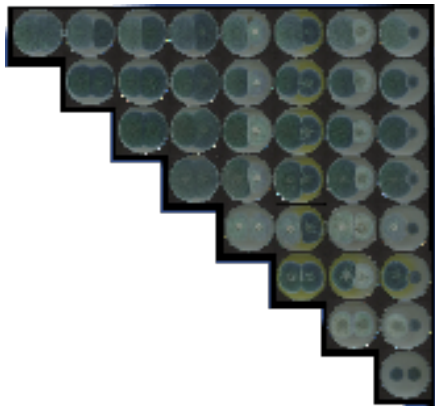


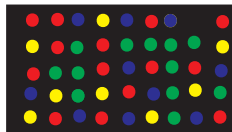
Experiments

Trait measurements



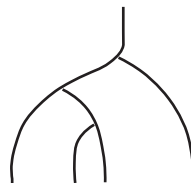
Bioinformatics

Sequencing and assembly



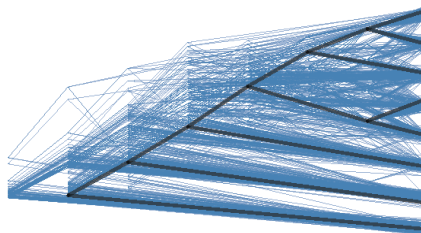
MSA and trees w/MAFFT and FastTree

AGTCTTACAAATATC
AGTCTTACAATGATC
ACTCTTACAAGGATC
ACTCTTAGAACGATC



Assess phylogenetic signal

Discordance between gene & species trees



P. SP. mb
P. polonicum
P. verrucosum
P. solitum
P. bioforme
P. chrysogenum
P. brevicompactum
P. cvjetkovicii

Trait:PD regressions for gene & species trees

