



# Fire, NOT Grazing

## influences soil microbial diversity in a semi-arid savanna

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WITWATERSRAND

**In African Savannas, fire and herbivory coexist**

**Prescribed fires are used as a management tool**

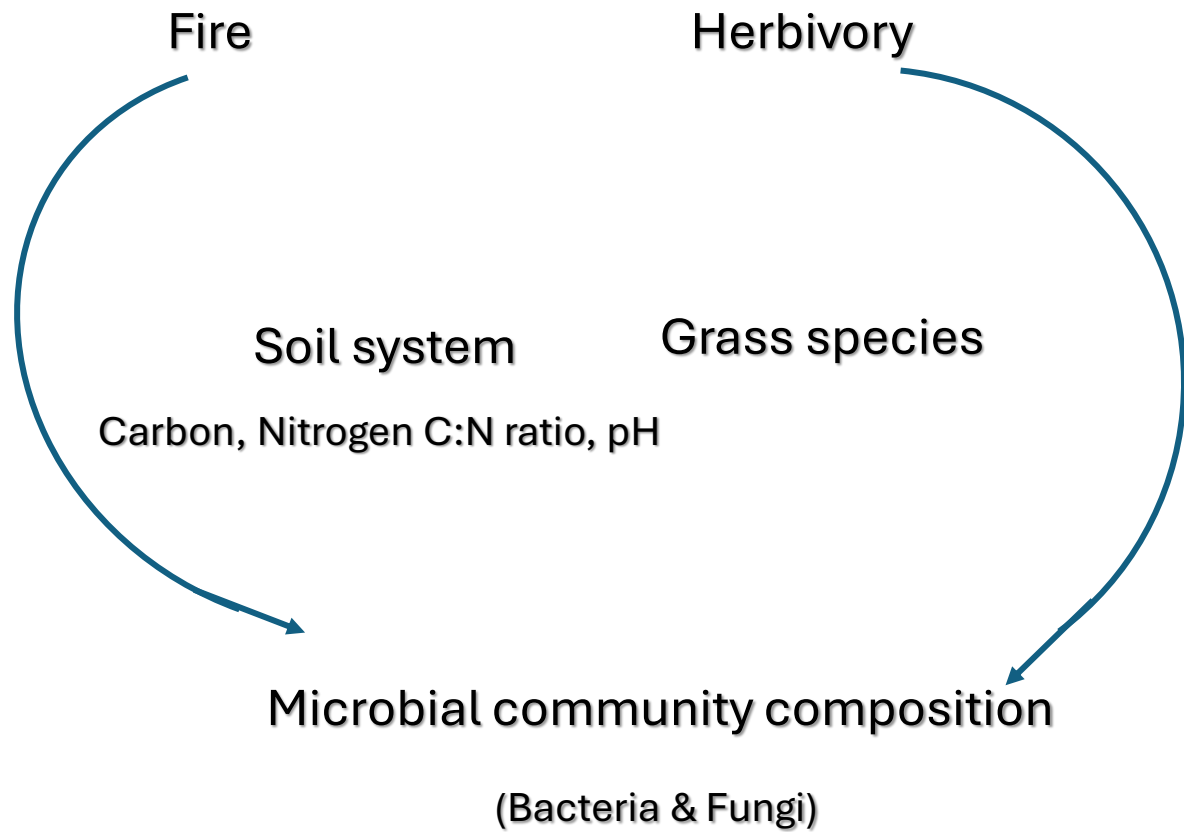
**Africa is the last continent with large mammal herbivores**



**How do fire and herbivory affect below-ground microbial communities?**

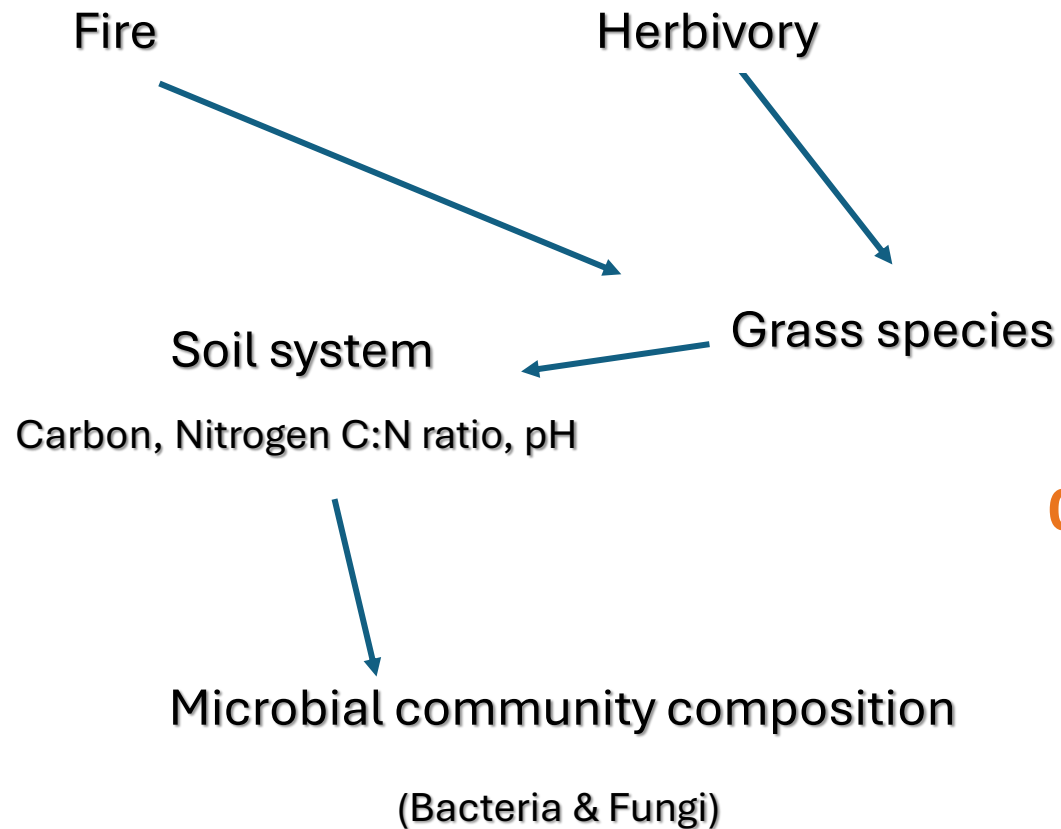


# Research questions & hypothesis



- 01** Are fire, herbivory, or plant species the strongest drivers of microbial community structure?

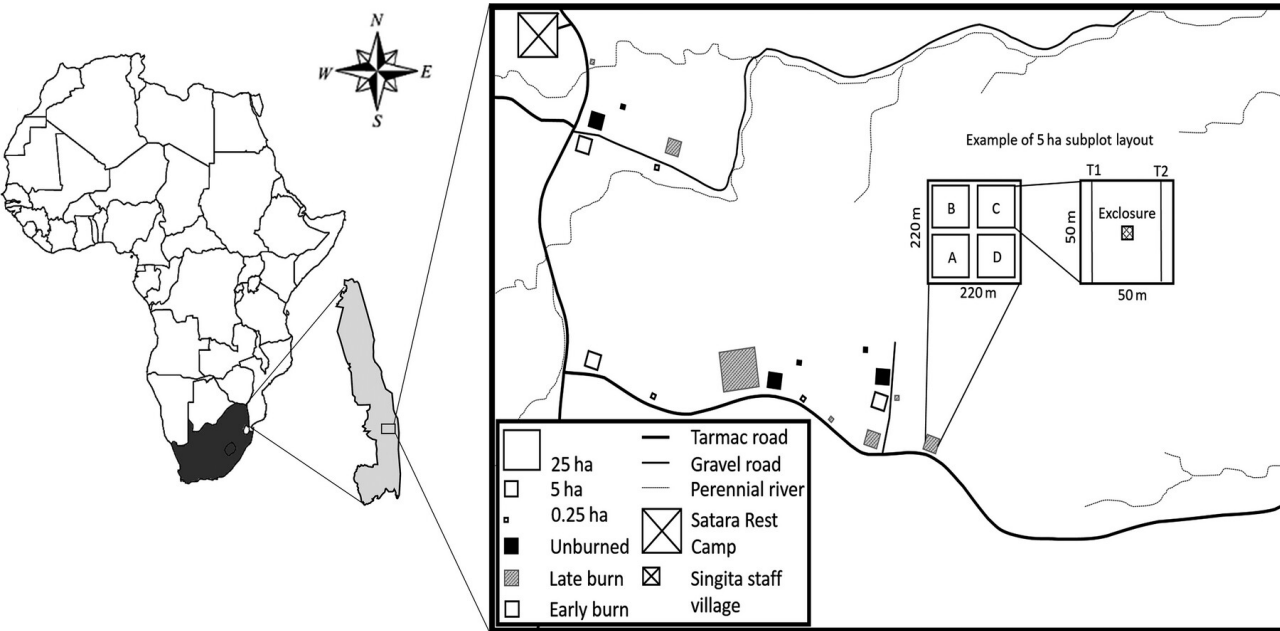
# Research questions & hypothesis



**02** Is there any evidence that differences in microbial composition is mediated by soil properties

**03** Which functional groups are most responsive to soil chemical properties?

# Satara Fire Experiment



**Figure 1:** Landscape-level experimental burn plot in the Satara, Kruger National Park (Donaldson et al. 2017)

# Data collection

Soils collected  
under each grass  
tussock



Used Deays® Power  
Soil® Kit (Qiagen) to  
extract the microbial  
DNA

Soil pH, % C, %N  
and C:N ratio were  
determined



Used Illumina  
Nextera tagged  
primers for Bacterial  
16S rRNA & fungal ITS  
regions

These are the assumptions of linear regression

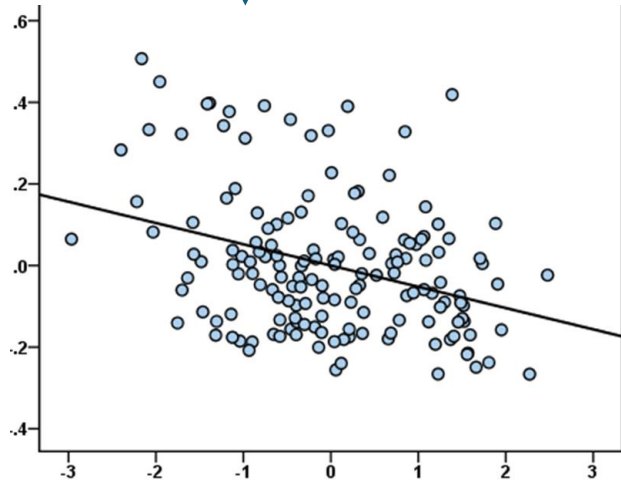
But where did they come from?



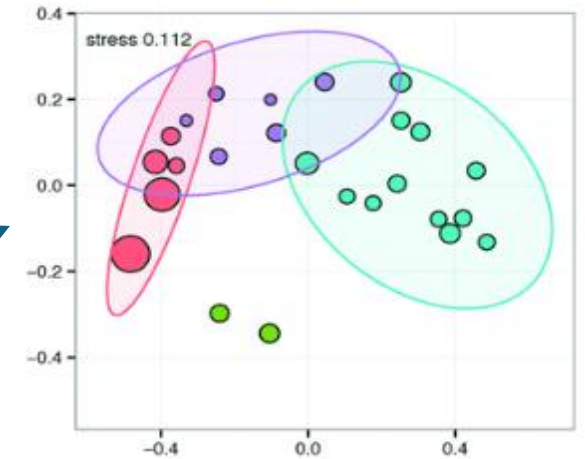
- Assumption 1
- Assumption 2
- Assumption 3
- Assumption 4

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`S = glmmTMB (S ~ Fire*Herbivory + (1 | Grass), family = poisson, data)`



Plot of best model



Bray-Curtis dissimilarity

#649 **Permanova**  
(`vegan::adonis2`)

**Table 1. The effect of fire and herbivory on soil properties**

Model	Effect	Coefficient	Confidence Intervals
<b>Soil properties</b>			
pH <sub>(KCl)</sub> ~ Herbivory + (1   Grass)	Ungrazed > Grazed	0.08	0.001 - 0.15
C ~ Fire*Herbivory + (1   Grass)	Control < Fire	-0.37	-2.46 -1.72
	Ungrazed > Grazed	3.90	1.56 - 6.24
	Fire*Control < Herbivory*Ungrazed	-2.77	-6.03 - 0.48
N ~Fire*Herbivory + (1   Grass)	Control < Fire	-0.06	-0.23 - 0.11
	Ungrazed > grazed	0.33	0.13 - 0.52
	Fire*Control < Herbivory*Ungrazed	-0.27	-0.54 - 0.01
CN ~ Fire + (1   Grass)	Control > Fire	0.77	0.24 - 1.29

# Soil properties

A

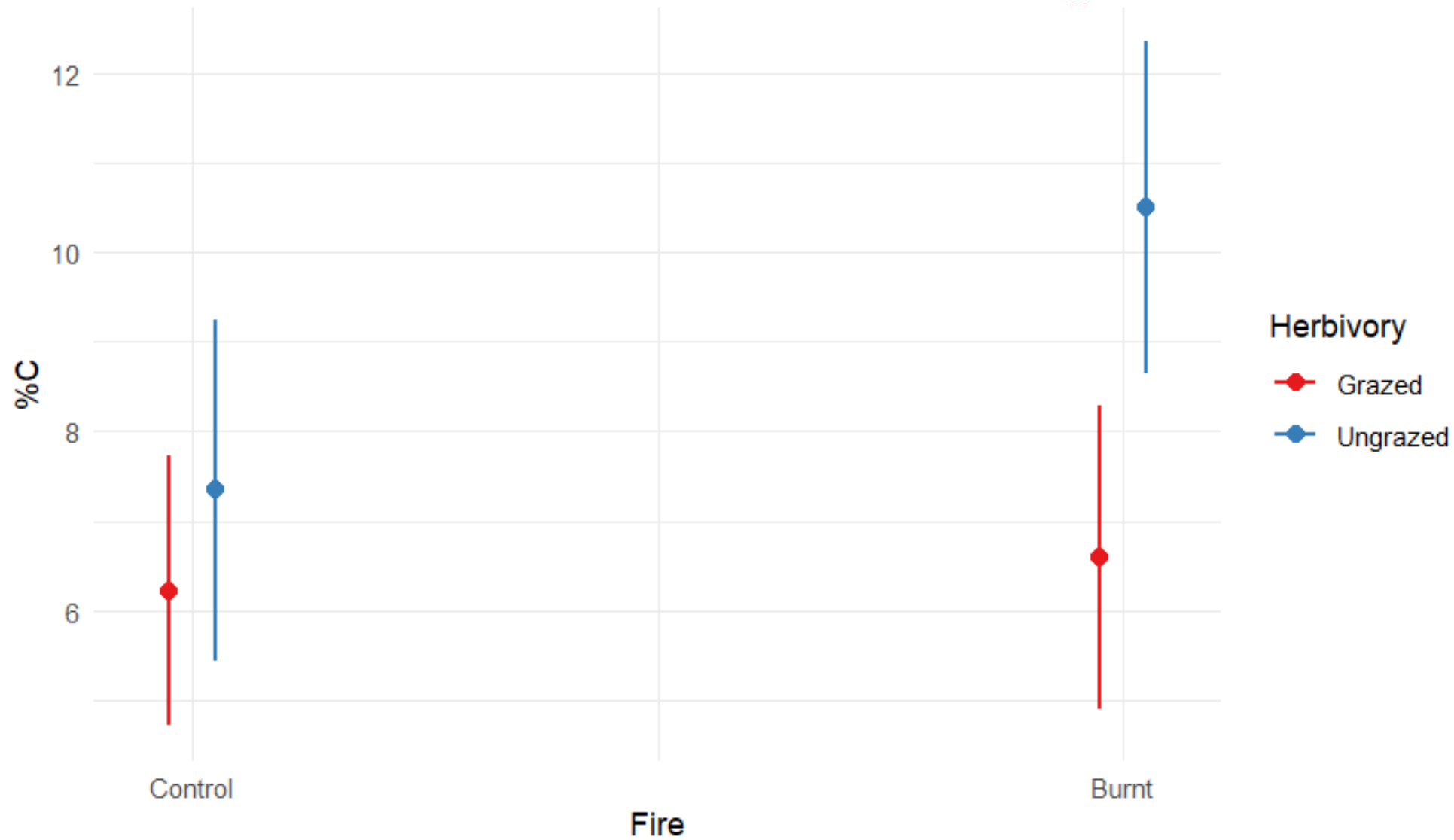


Figure 2. Visualisation of mixed effects of Influence of (A) grazing on soil pH and (B) % total Carbon

# Soil properties

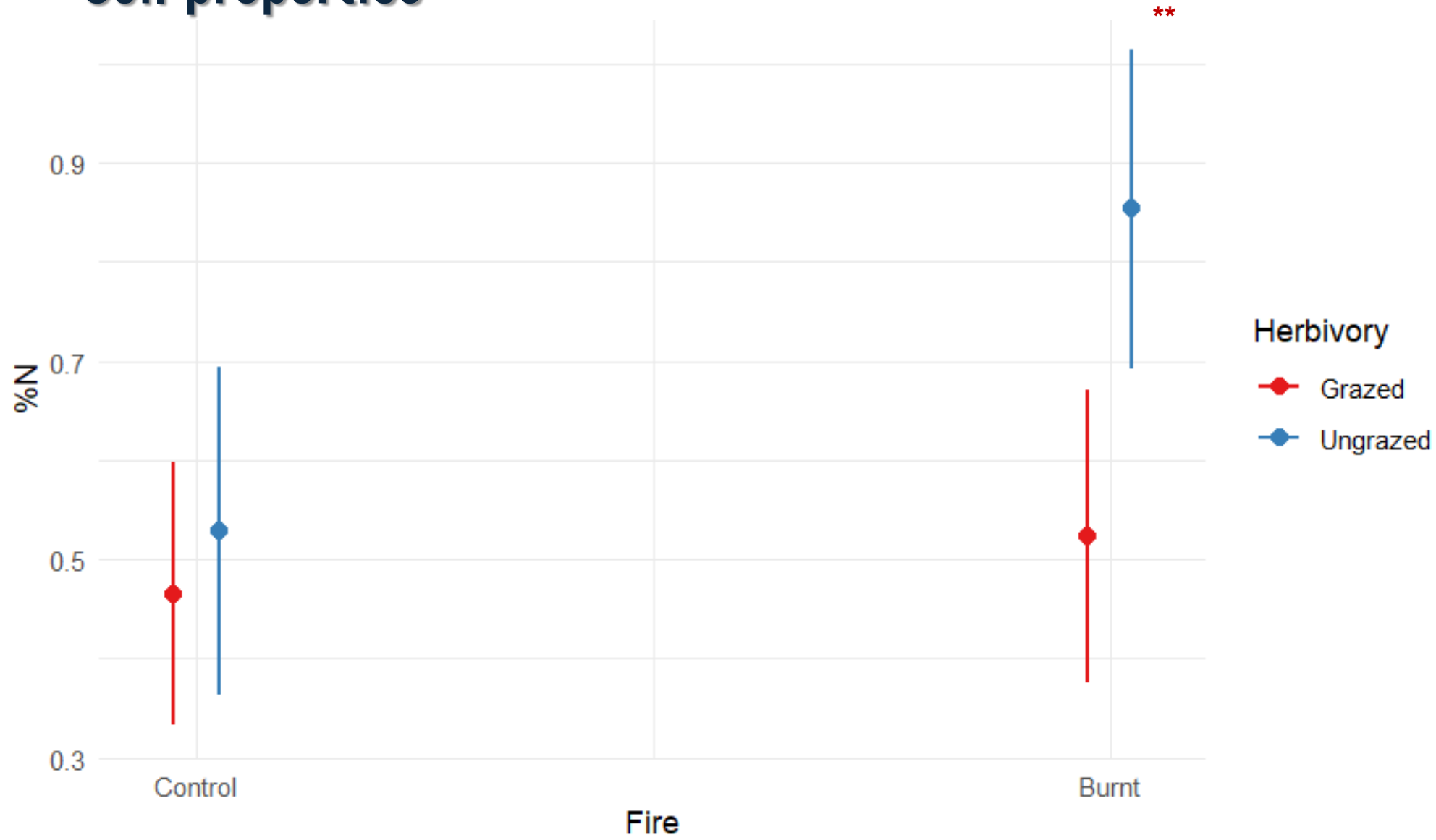
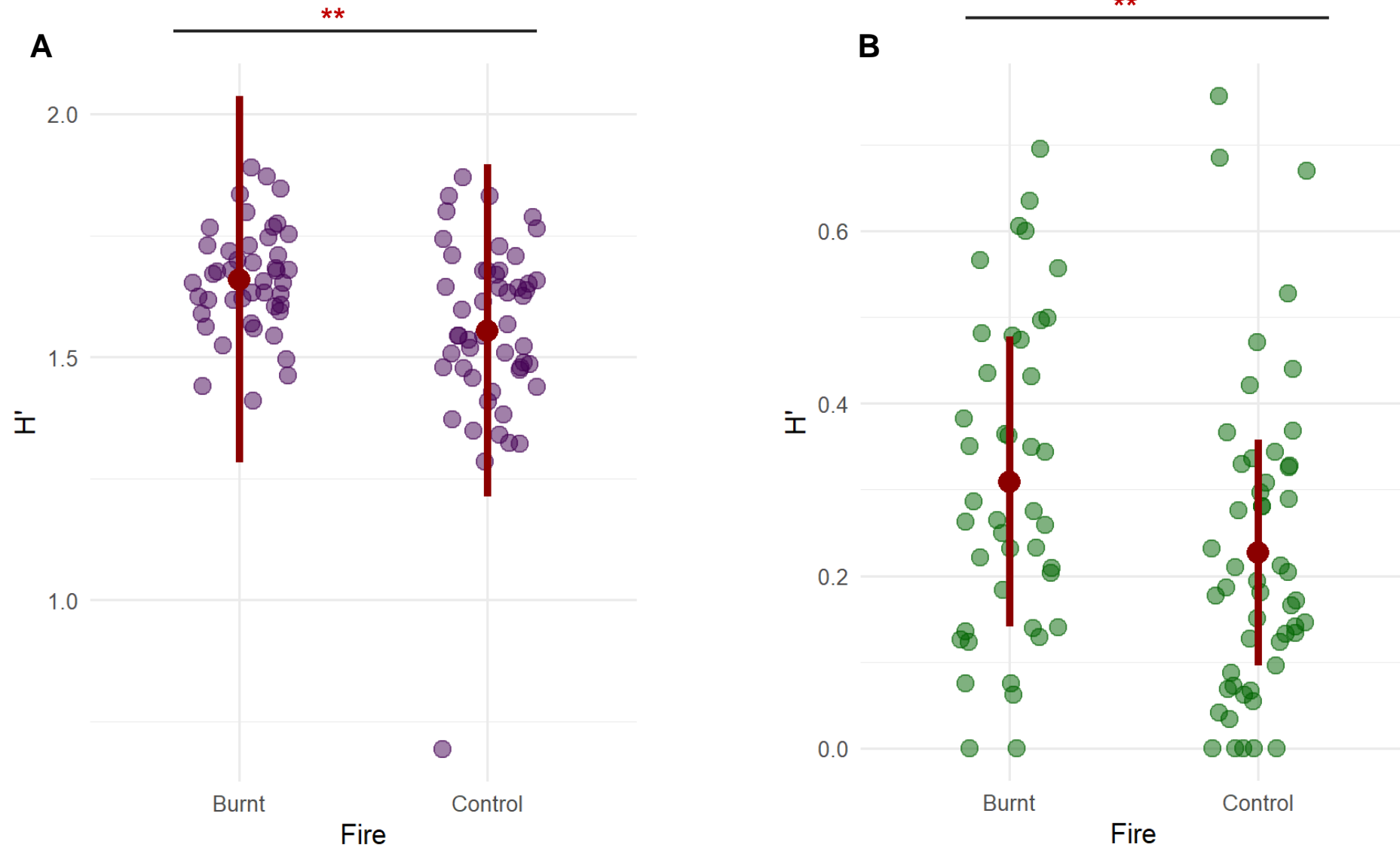


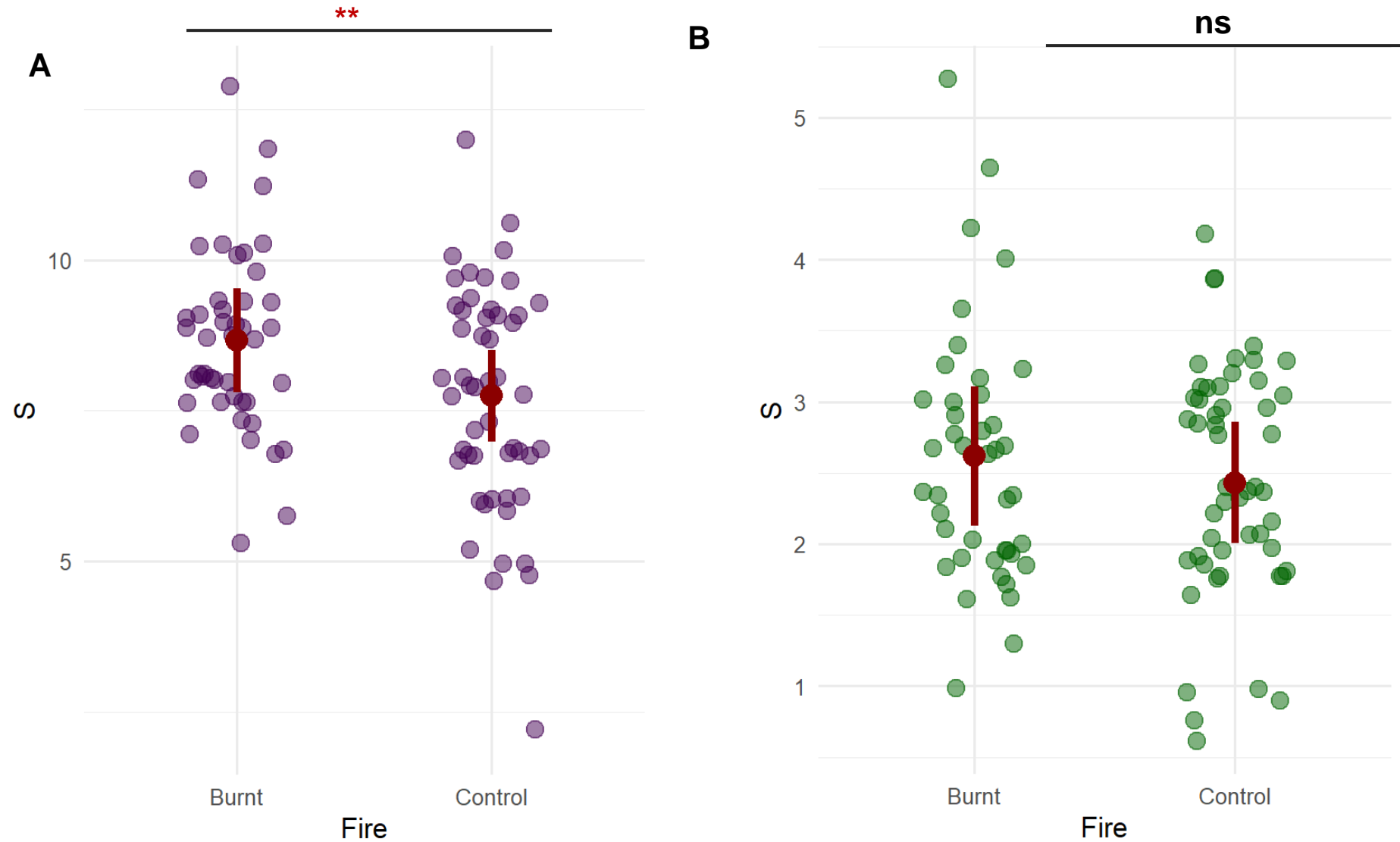
Figure 3. Visualisation of mixed effects of Influence of grazing on (A) % total Nitrogen and (B) C: N ratio

**Table 1. The effect of fire and herbivory on alpha diversity**

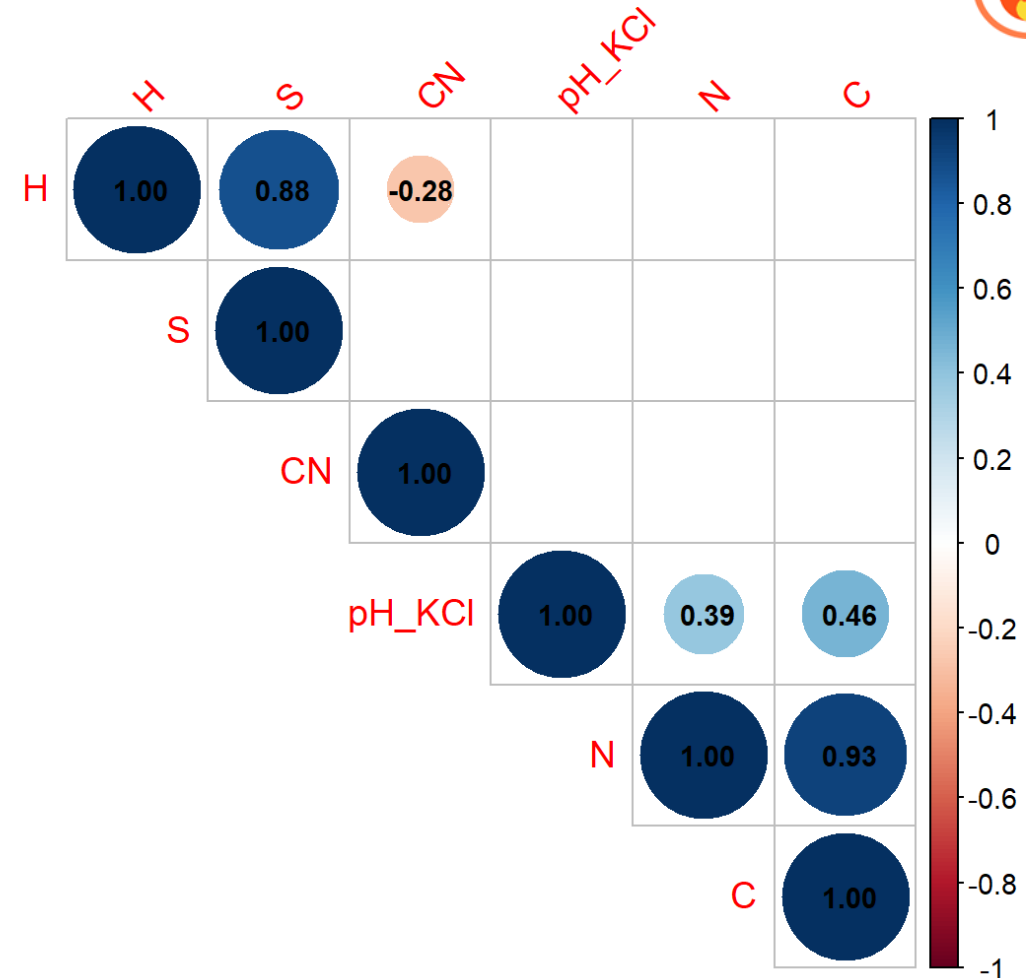
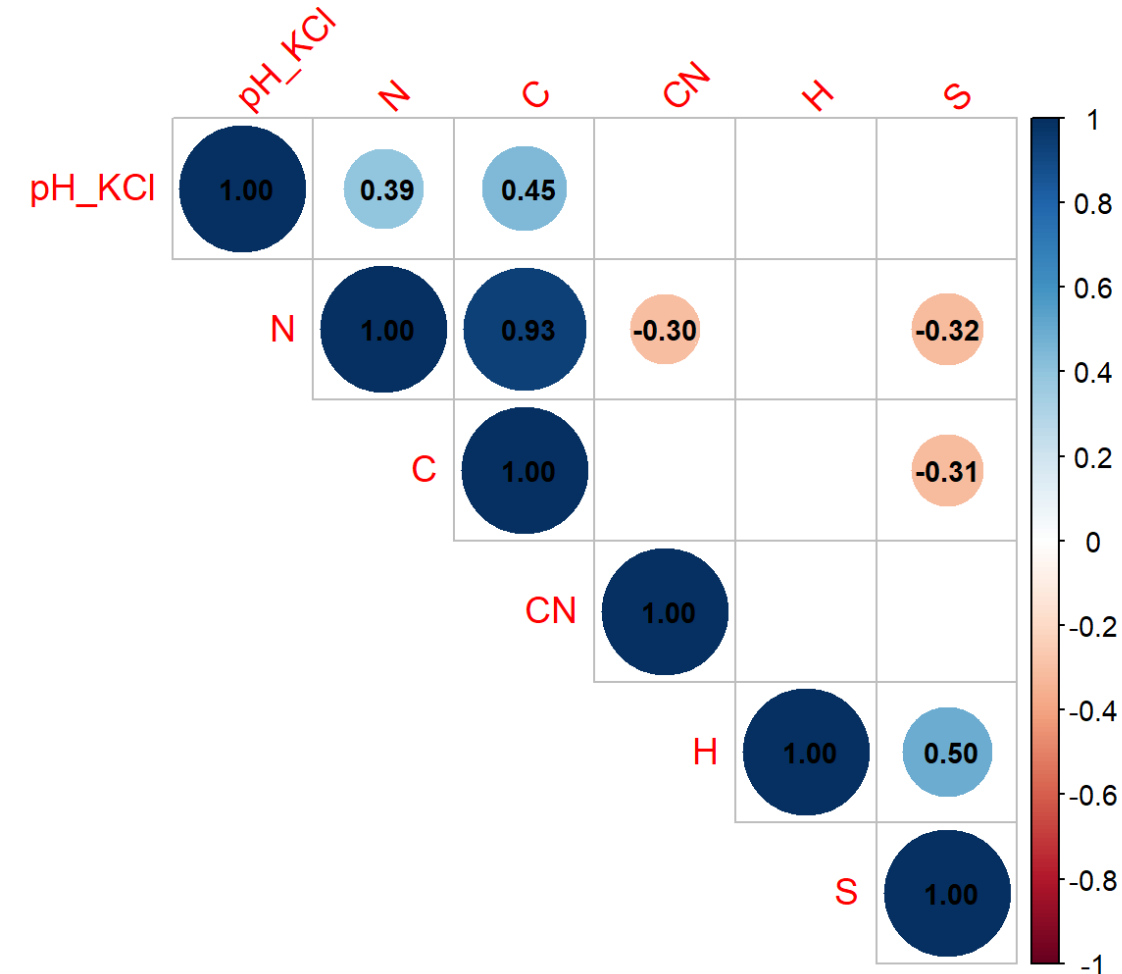
Model	Effect	Coefficient	Confidence Intervals
<b>Bacterial Alpha Diversity</b>			
H ~ Fire + (1   Grass)	Control < Fire	-0.11	-0.11-0.04
S ~ Fire + (1   Grass)	Control < Fire	-0.89	-0.25 -0.027
<b>Fungal Alpha diversity</b>			
H ~ Fire + (1   Grass)	Control < Fire	-0.08	-0.15-0.01
S ~ Fire + (1   Grass)	Control < Fire	-0.07	-0.33-0.18

 $\alpha$ 

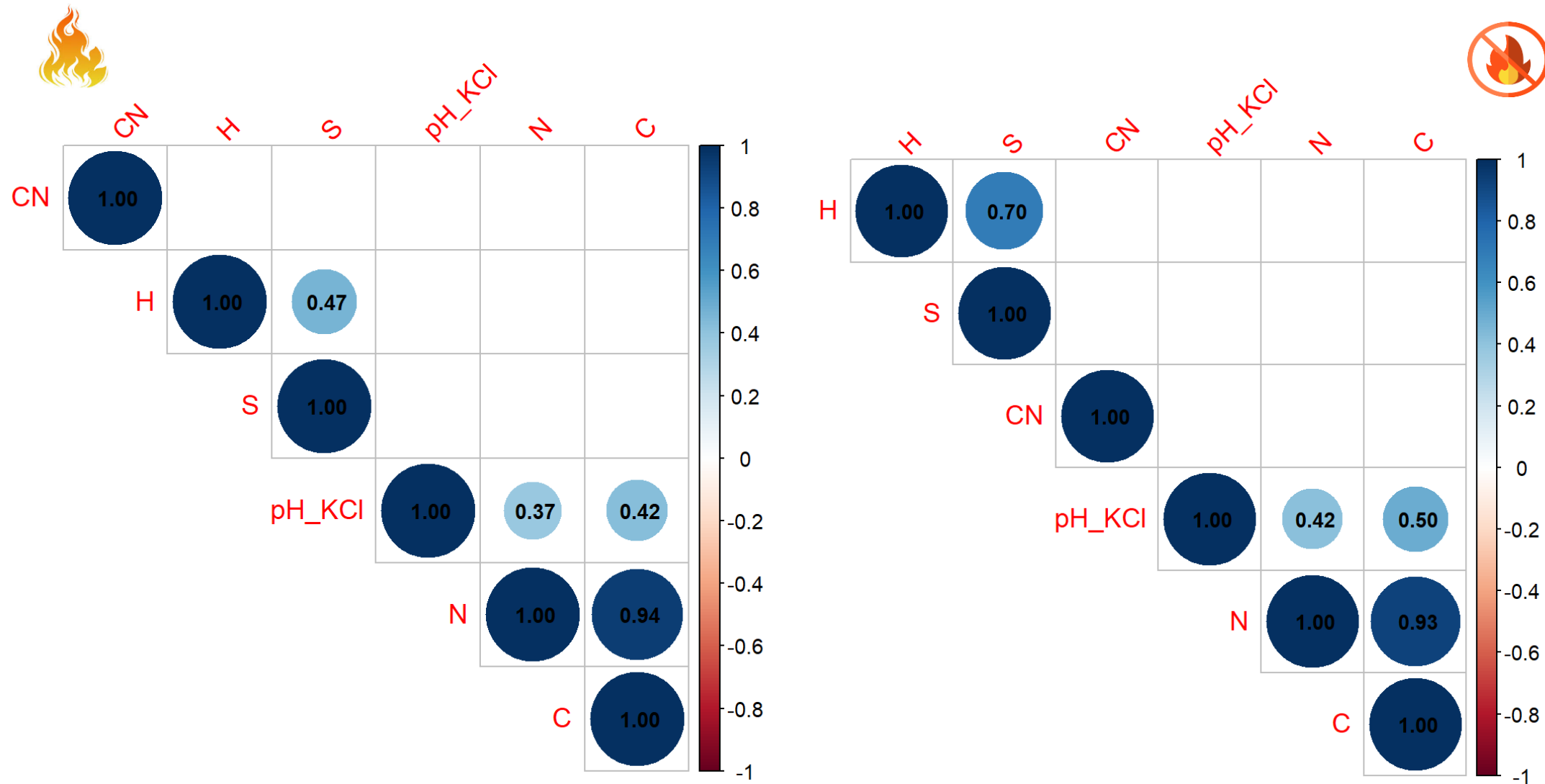
**Figure 6.** Shannon diversity ( $H'$ ) index for **(A)** Bacterial and **(B)** Fungal communities associated with the burned plots



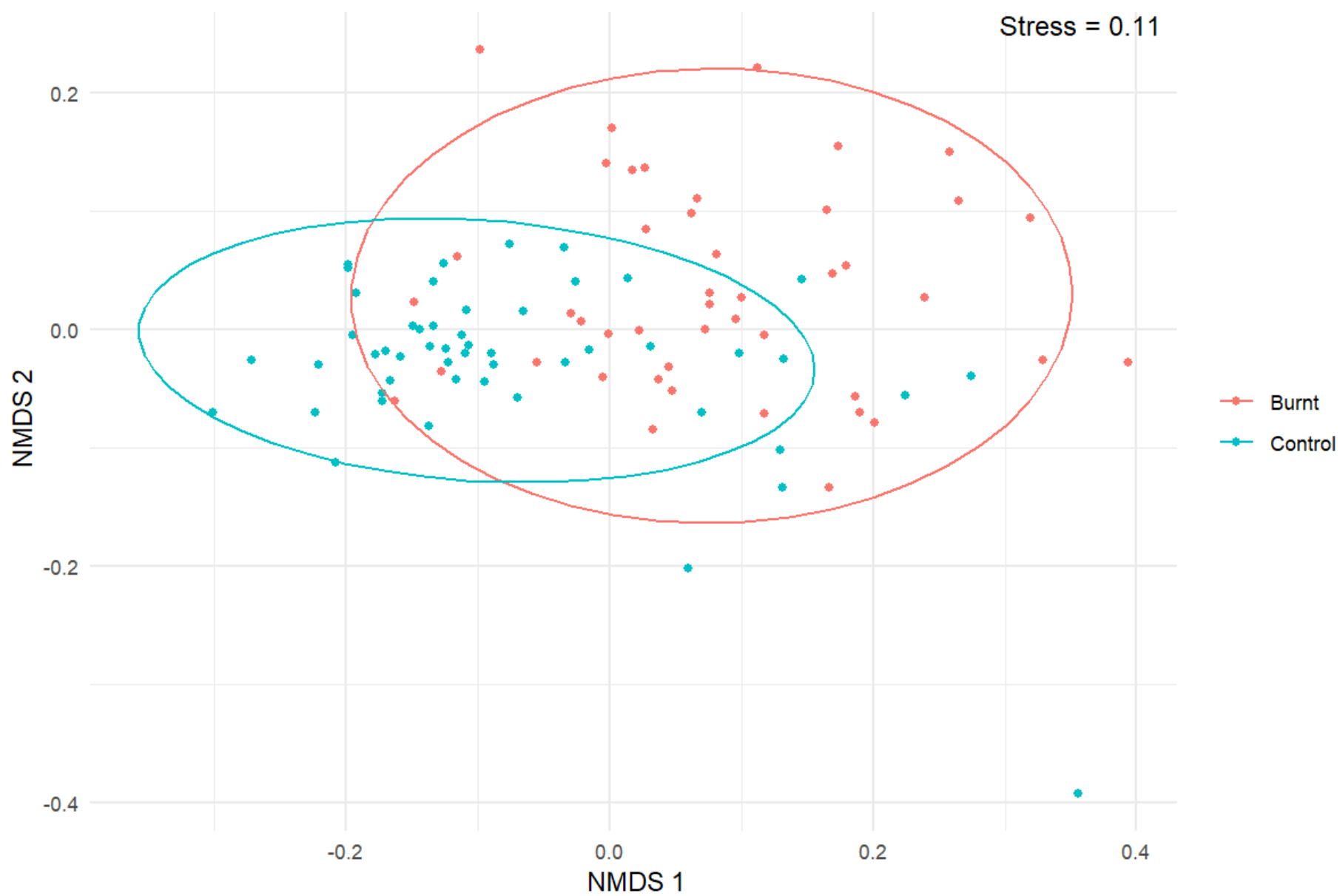
**Figure 7.** Microbial richness (S) index for **(A)** Bacterial and **(B)** Fungal communities associated with the burned plots



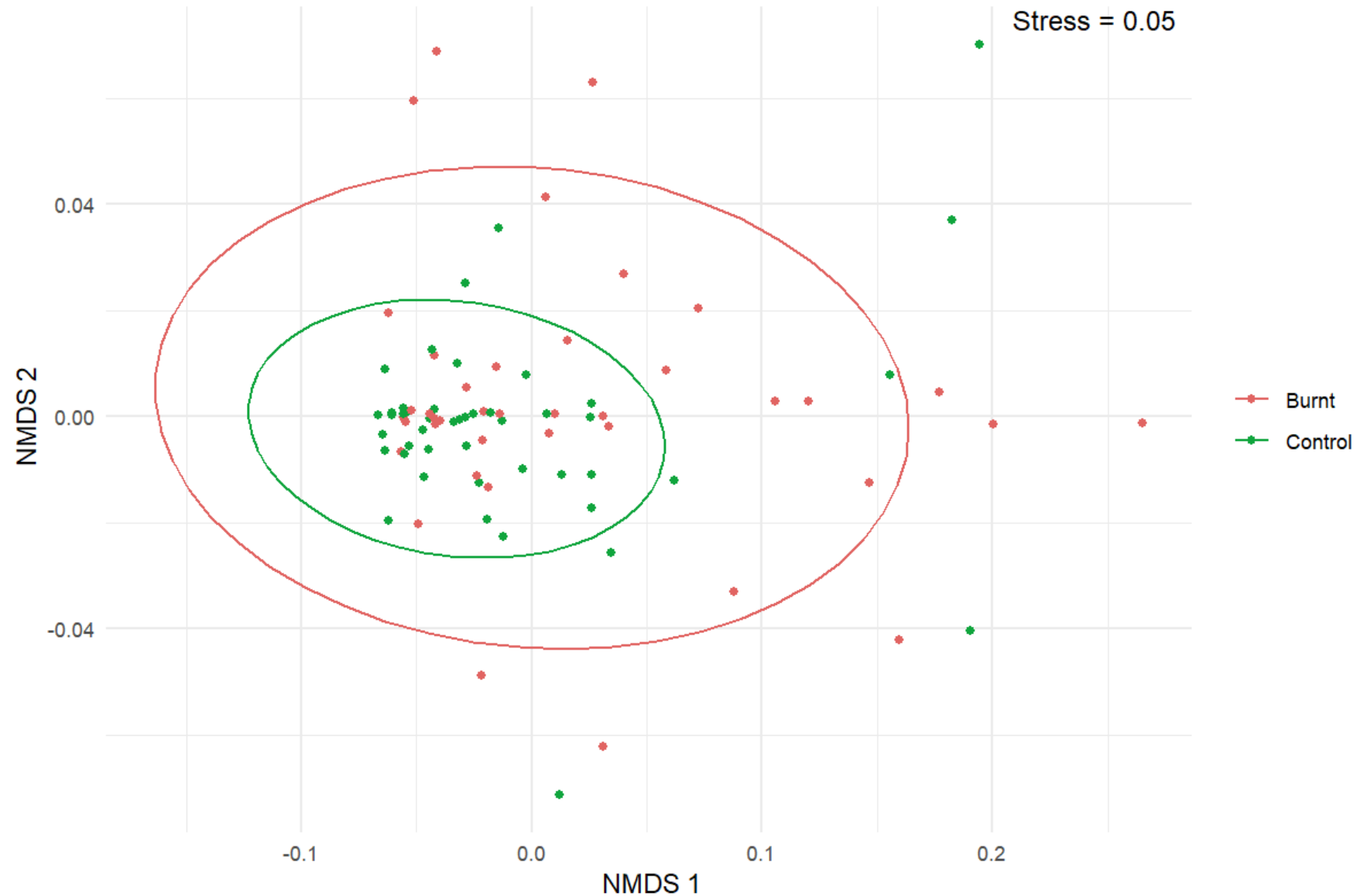
**Figure 8.** Spearman correlation between soil properties and bacterial alpha diversity index for the burned treatment in Satara



**Figure 9.** Spearman correlation between soil properties and fungal alpha diversity index for the burned treatment in Satara



**Figure 10.** Non-Metric Multi-Dimensional Scaling (NMDS) plot of Bacterial communities associated with soils collected from the Satara fire experimental burned plots (PERMANOVA:  $p$  value = 0.001)

$\beta$ 

**Figure 11.** Non-Metric Multi-Dimensional Scaling (NMDS) plot of **(A)** Bacterial and **(B)** Fungal communities associated with soils collected from the Satara fire experimental burned plots (p value =0.07)



# SUMMARY

1. The presence of **herbivores** decreases soil pH, total Carbon and Nitrogen
2. **Fire and not herbivory** affects microbial diversity and community composition
3. Soil properties do mediate changes in microbial composition

