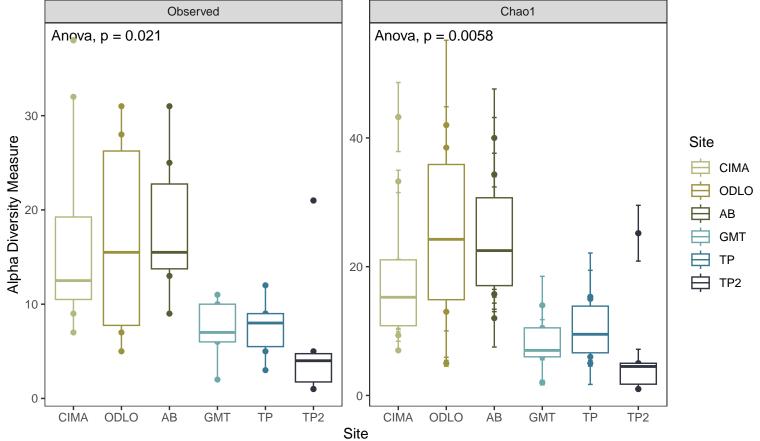
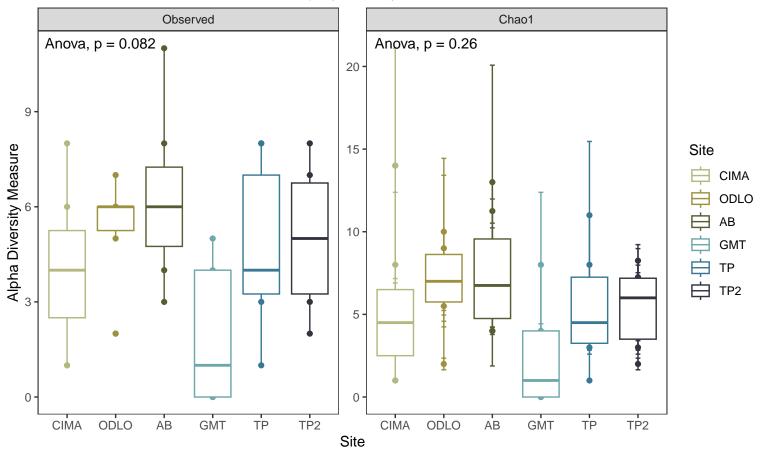
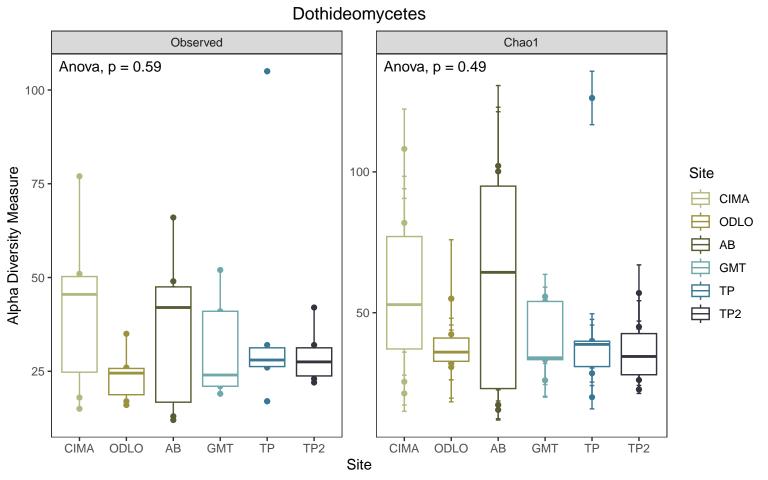
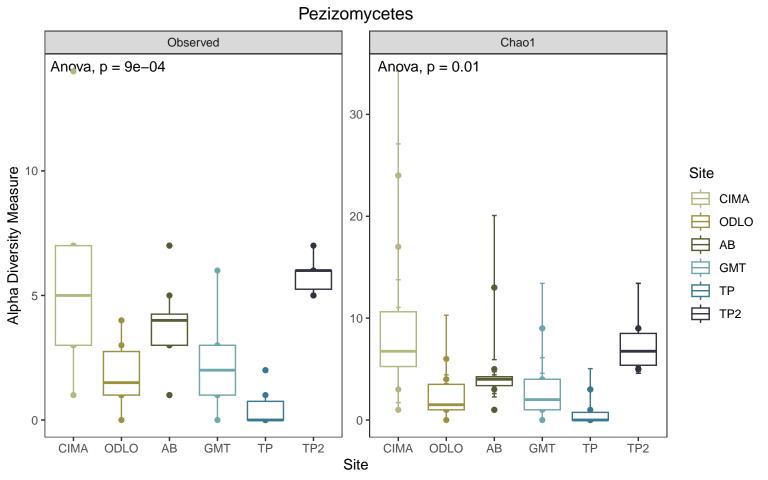
Microbotryomycetes

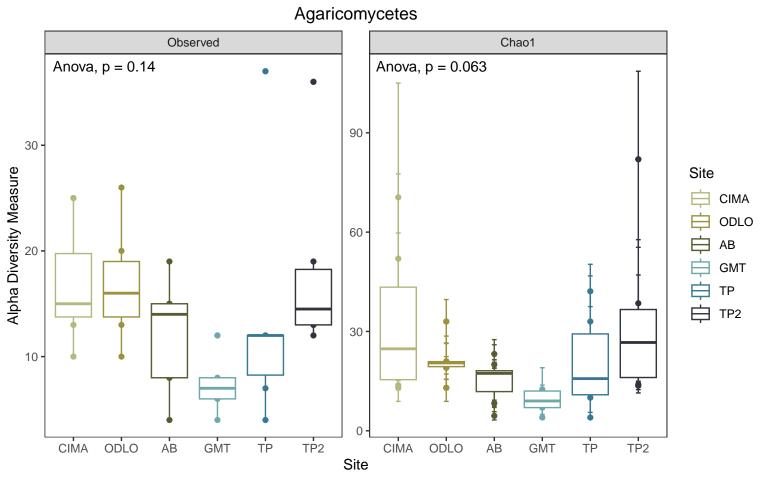


Rhizophlyctidomycetes



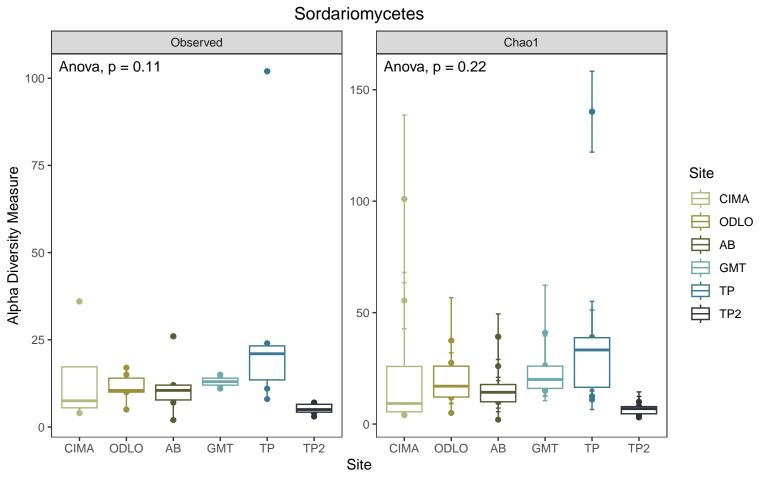




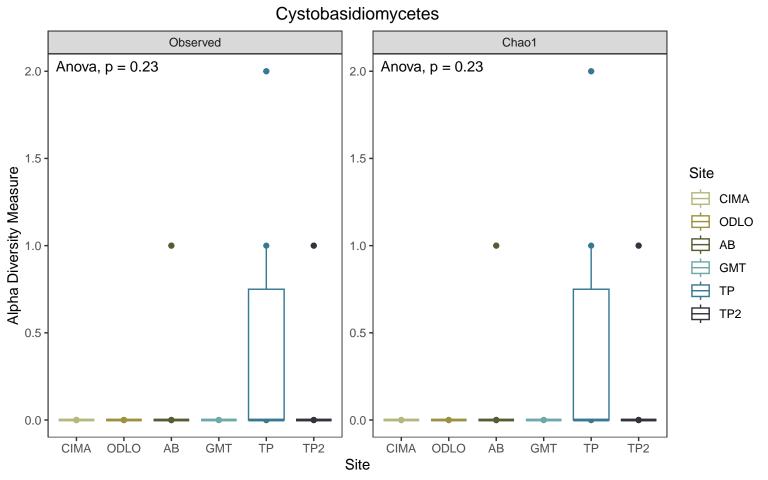


Tremellomycetes Observed Chao1 40 -Anova, p = 0.39Anova, p = 0.5315 -30 -Alpha Diversity Measure Site CIMA ODLO AΒ 20 -**GMT** TP TP2 10 -5 -ODLO ΑB GMT ΤP TP2 ODLO AB ΤP CIMA GMT TP2 CIMA Site

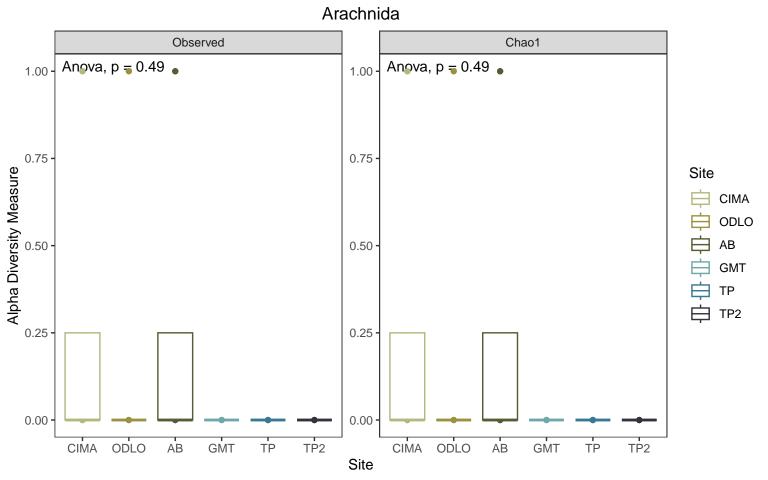
Leotiomycetes Observed Chao1 Anova, p = 0.0011Anova, p = 0.0003910 -Alpha Diversity Measure Site CIMA ODLO AΒ **GMT** 5 TP TP2 0 -0 ODLO ΑB GMT TP2 ODLO ΑB Τ̈́P TP2 GMT Τ̈́P CIMA CIMA



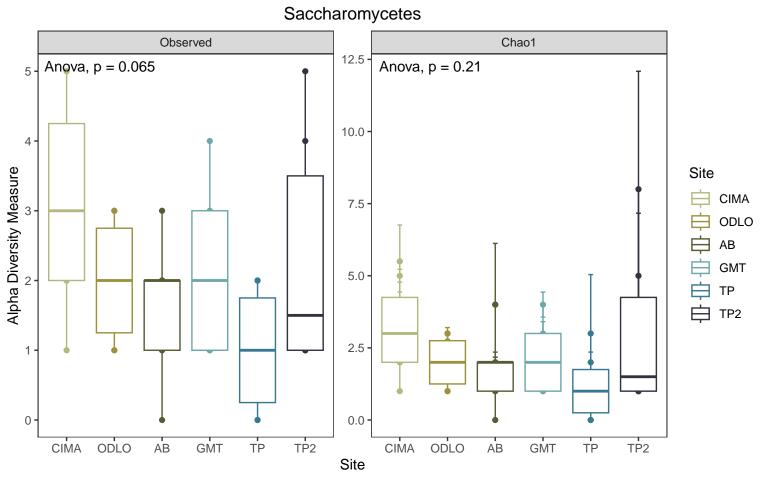
Eurotiomycetes Observed Chao1 Anova, p = 0.00027Anova, p = 1.9e - 0640 -40 -Alpha Diversity Measure Site CIMA 30 -ODLO AΒ **GMT** 20 -TP TP2 10-10 -ODLO ΑB AB CIMA GMT Τ'n ODLO GMT TP2 CIMA TP TP2 Site



Lecanoromycetes Observed Chao1 Anova, p = 0.043Anova, p = 0.0577.5 9 · Site Alpha Diversity Measure CIMA ODLO 6 -AΒ GMT TP TP2 3 0.0 0 . TP2 ΑB ΤP CIMA ODLO AB ΤP ODLO GMT GMT TP2 CIMA

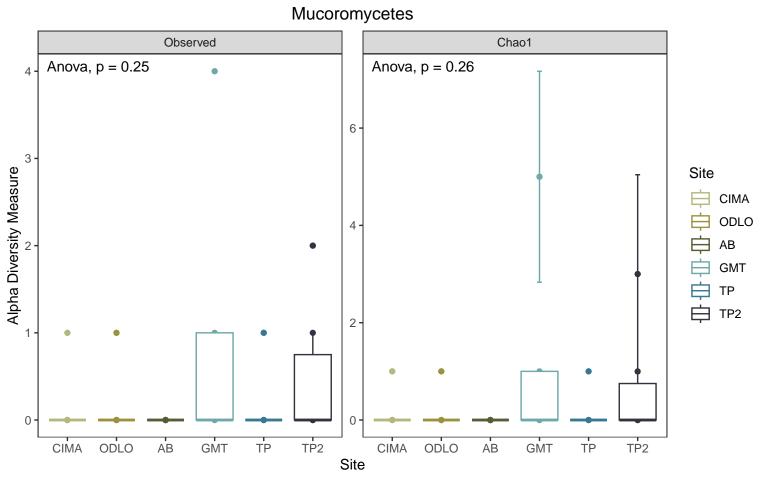


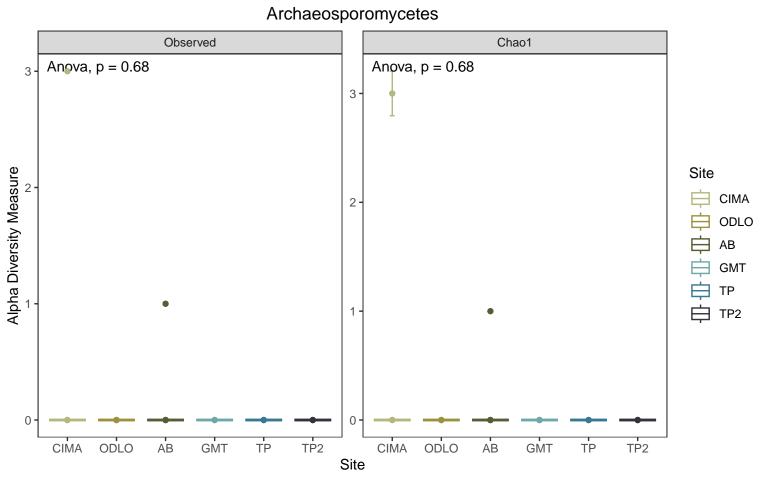
Blastocladiomycetes Observed Chao1 $_{10.0}$ Anova, p = 0.19 Anova, p = 0.167.5 -10 · Alpha Diversity Measure Site CIMA ODLO 5.0 -AΒ **GMT** 5 TP TP2 2.5 -0.0 0 -TP2 ODLO AΒ GMT TP CIMA ODLO AB GMT TP2 CIMA TP

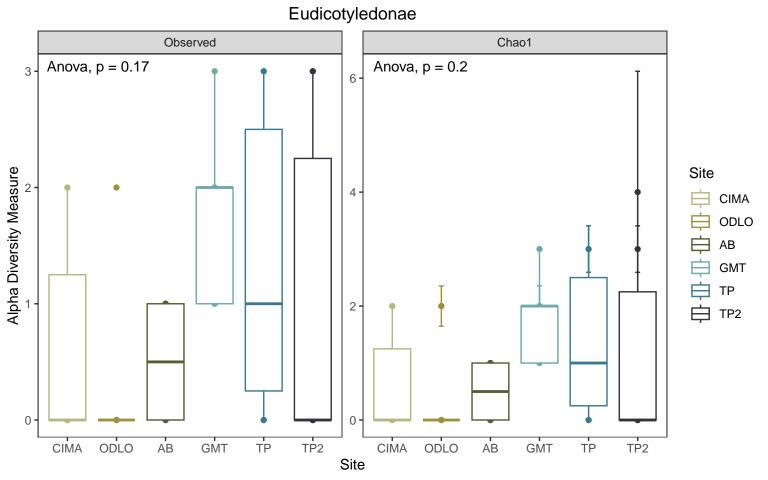


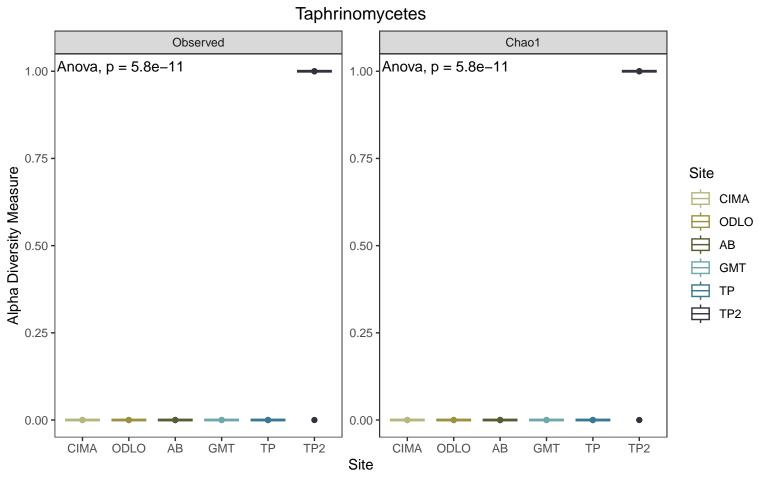
Glomeromycetes Observed Chao1 60 -Anova, p = 0.05Anova, p = 0.019125 **-**100 -Alpha Diversity Measure Site CIMA 75 -ODLO AΒ **GMT** 50 -TP TP2 25 -0 -0 -TP2 ODLO ΑB ΤP CIMA ODLO AB GMT GMT CIMA TP TP2

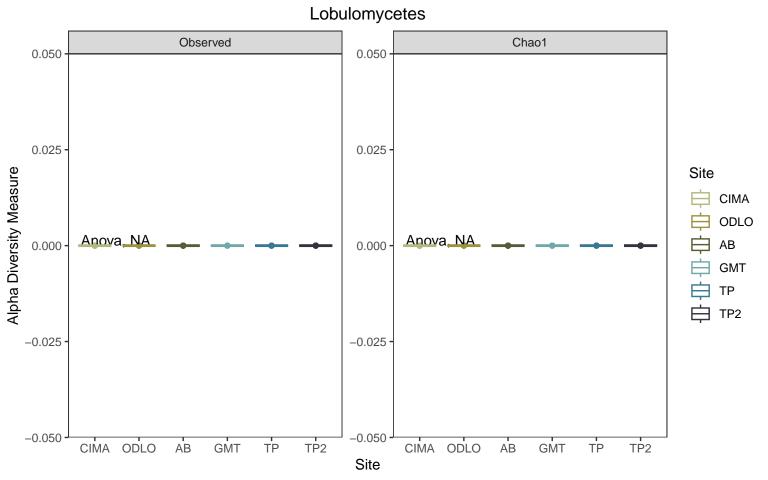
Mortierellomycetes Observed Chao1 Anova, p = 0.00013Anova, p = 0.001610.0 -6 -7.5 -Alpha Diversity Measure Site CIMA ODLO AΒ 5.0 -**GMT** TP TP2 2.5 0 . 0.0 TP2 ΑB ΤP ODLO AB TP ODLO GMT GMT CIMA TP2 CIMA Site

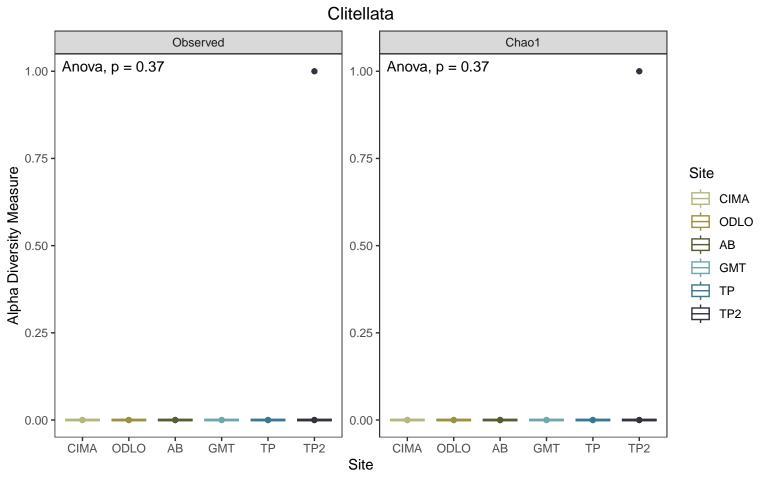


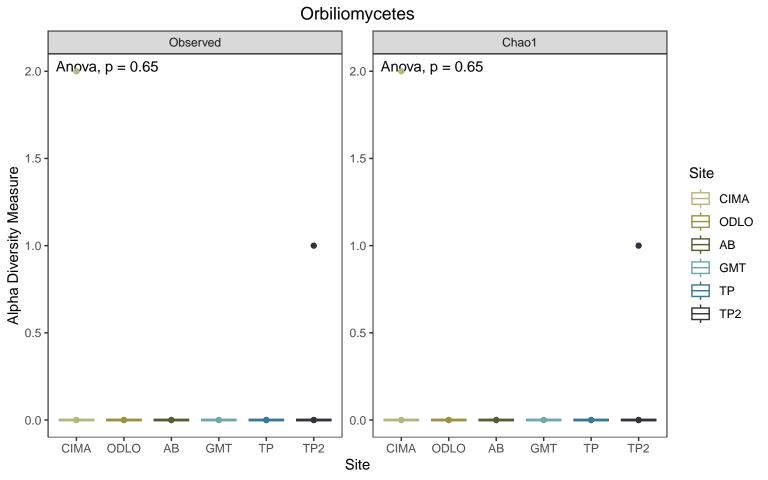


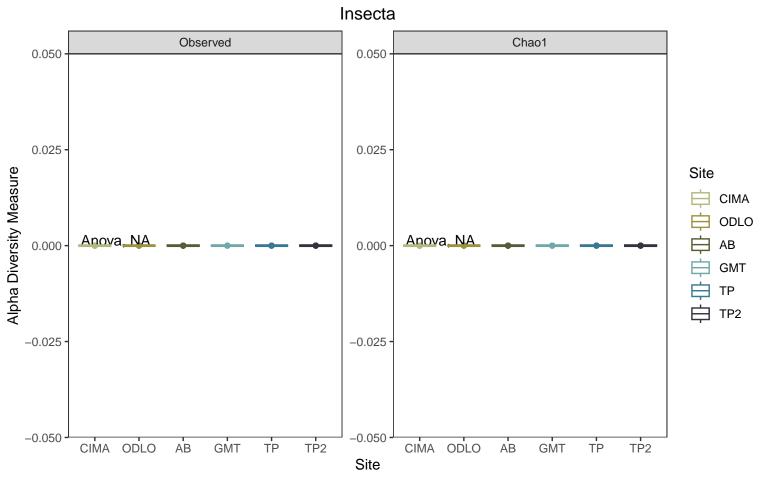




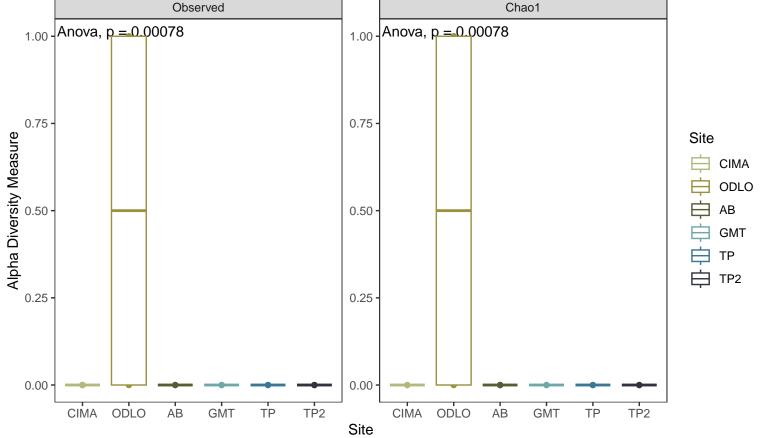


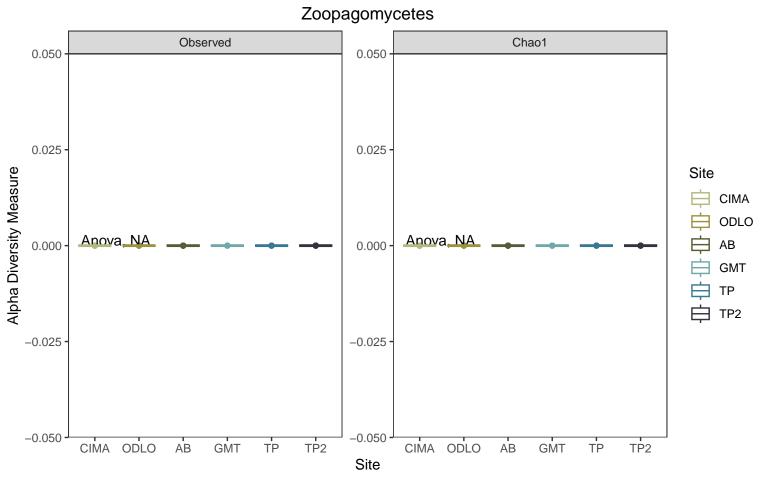






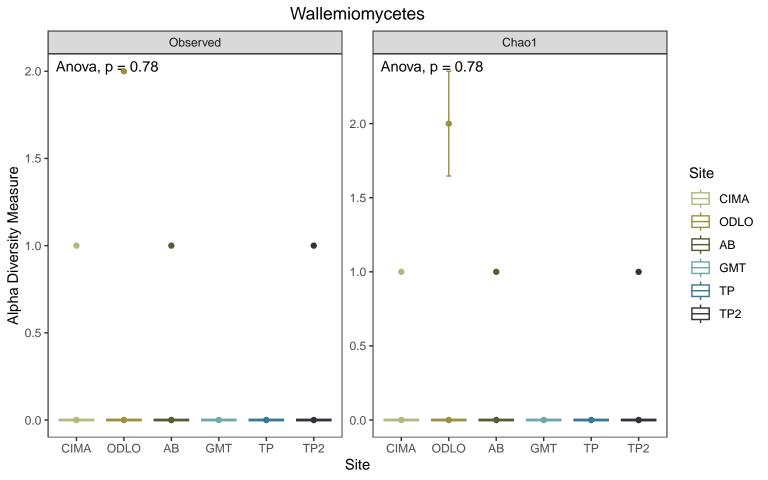
Archaeorhizomycetes





Monocotyledonae Observed Chao1 $_{1.00}$ Anova, p = 0.73 • Anova, p = 0.730.75 -0.75 -Alpha Diversity Measure Site CIMA ODLO 0.50 -AΒ **GMT** TP TP2 0.25 -0.25 -0.00 -0.00 -TP2 GMT ΤP ODLO GMT ODLO ΑB CIMA ΑB ΤP CIMA TP2 Site

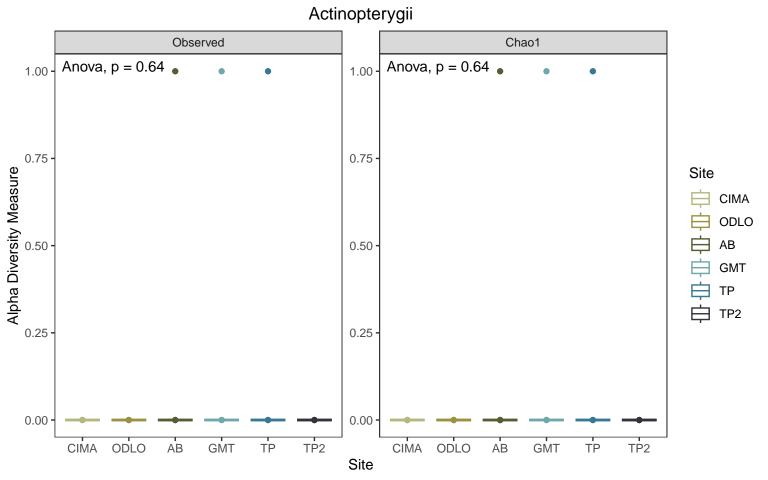
Geminibasidiomycetes Observed Chao1 $_{1.00}$ Anova, p = 0.00078 $_{1.00}$ Anova, p = 0.00078 0.75 -0.75 -Alpha Diversity Measure Site CIMA ODLO 0.50 -AΒ 0.50 -**GMT** TP TP2 0.25 -0.25 -0.00 -0.00 -GMT ΤP TP2 ODLO ODLO GMT CIMA CIMA AB AB TP TP2 Site

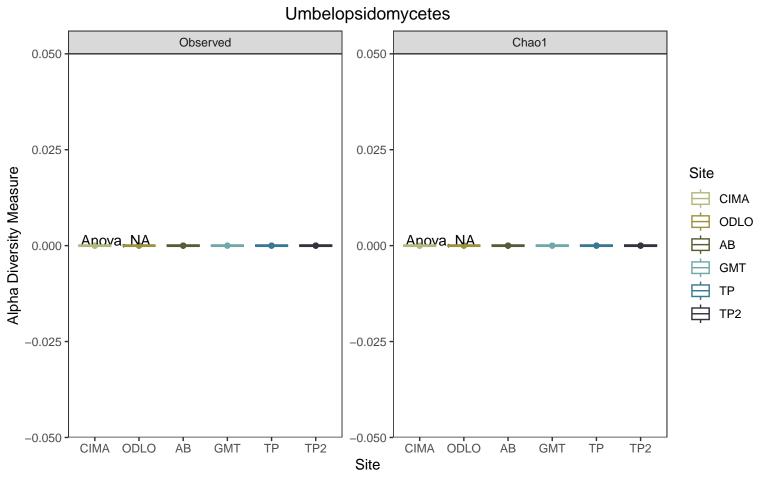


Lichinomycetes Observed Chao1 $_{1.00}$ Anova, p = 0.37 Anova, p = 0.370.75 -0.75 -Alpha Diversity Measure Site CIMA ODLO 0.50 -AΒ **GMT** TP TP2 0.25 -0.25 -0.00 -0.00 -TP2 GMT ΤP ODLO GMT ODLO ΑB CIMA ΑB ΤP CIMA TP2 Site

Agaricostilbomycetes Observed Chao1 $_{1.00}$ Anova, p = 0.17 Anova, p = 0.170.75 -0.75 -Alpha Diversity Measure Site CIMA ODLO 0.50 -AΒ **GMT** TP TP2 0.25 -0.25 -0.00 -0.00 -GMT ΤP TP2 ODLO ΑB ODLO GMT ΑB CIMA CIMA TP TP2 Site

Basidiobolomycetes Chao1 Observed Anova, p = 0.18Anova, p = 0.183 3 Alpha Diversity Measure Site CIMA ODLO AΒ 2 -**GMT** TP TP2 1 0 0 ODLO ΑB TP TP2 CIMA ΑB ΤP GMT ODLO GMT TP2 CIMA Site





Malasseziomycetes Observed Chao1 _{1.00} Anova, p = 0.37 Anova, p = 0.37 0.75 -0.75 -Alpha Diversity Measure Site CIMA ODLO AΒ 0.50 -**GMT** TP TP2 0.25 -0.25 -0.00 -0.00 -

ODLO

CIMA

Site

GMT

TP

TP2

ΑB

TP2

ΤP

GMT

ΑB

ODLO

CIMA