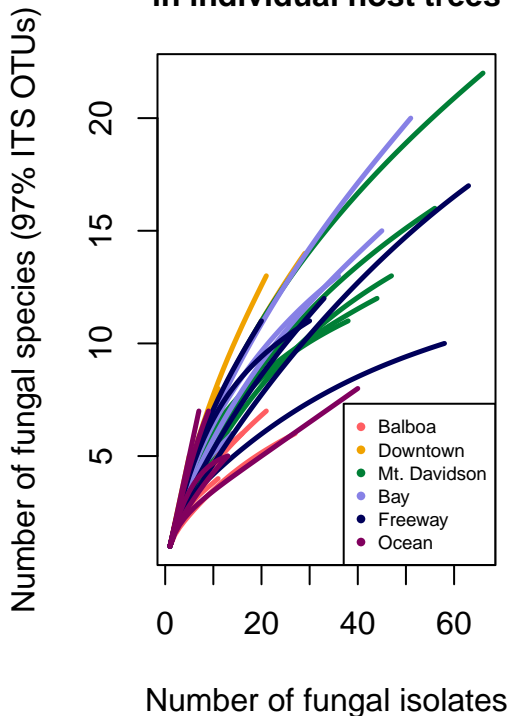


A

**Fungal community richness  
in individual host trees**

B

**Fungal community richness  
for each site**