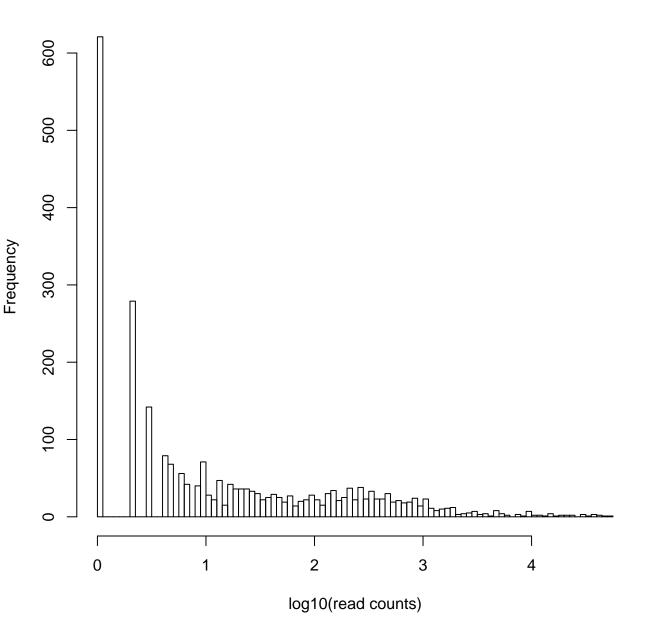
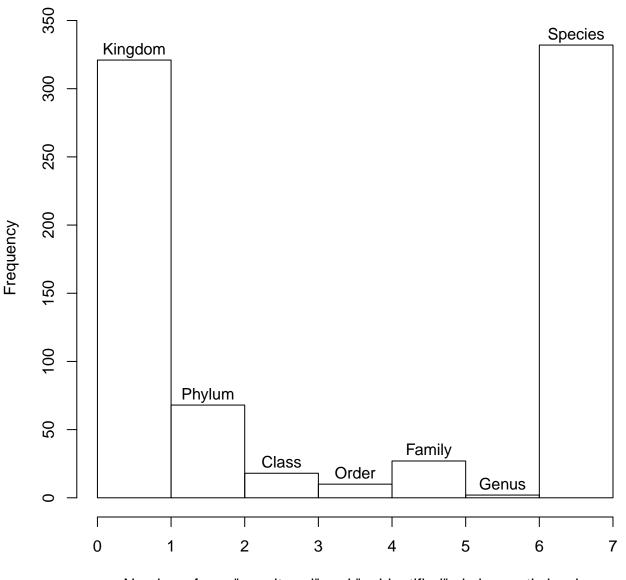
## Histogram of reads per sample/OTU



## Histogram of INFERRED most specific rank



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

