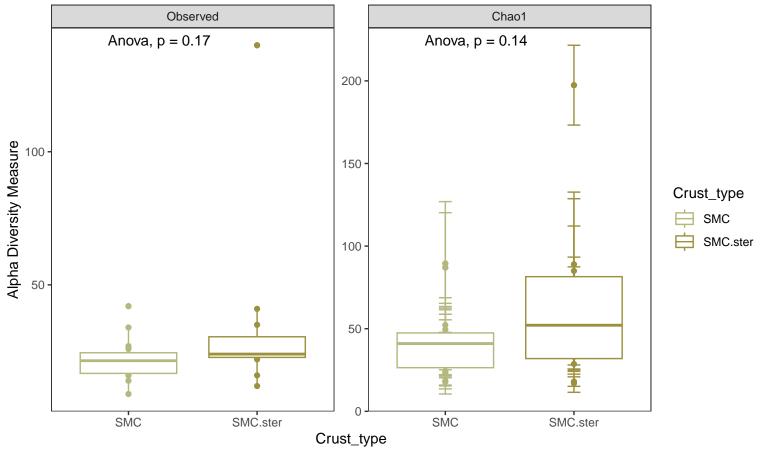
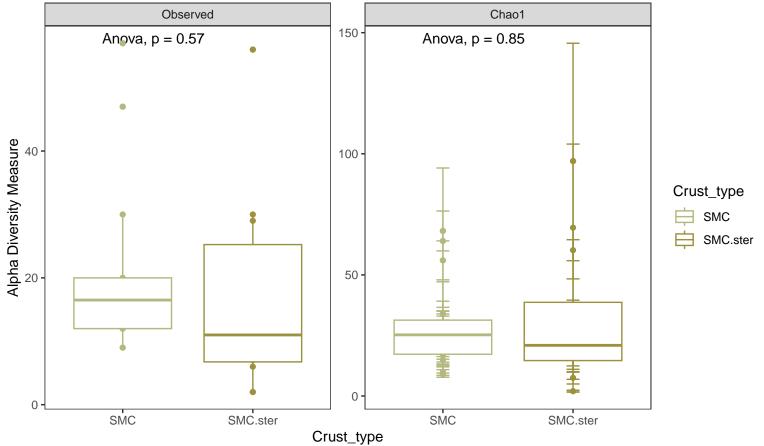
Sordariomycetes

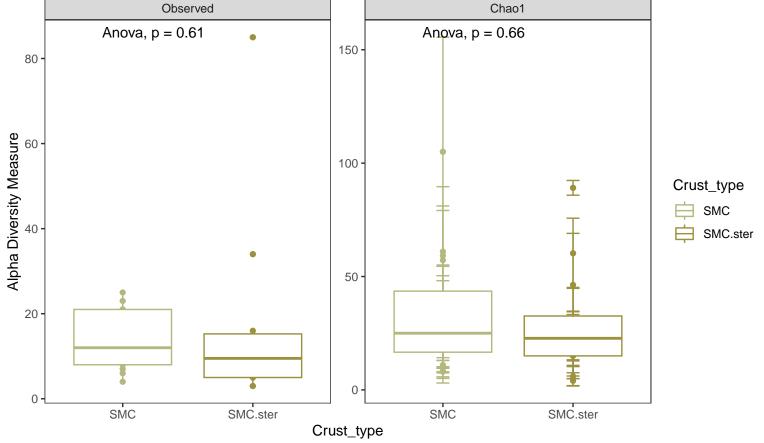


Agaricomycetes Observed Chao1 80 . Anova, p = 0.51Anova, p = 0.2250 -40 -60 -Alpha Diversity Measure Crust_type SMC 40 -SMC.ster 20 -10 -SMC SMC SMC.ster SMC.ster Crust_type

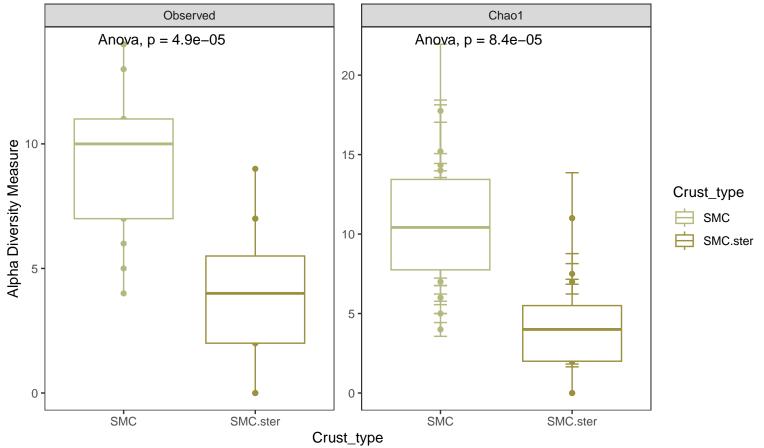
Eurotiomycetes

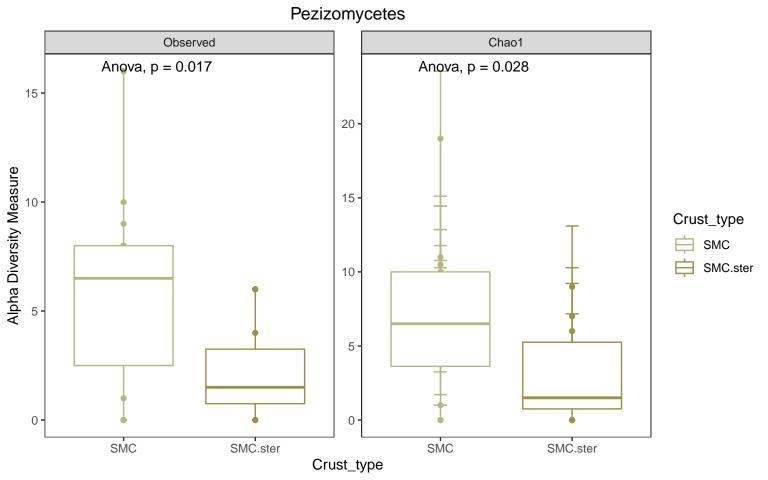


Glomeromycetes



Rhizophlyctidomycetes

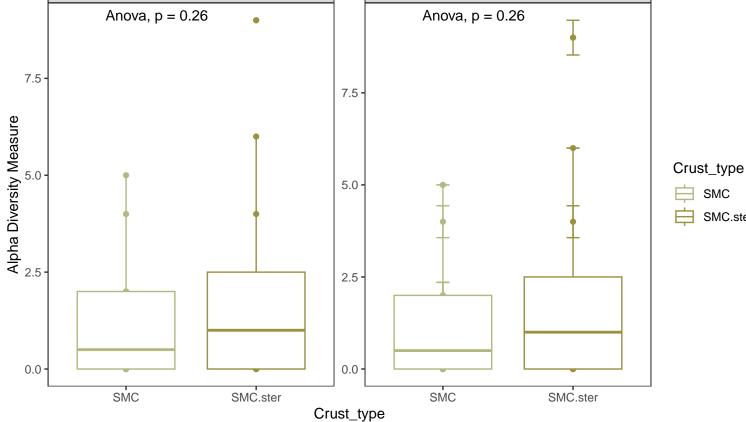




Dothideomycetes Observed Chao1 150 **-**Anova, p = 0.68Anova, p = 0.87200 -Alpha Diversity Measure 100 -150 -Crust_type SMC SMC.ster 100 -50 -50 -SMC SMC.ster SMC SMC.ster

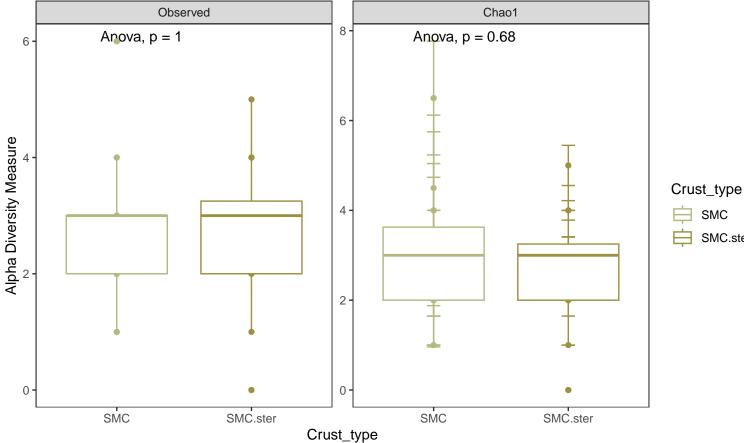
Crust_type

Lecanoromycetes Observed Chao1 Anova, p = 0.26Anova, p = 0.267.5 7.5 -

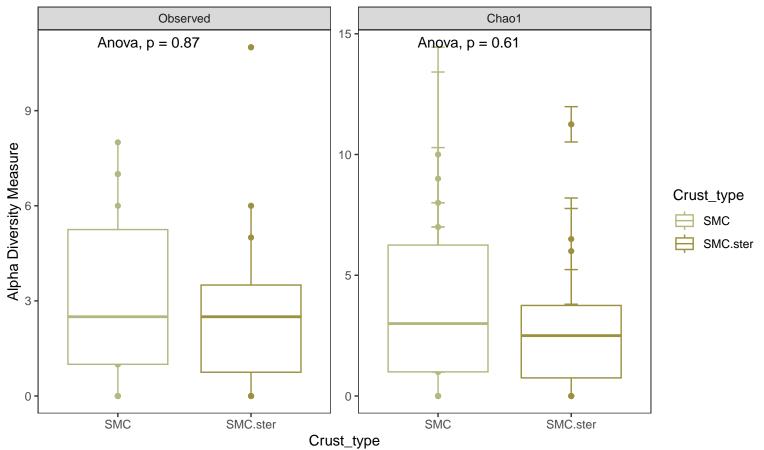


SMC SMC.ster

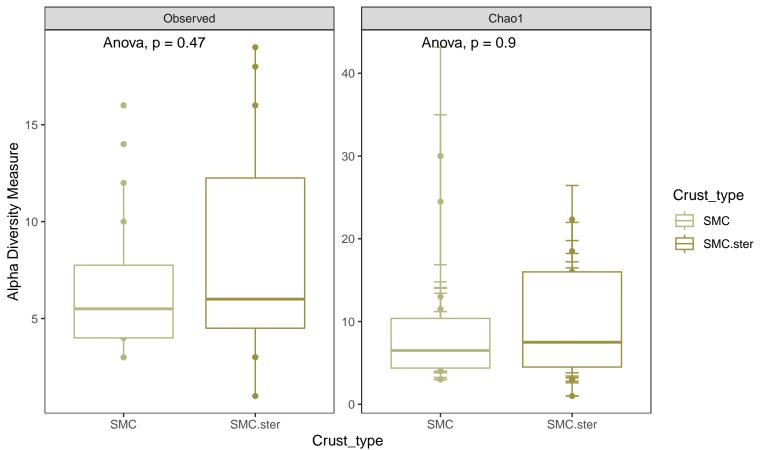
Saccharomycetes Observed Chao1 8 -Anova, p = 1Anova, p = 0.686 -6 -



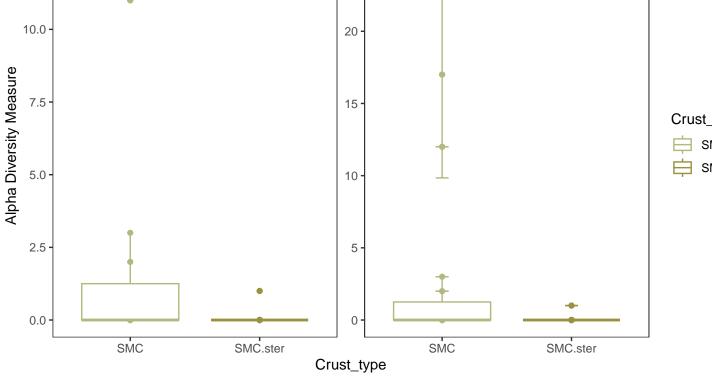
SMC SMC.ster Leotiomycetes

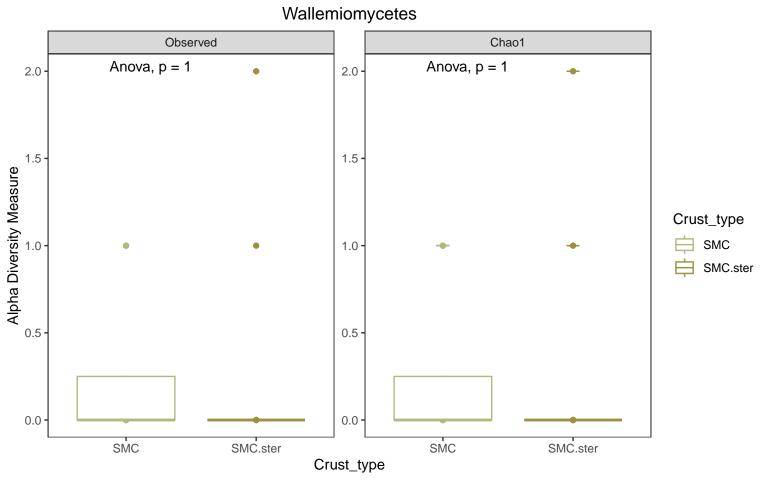


Tremellomycetes

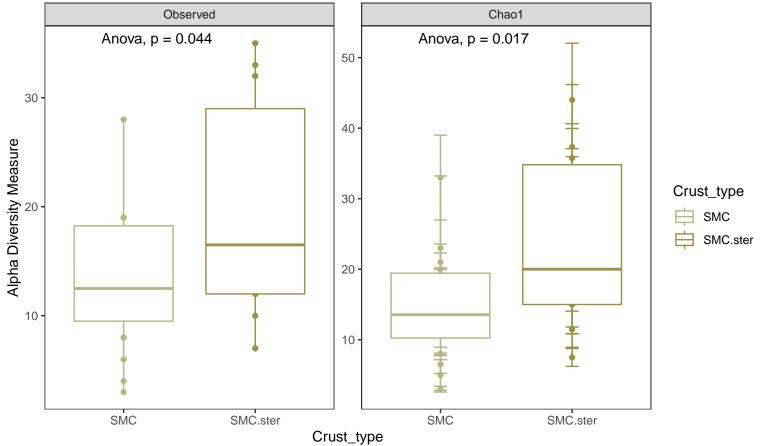


Blastocladiomycetes Observed Chao1 12.5 -25 Anova, p = 0.12Anova, p = 0.1410.0 -20 7.5 -15 **-**Crust_type SMC SMC.ster 5.0 -10 -

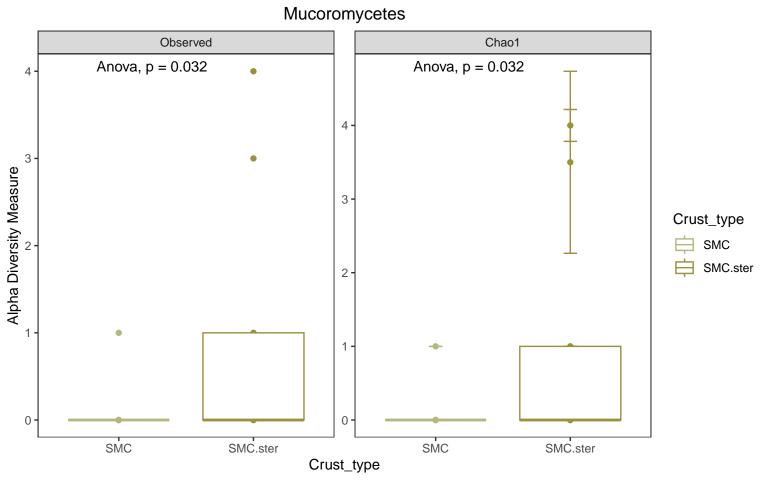


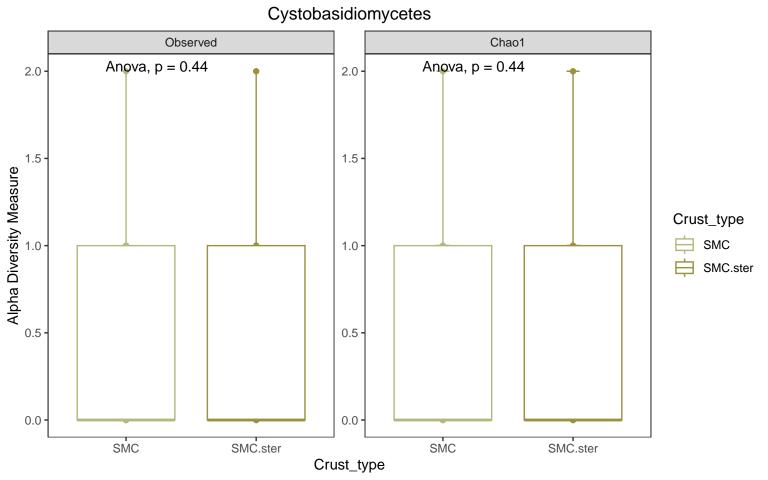


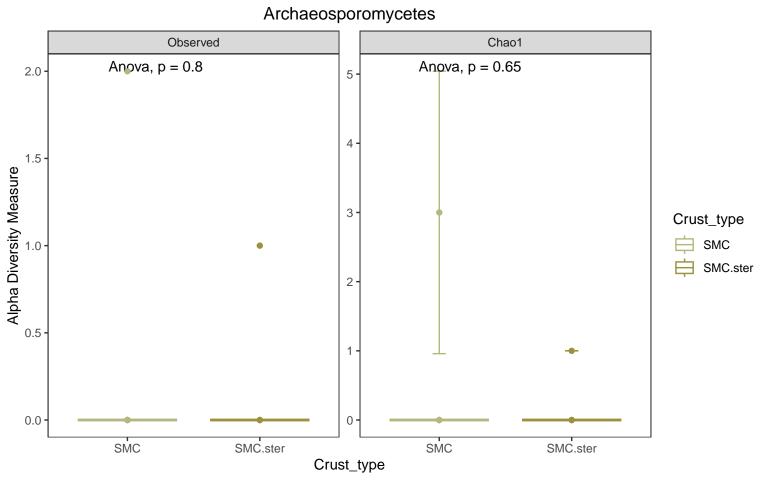
Microbotryomycetes

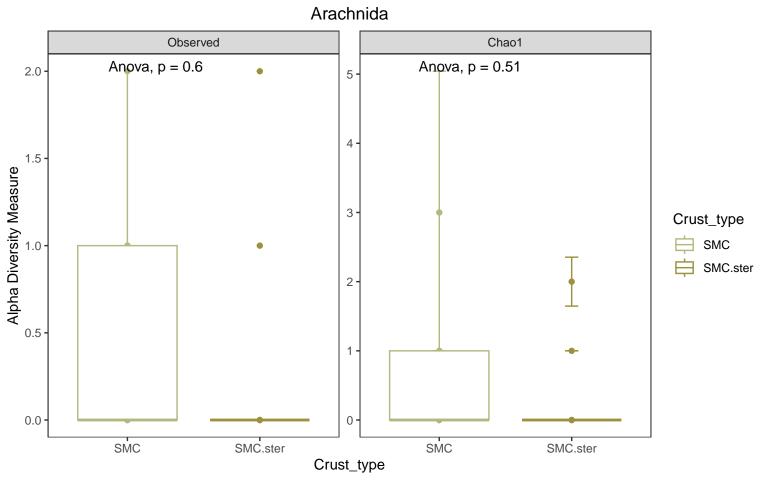


Eudicotyledonae Observed Chao1 Anova, p = 0.019Anova, p = 0.0215 -8 -4 -6 -Alpha Diversity Measure Crust_type SMC 4 -SMC.ster 2 0 -0 -SMC SMC.ster SMC SMC.ster Crust_type





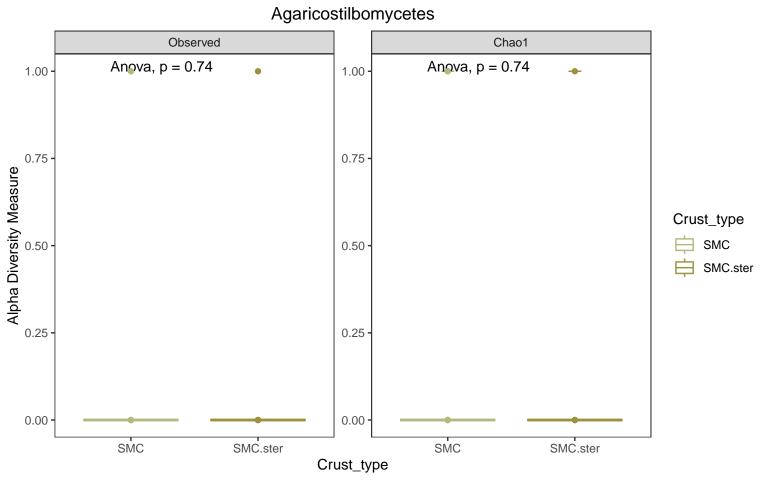




Mortierellomycetes Observed Chao1 Anova, p = 0.15Anova, p = 0.153 -3 Alpha Diversity Measure 2 -Crust_type SMC SMC.ster 0 -0 -SMC SMC.ster SMC

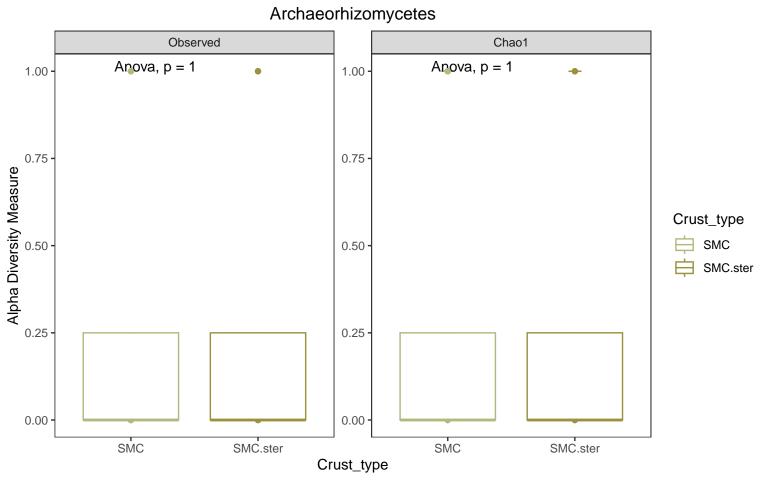
Crust_type

SMC.ster



Geminibasidiomycetes Observed Chao1 Anova, p = 0.84Anova, p = 0.841.00 -1.00 -0.75 -0.75 Alpha Diversity Measure Crust_type SMC 0.50 -SMC.ster 0.25 -0.25 0.00 -0.00 SMC SMC.ster SMC SMC.ster

Crust_type



Basidiobolomycetes Observed Chao1 Anova, p = 0.93Anova, p = 0.932.0 -2.0 -1.5 Alpha Diversity Measure 1.5 -Crust_type SMC 1.0 -SMC.ster 0.5 0.5 -0.0 -0.0 -SMC SMC SMC.ster SMC.ster

Crust_type

