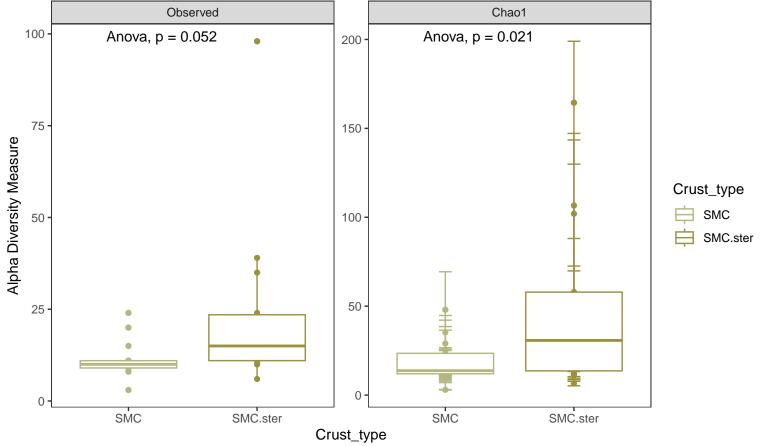
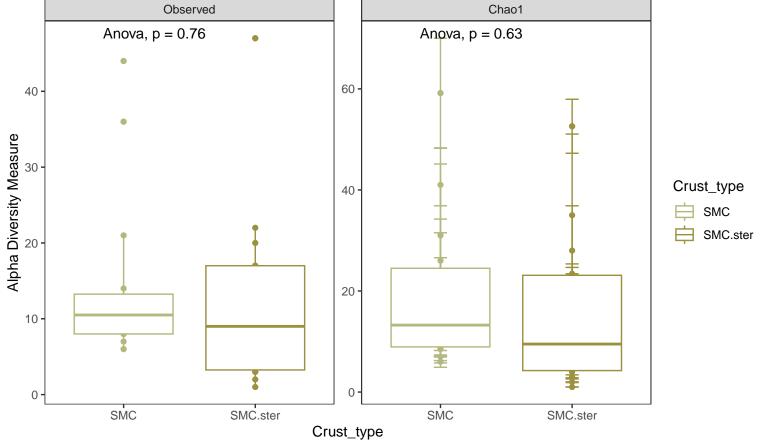
Sordariomycetes

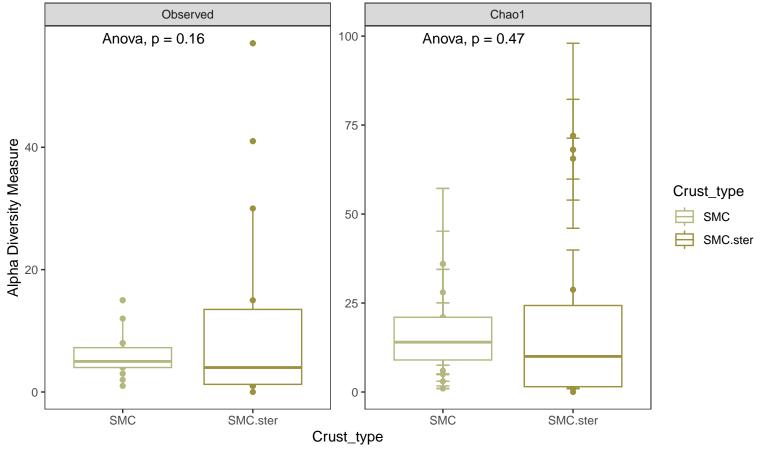


Agaricomycetes Observed Chao1 Anova, p = 0.7Anova, p = 0.9830 -75 -Alpha Diversity Measure Crust_type 50 -SMC SMC.ster 25 -10 -0 -SMC SMC SMC.ster SMC.ster

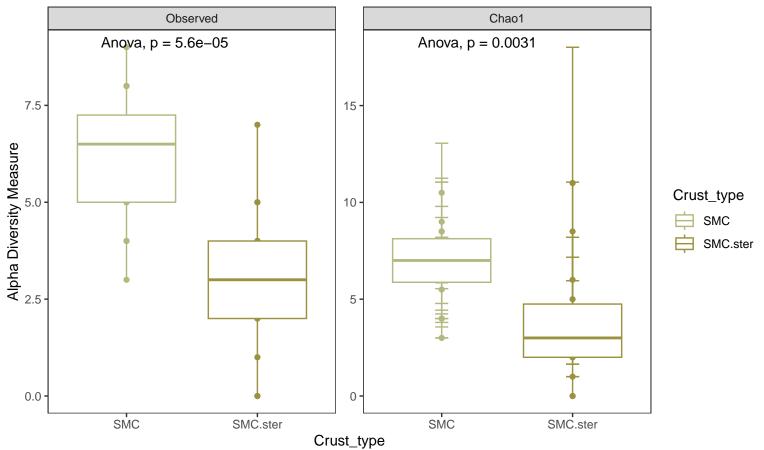
Eurotiomycetes

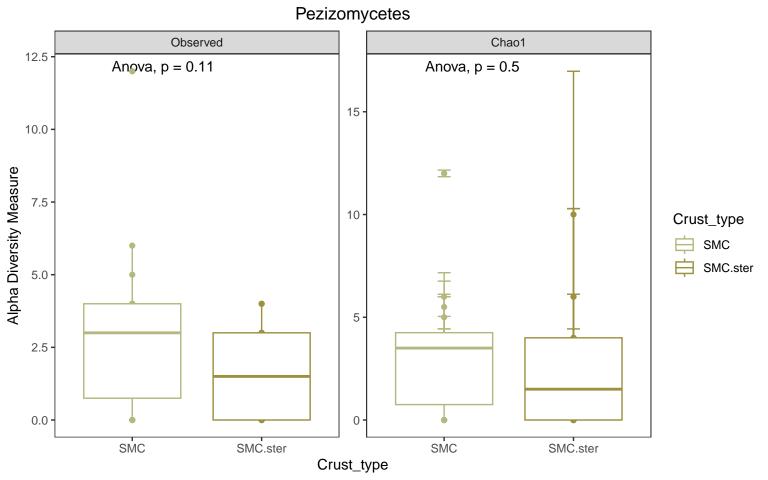


Glomeromycetes

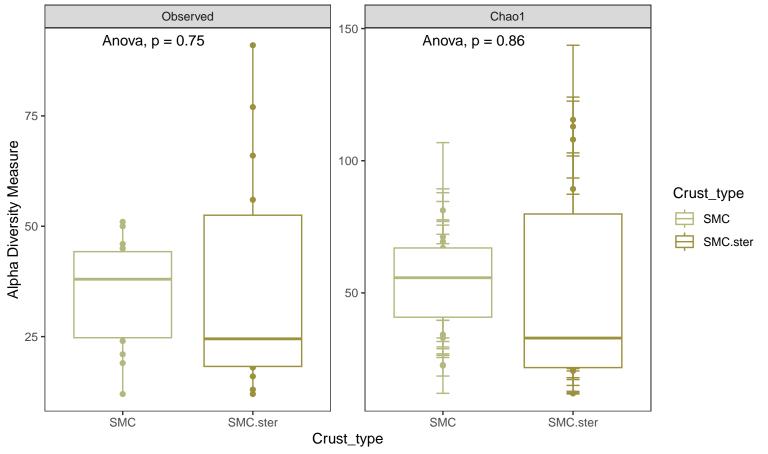


Rhizophlyctidomycetes

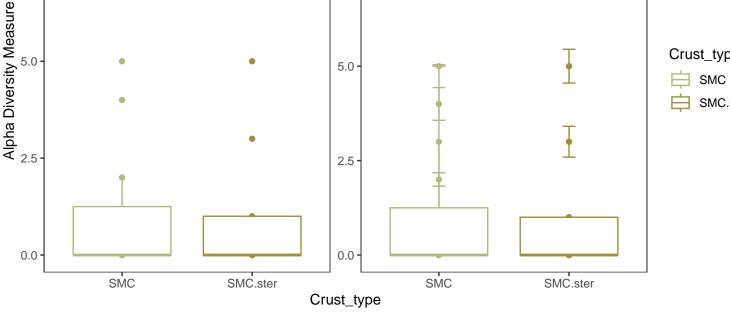




Dothideomycetes



Lecanoromycetes Observed Chao1 Anova, p = 0.58Anova, p = 0.647.5 7.5 -Alpha Diversity Measure Crust_type 5.0 -SMC SMC.ster 2.5 -

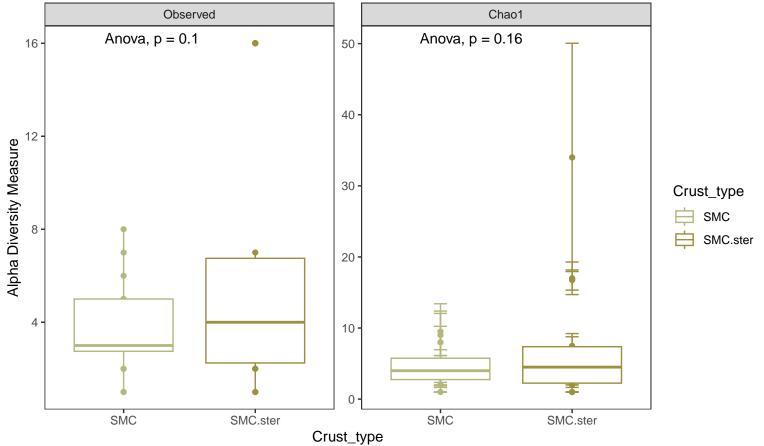


Saccharomycetes Observed Chao1 Anova, p = 0.065Anova, p = 0.0475 -8 -4 -6 -Alpha Diversity Measure Crust_type SMC 4 -SMC.ster 2 0 -0 -SMC.ster SMC SMC SMC.ster Crust_type

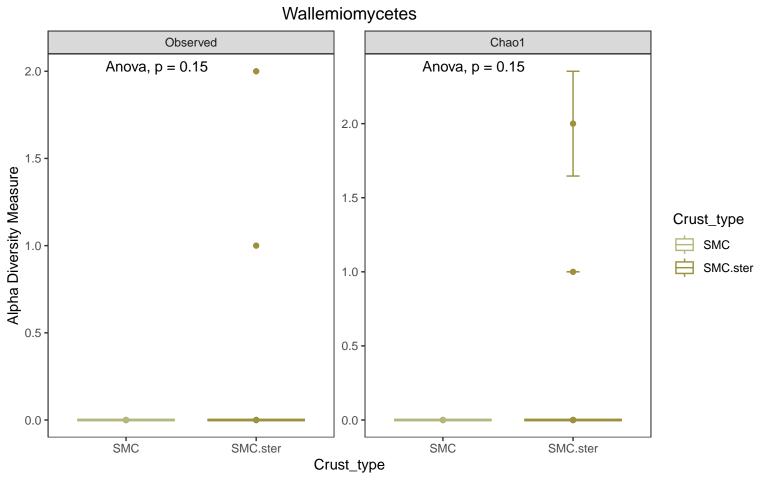
Leotiomycetes Observed Chao1 Anova, p = 0.79Anova, p = 0.728 -20 -6 Alpha Diversity Measure 15 -Crust_type SMC SMC.ster 10 -5 -0 -0 -SMC SMC SMC.ster SMC.ster

Tremellomycetes

Observed



Blastocladiomycetes Observed Chao1 12.5 -Anova, p = 0.11Anova, p = 0.19 -10.0 -Alpha Diversity Measure 7.5 -Crust_type SMC SMC.ster 5.0 -2.5 -0 -0.0 -SMC SMC SMC.ster SMC.ster



Microbotryomycetes Observed Chao1 Anova, p = 0.021Anova, p = 0.009130 -40 -Alpha Diversity Measure 30 -Crust_type SMC SMC.ster 20 -10 -

0 .

Crust_type

SMC.ster

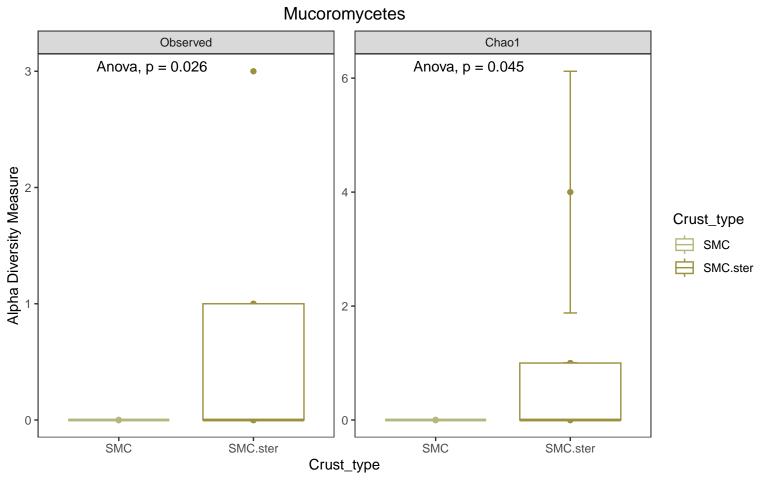
SMC

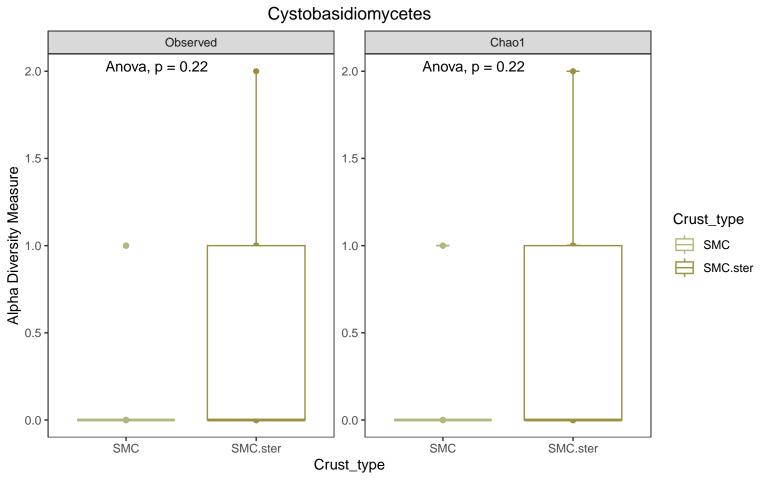
SMC.ster

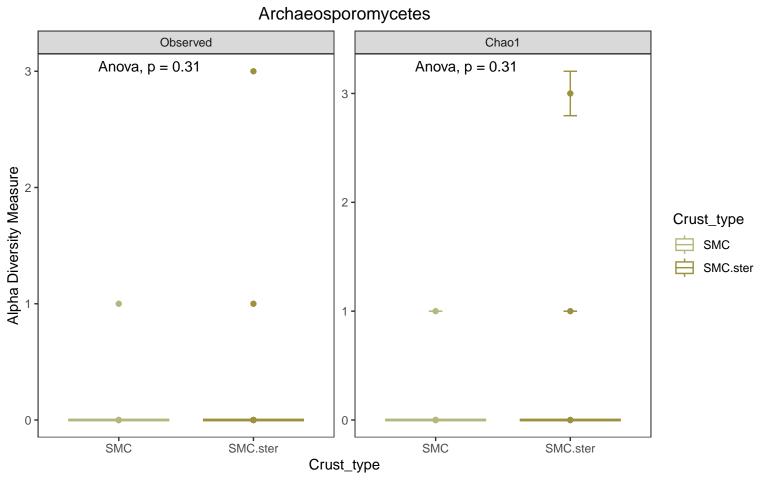
0 -

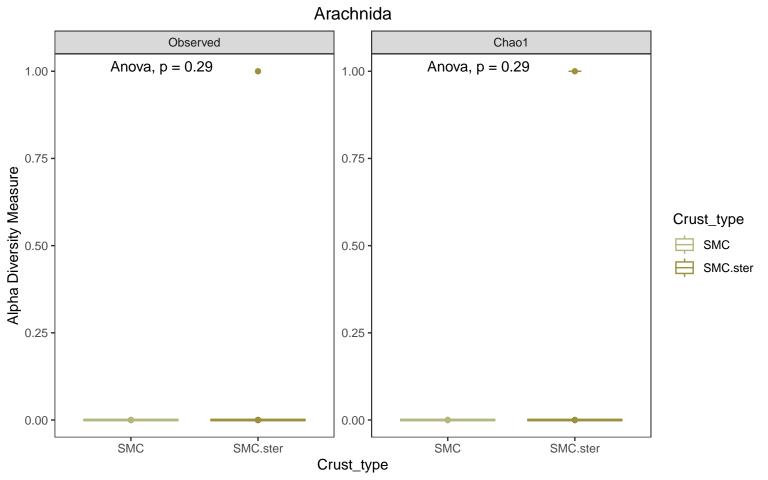
SMC

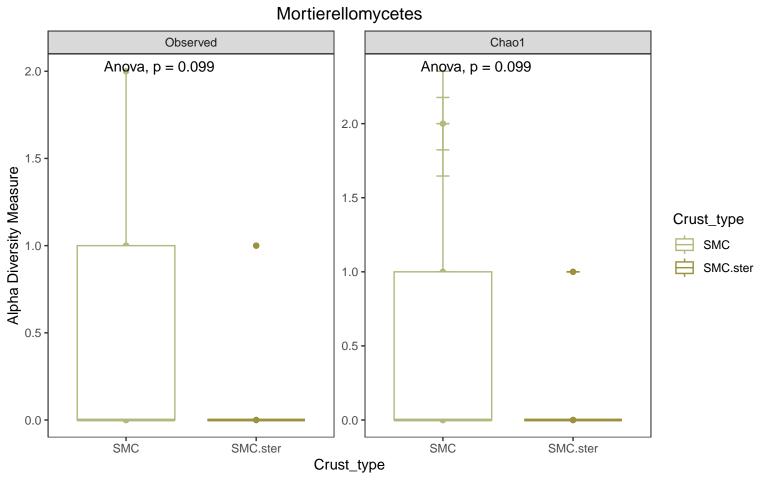
Eudicotyledonae Observed Chao1 Anova, p = 0.041Anova, p = 0.0343 -5 -4 -Alpha Diversity Measure 3 -Crust_type SMC SMC.ster 2 -1 -0 -0 -SMC SMC.ster SMC SMC.ster Crust_type

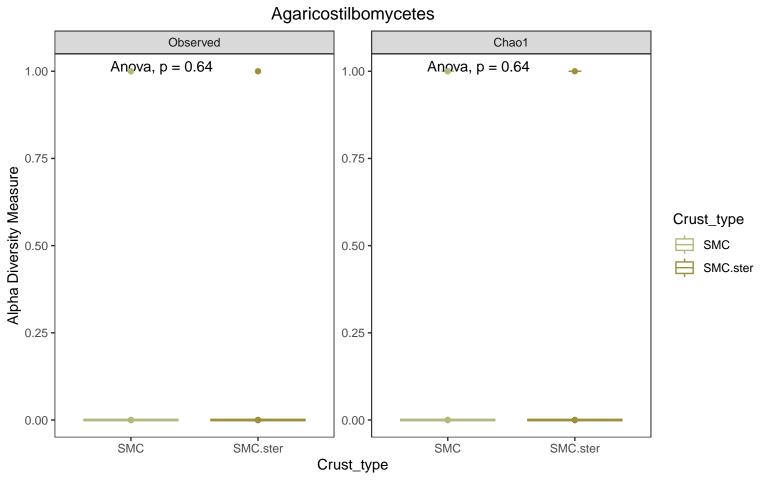




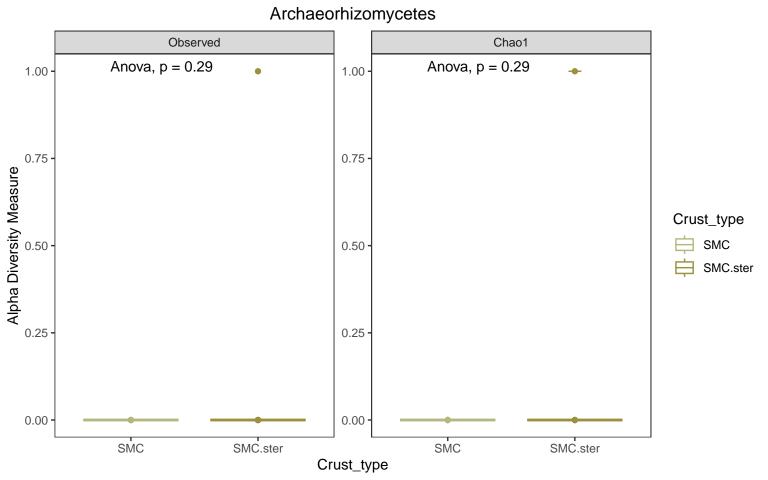








Geminibasidiomycetes Observed Chao1 Anova, p = 0.29Anova, p = 0.291.00 -1.00 -0.75 -0.75 Alpha Diversity Measure Crust_type SMC 0.50 -SMC.ster 0.25 -0.25 0.00 -0.00 -SMC SMC.ster SMC SMC.ster



Basidiobolomycetes Observed Chao1 Anova, p = 0.16Anova, p = 0.162.0 -2.0 -1.5 Alpha Diversity Measure 1.5 -Crust_type SMC SMC.ster 1.0 -0.5 0.5 -0.0 -0.0 -SMC SMC SMC.ster SMC.ster Crust_type

