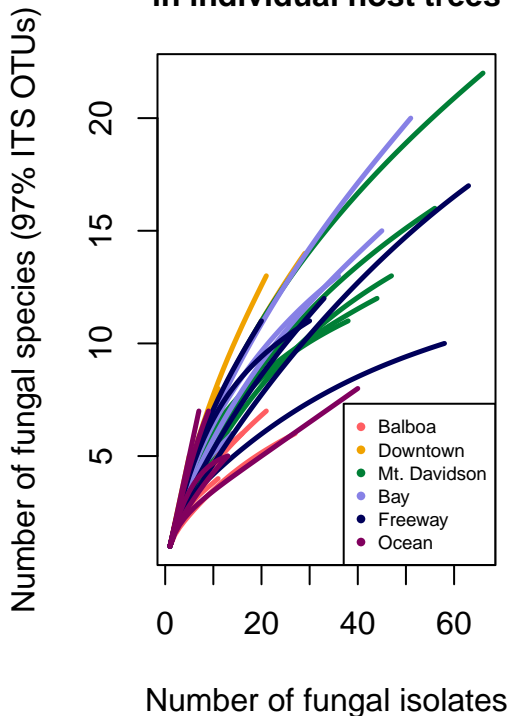


A

**Fungal community richness
in individual host trees**

B

**Fungal community richness
for each site**