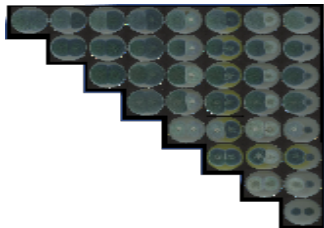


# Experiments

Competition experiments  
& trait measurements



# Bioinformatics

Sequencing and assembly



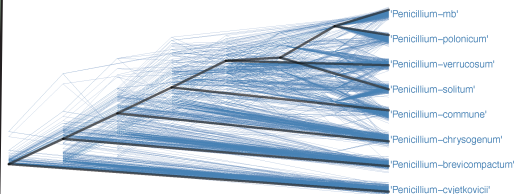
MSA and trees w/MAFFT and FastTree

```
AGTCTTACAAATATC
AGTCTTACAATGATC
ACTCTTACAAGGATC
ACTCTTAGAACGATC
```



Assess phylogenetic signal

Discordance between gene & species trees



PD:RCC for gene and species trees

