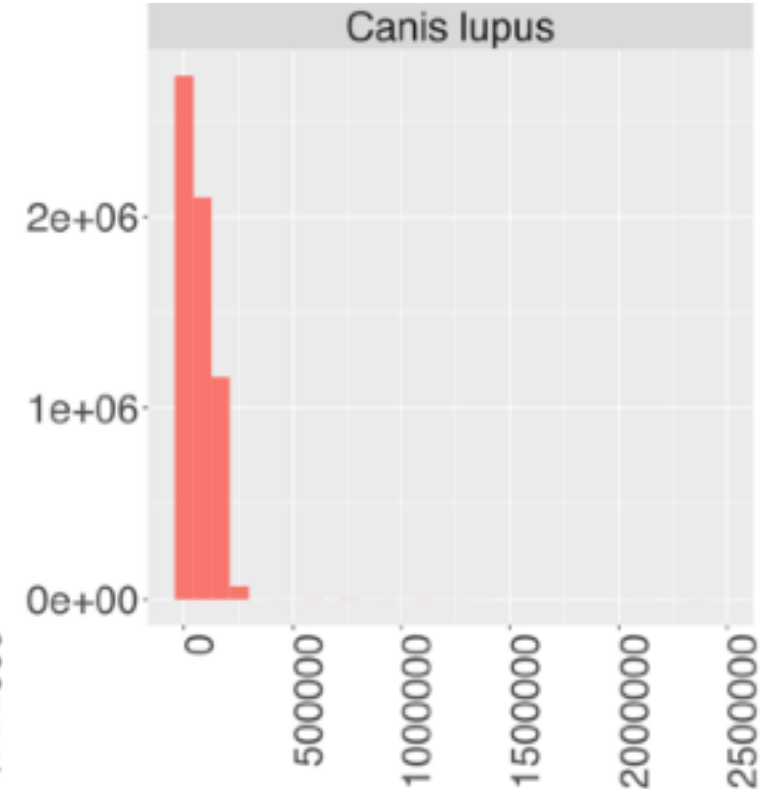
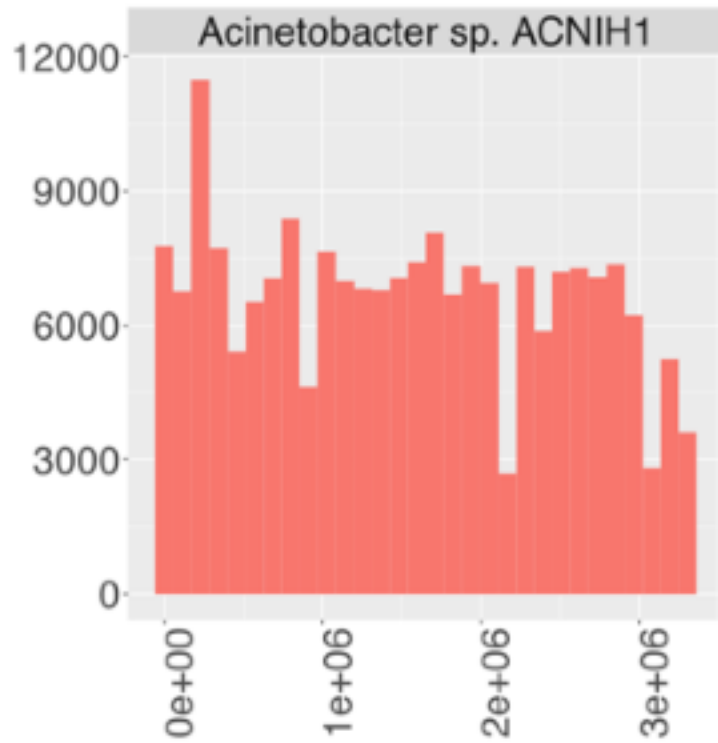


Evaluating results



Genome Position

- *Acinetobacter* sp. ACNIH1
- *Cutibacterium granulosum*
- *Streptococcus pneumoniae*
- *Neisseria meningitidis*

- *Malassezia restricta*
- *Helianthus annuus*

- *Canis lupus* + 26 other eukaryotes
- *Staphylococcus epidermidis*

Adding genomes to database

- *Cutibacterium granulorum* +4
- Basidiomycota (*Malassezia*) +18
- Actinobacteria (*Corynebacterium*) +22
- Firmicutes (*Staphylococcus*) + 23

