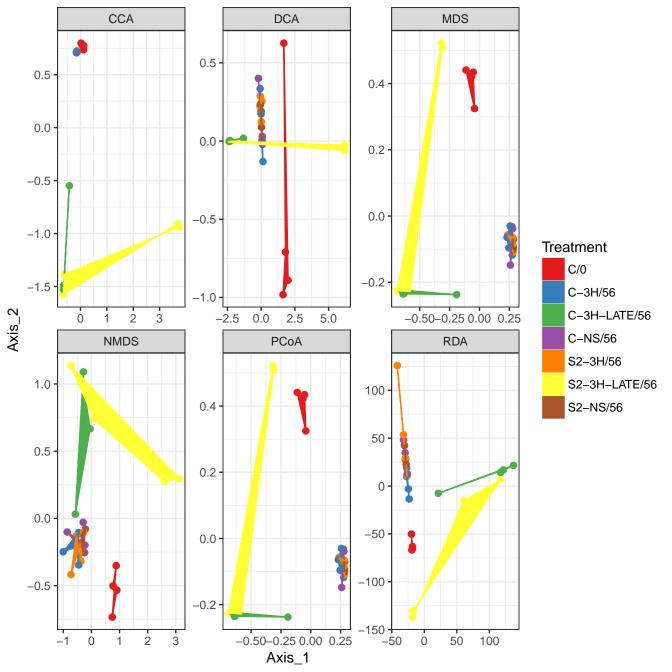


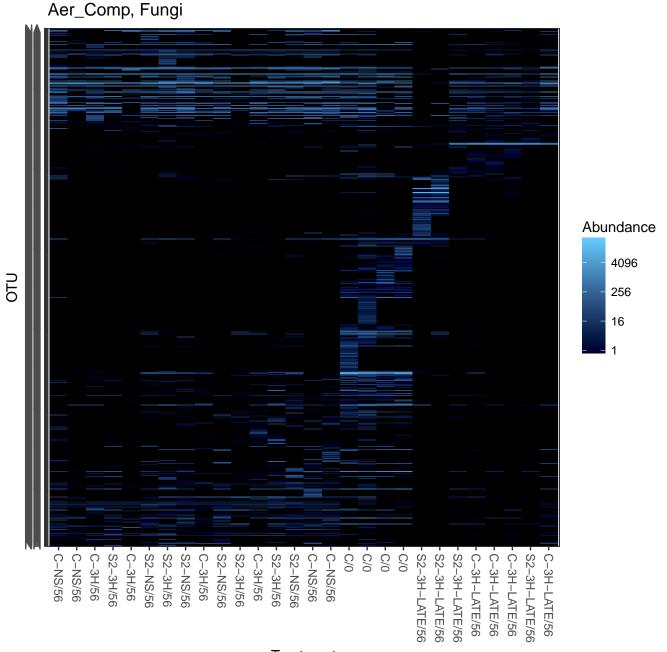












Treatment

