

# Bacteriomes of Peaches and Cover Crops

By Derek Newberger

Ph.D. Candidate



# Acknowledgements

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- CSU: Minas and Vivanco Lab Collaboration
- USDA: Dr. Daniel Manter, Soil Management and Sugarbeet Research
- Producer Representative: Bruce Talbott, Talbott Farms, Inc
- Western Sustainable Agriculture Research and Education (SARE) project #SW20-910 with title 'Developing sustainable peach orchard soil microbiome management practices to control replant disease syndrome'



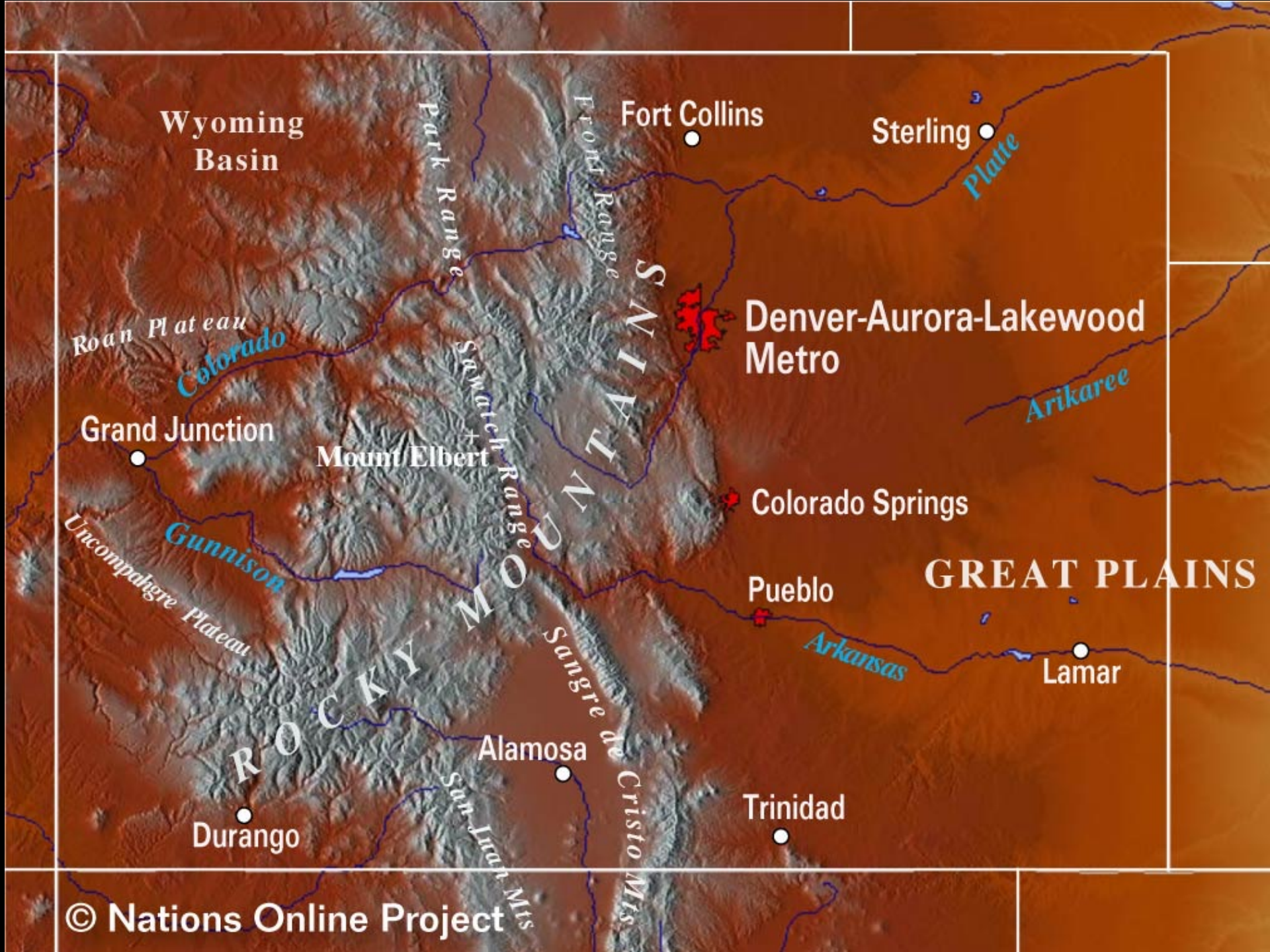
United States  
Department of  
Agriculture







\$40 M annually





# The problem

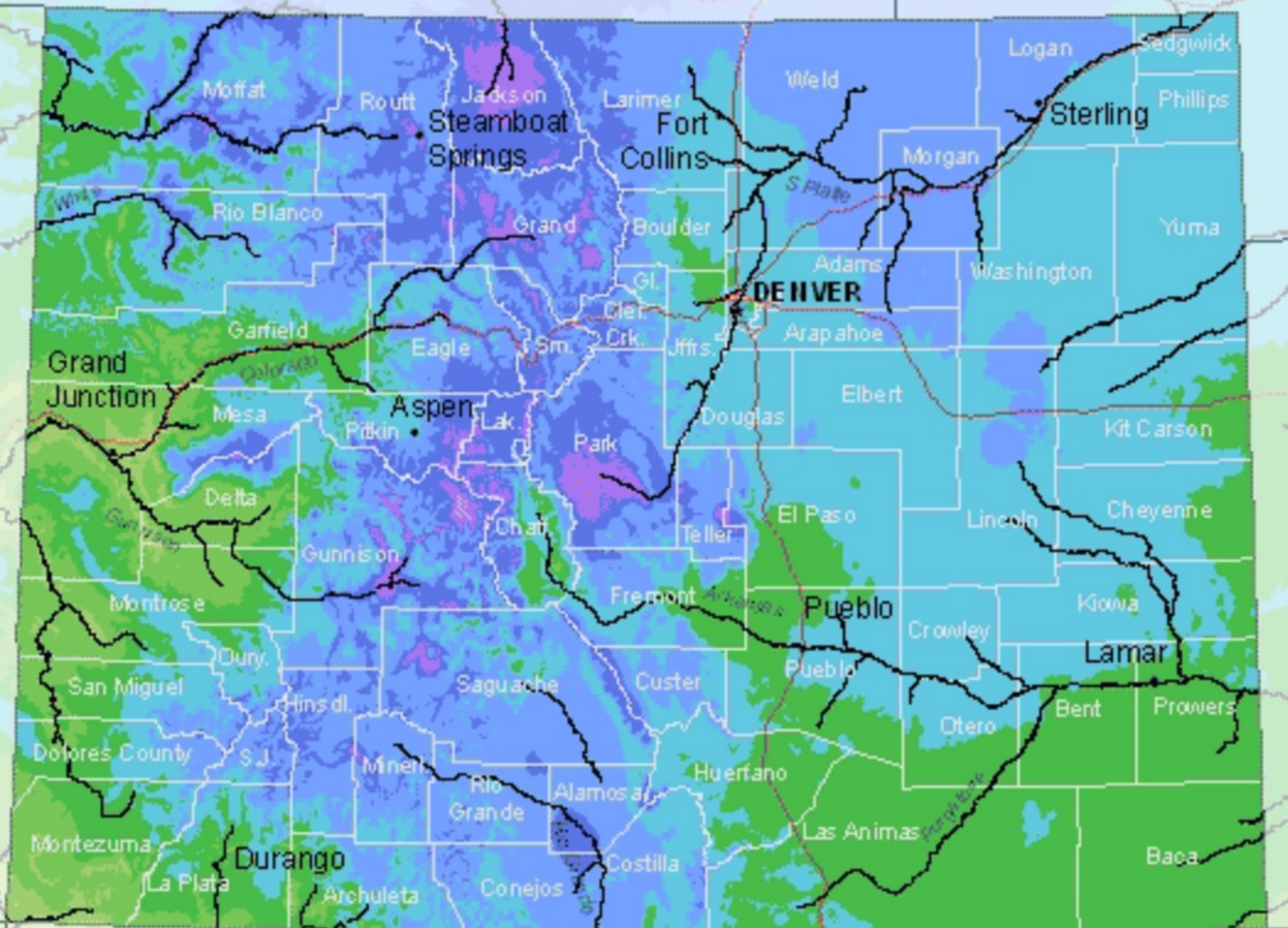




# Colorado

## Average Annual Extreme Minimum Temperature 1976-2005

Temp (F)	Zone	Temp (C)
40 to -35	3a	40 to -37.2
-35 to -30	3b	-37.2 to -34.4
-30 to -25	4a	-34.4 to -31.7
-25 to -20	4b	-31.7 to -28.9
-20 to -15	5a	-28.9 to -26.1
-15 to -10	5b	-26.1 to -23.3
-10 to -5	6a	-23.3 to -20.6
-5 to 0	6b	-20.6 to -17.8
0 to 5	7a	-17.8 to -15





# Peach Replant Syndrome

**Non-replant site**



**Replant site**

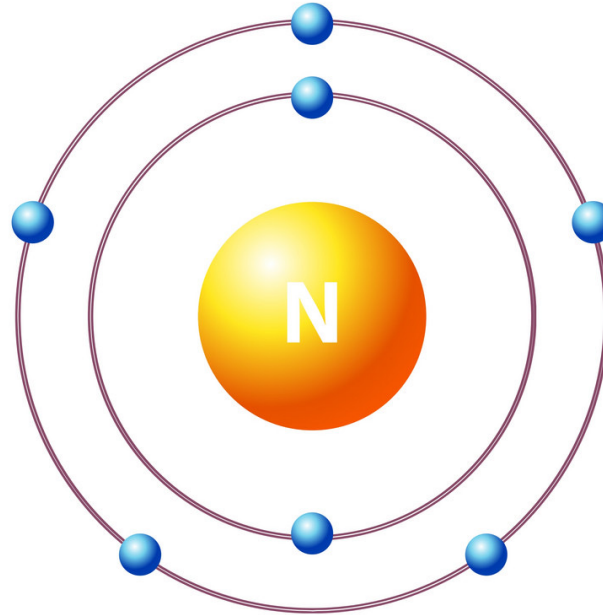




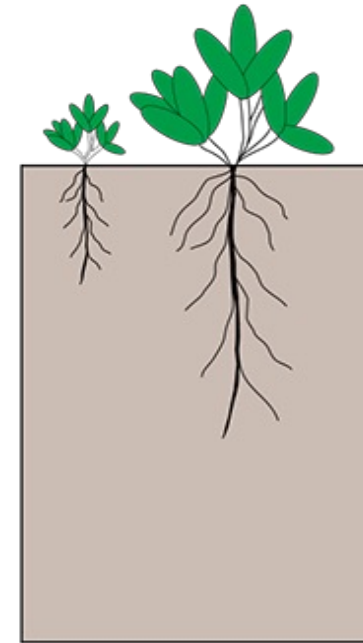
**Monocropping:**  
Orchard: 15 years



**Abiotic Factors:**  
Soil Nutrients



**Intraspecific Allelopathy:**  
Autotoxicity





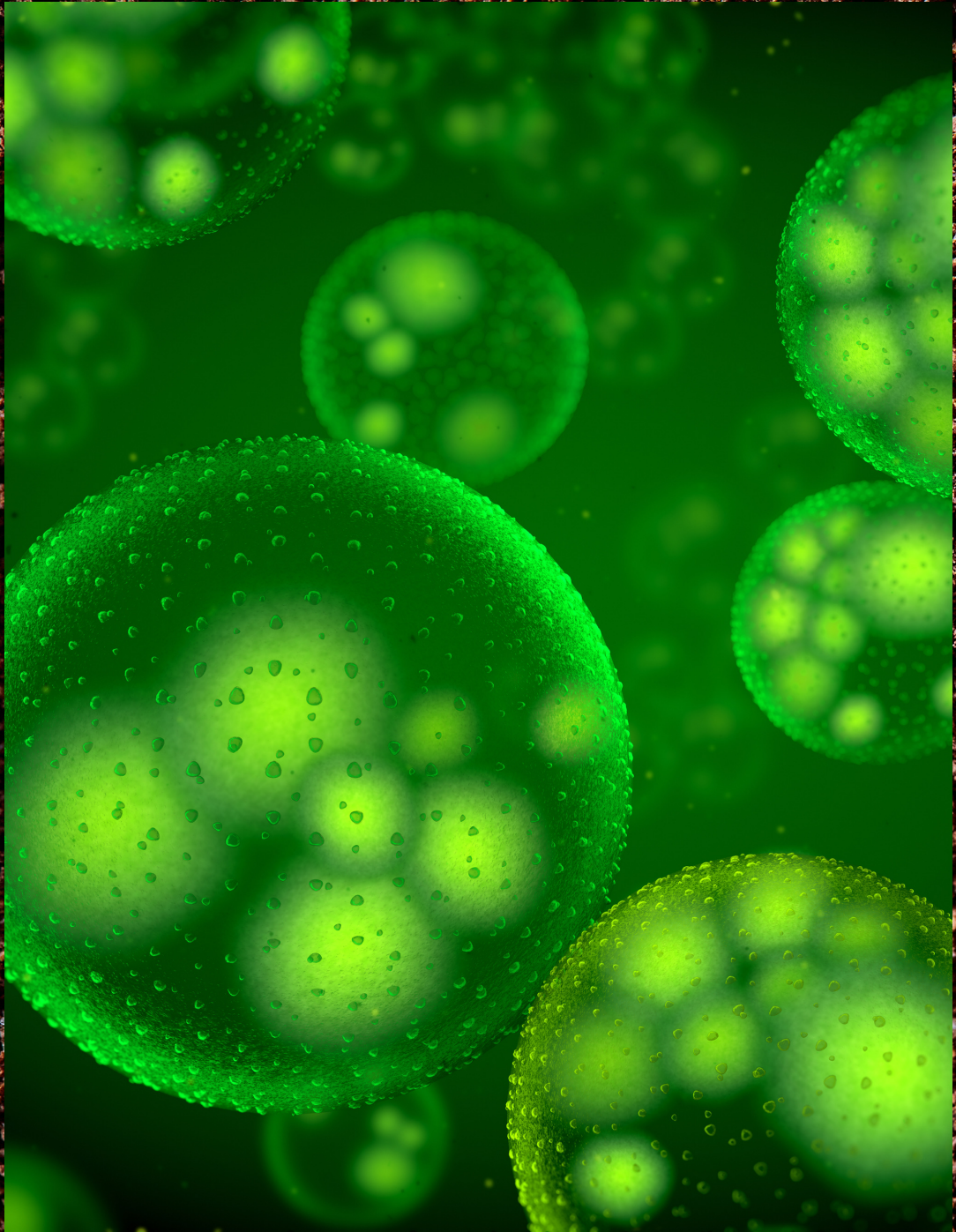




A close-up photograph of dark brown soil, showing a granular and crumbly texture. The soil is the background for the text.

1g of soil =  
10 billion microbes









PGPR

vs



Pathogen









# Our Study

Issue

Remedy

**Monocropping**

Genetically different cover crops

**Nutrient depletion**

Cover Crop reincorporation

**Pathogen buildup**

We hypothesize that these sustainable practices could beneficially alter the microbiome of replant soils and empower peaches to recruit beneficial bacteria



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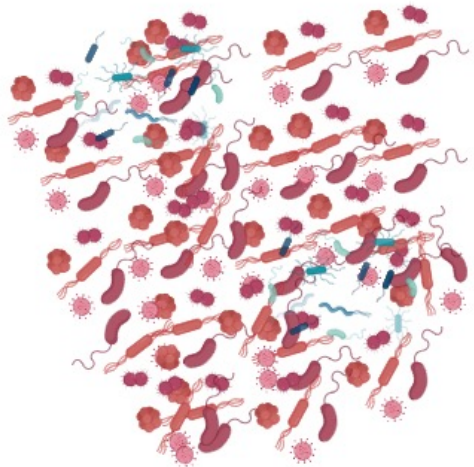
**Soil disinfection to lower microbial load**

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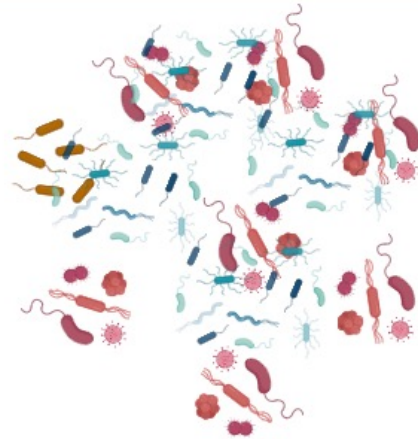


# Conceptual Replant Syndrome Solution

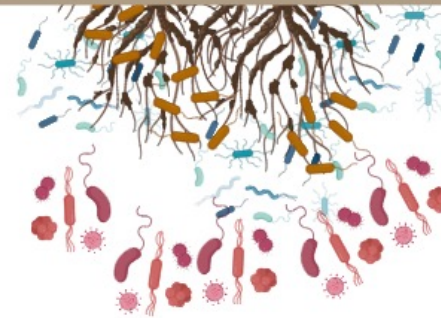
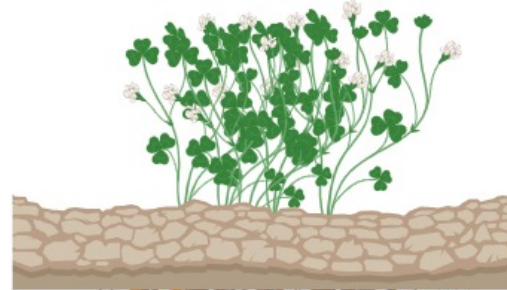
**Replant Soil**



**Step 1: Disinfect Soil**



**Step 2: Cover Crop**



**Step 3: Plant Tree**







# 4 Cover Crops in Peach Orchard Soil









Aaron Stock # 45053320

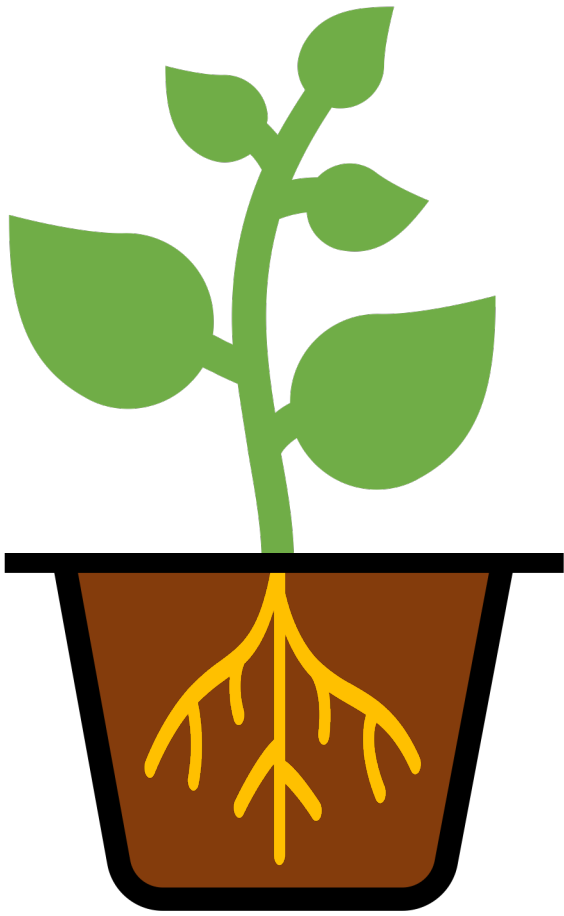




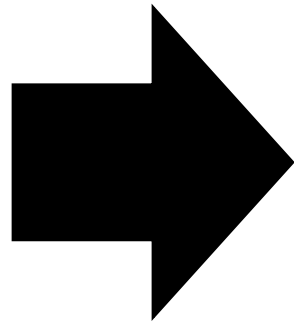
Aaron Stock # 45053320



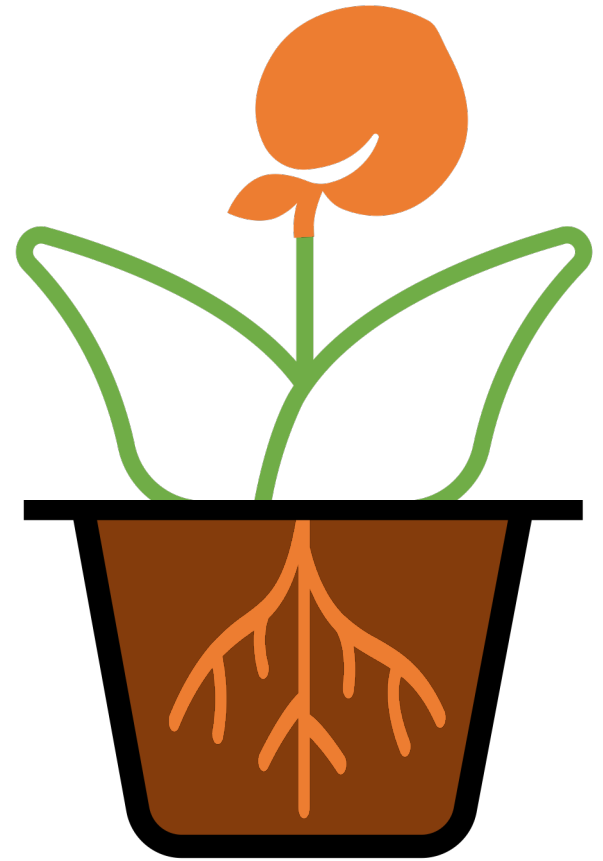
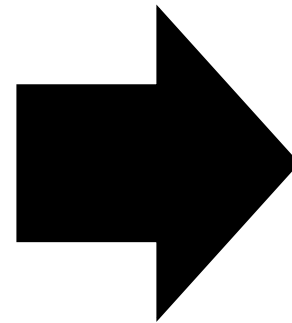




12 Weeks



2 Weeks



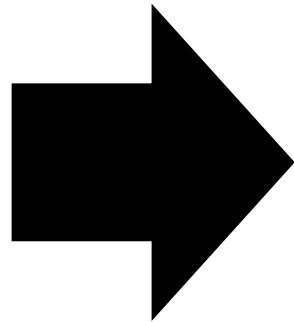
22 Weeks



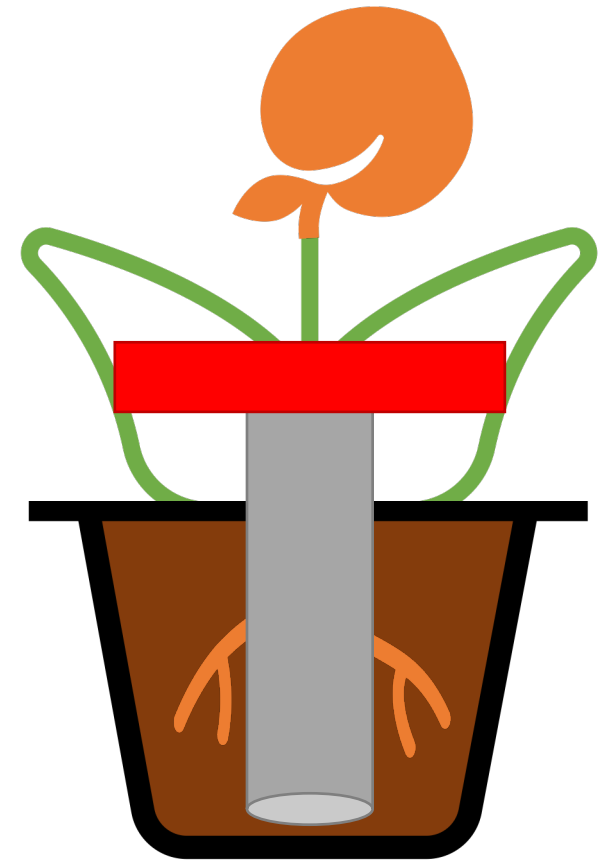
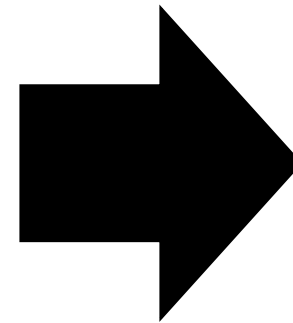
Bulk Soil  
3 Time Points



Cover Crop

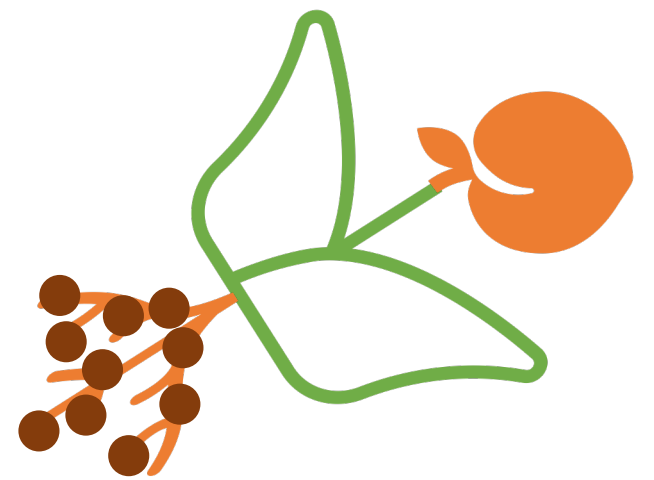
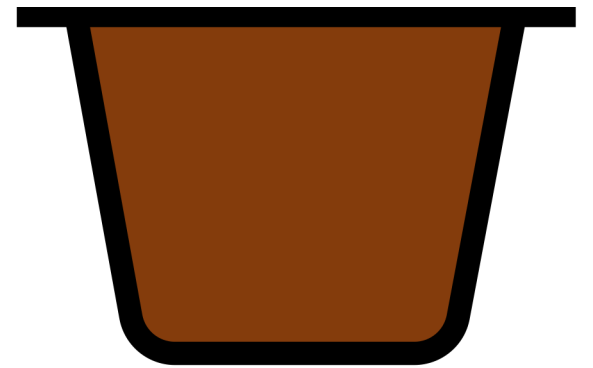
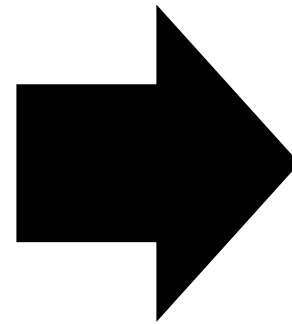
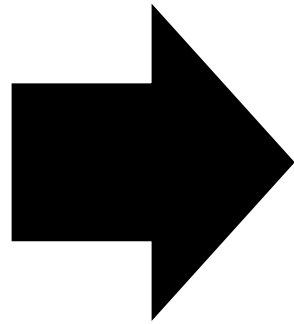
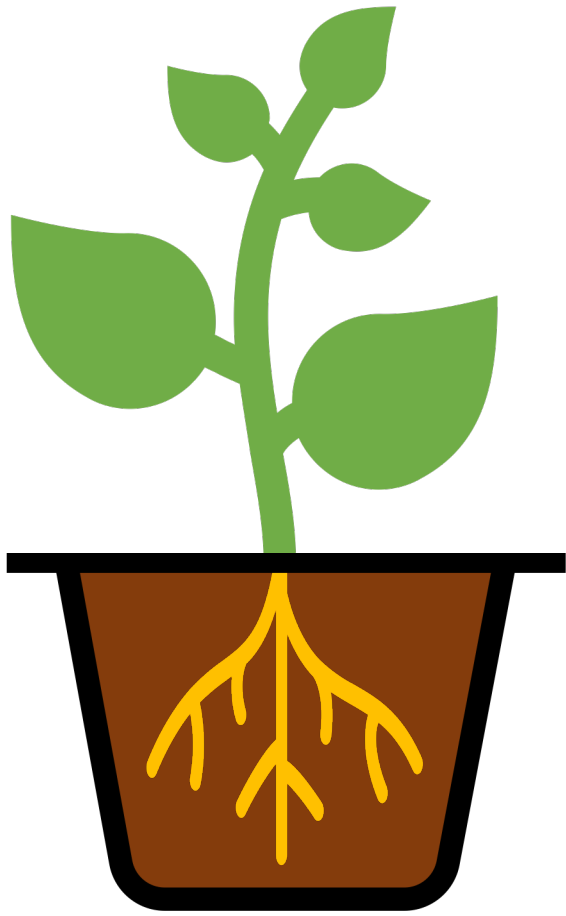


Incorporated



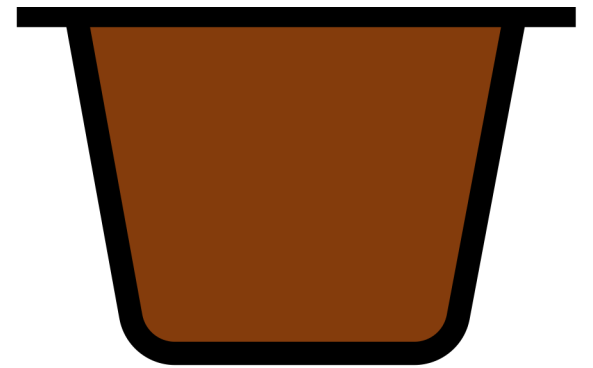
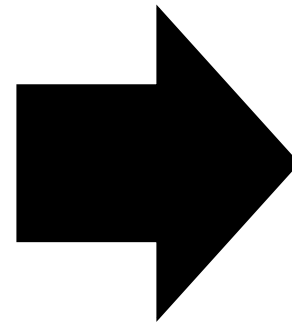
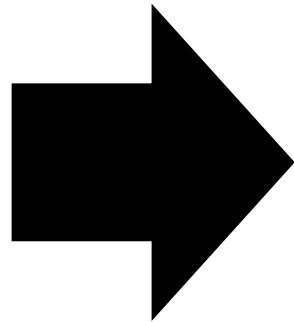
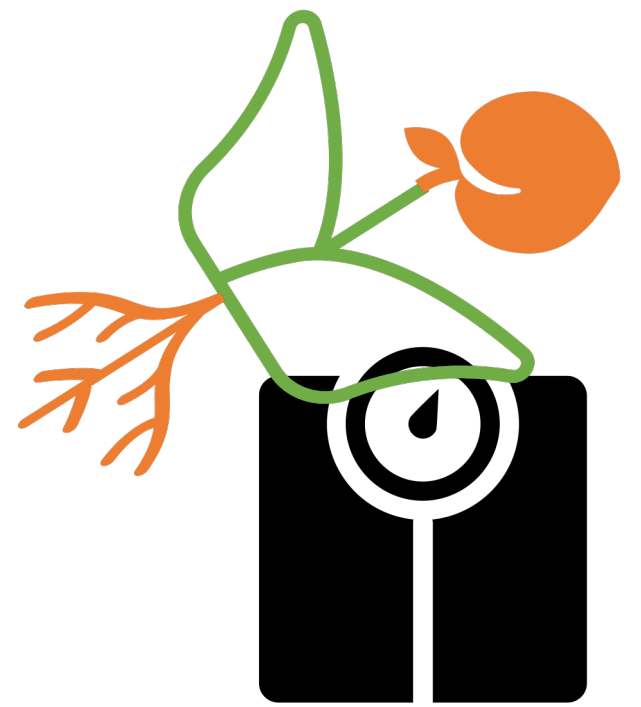
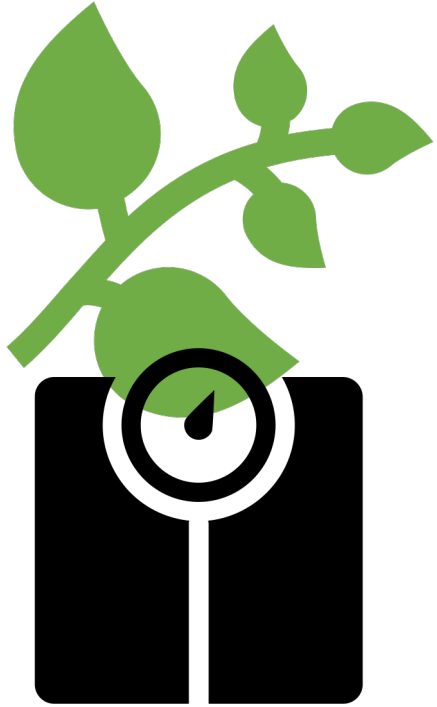
Peach





Rhizosphere

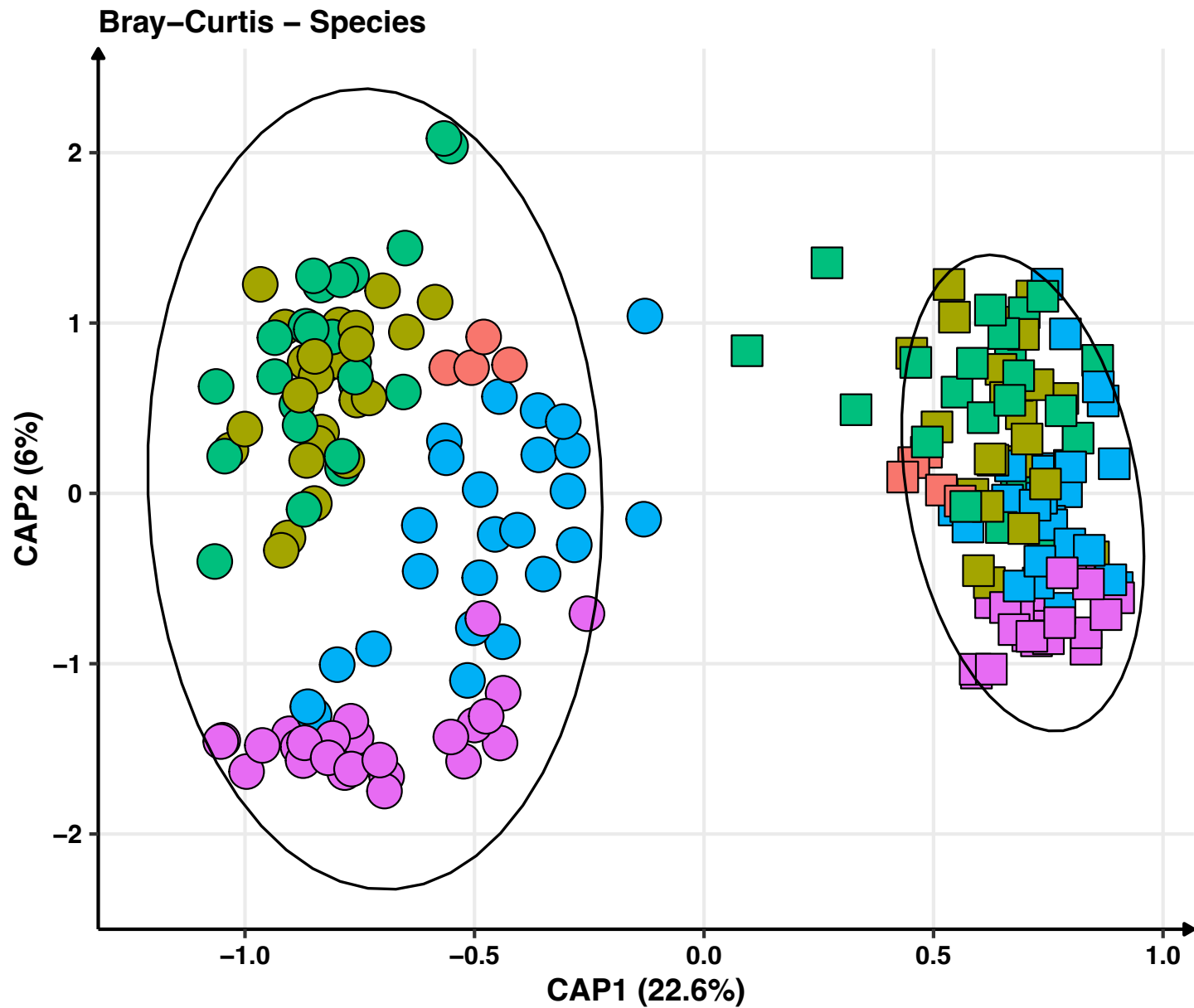




12 Weeks

22 Weeks





**Soil Treatment**

○ Autoclaved

□ Non-Autoclaved

**Crop Phase**

● Bulk Soil Control

● Rotation Crop

● Rotation Crop Incorporated

● Peach

● Peach Rhizosphere



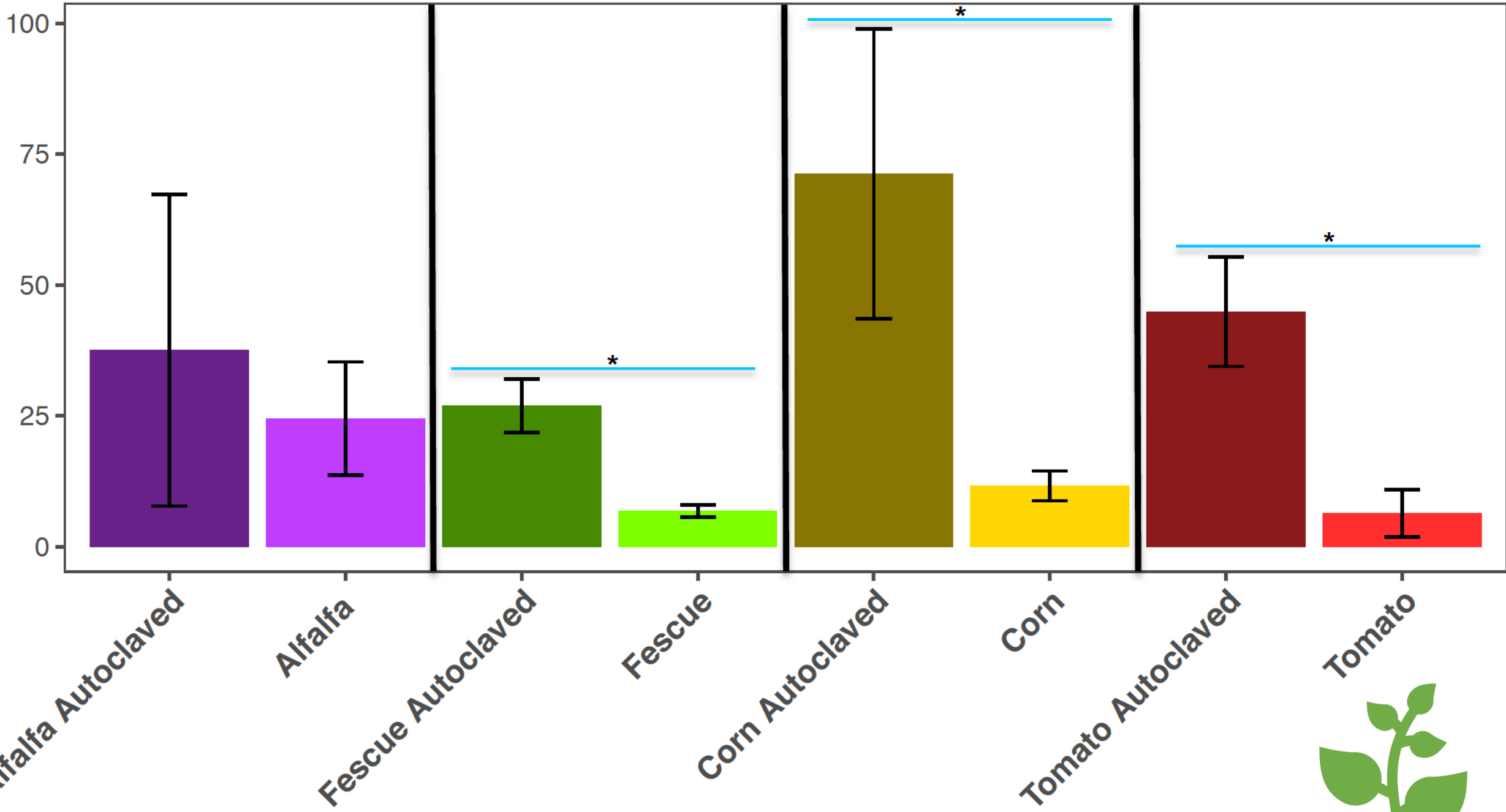
Non-autoclaved



Autoclaved

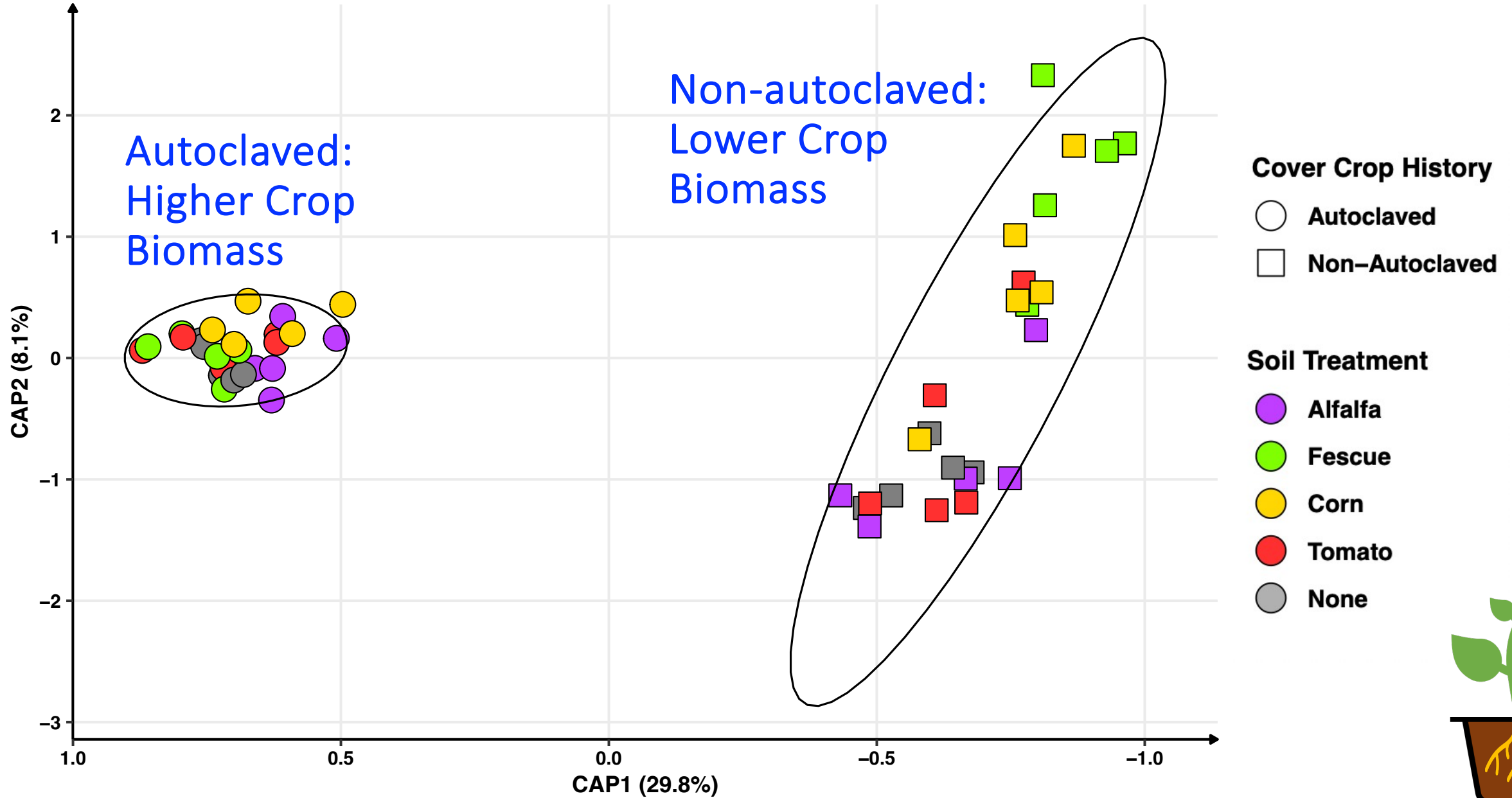


# Mean Fresh Above Biomass (g)



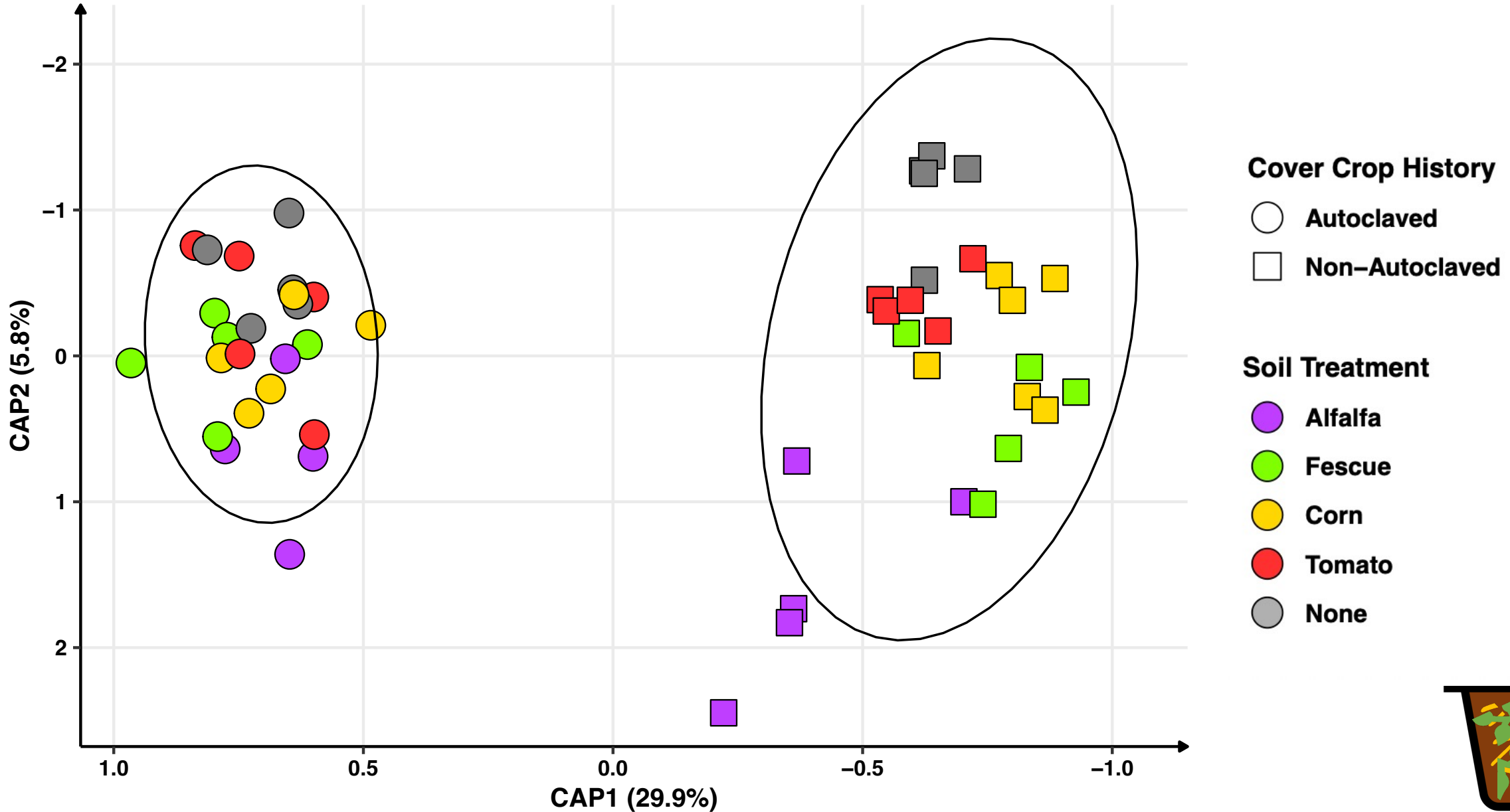


# Cover crop bulk soil microbiomes





# Cover crop incorporated bulk soil microbiomes





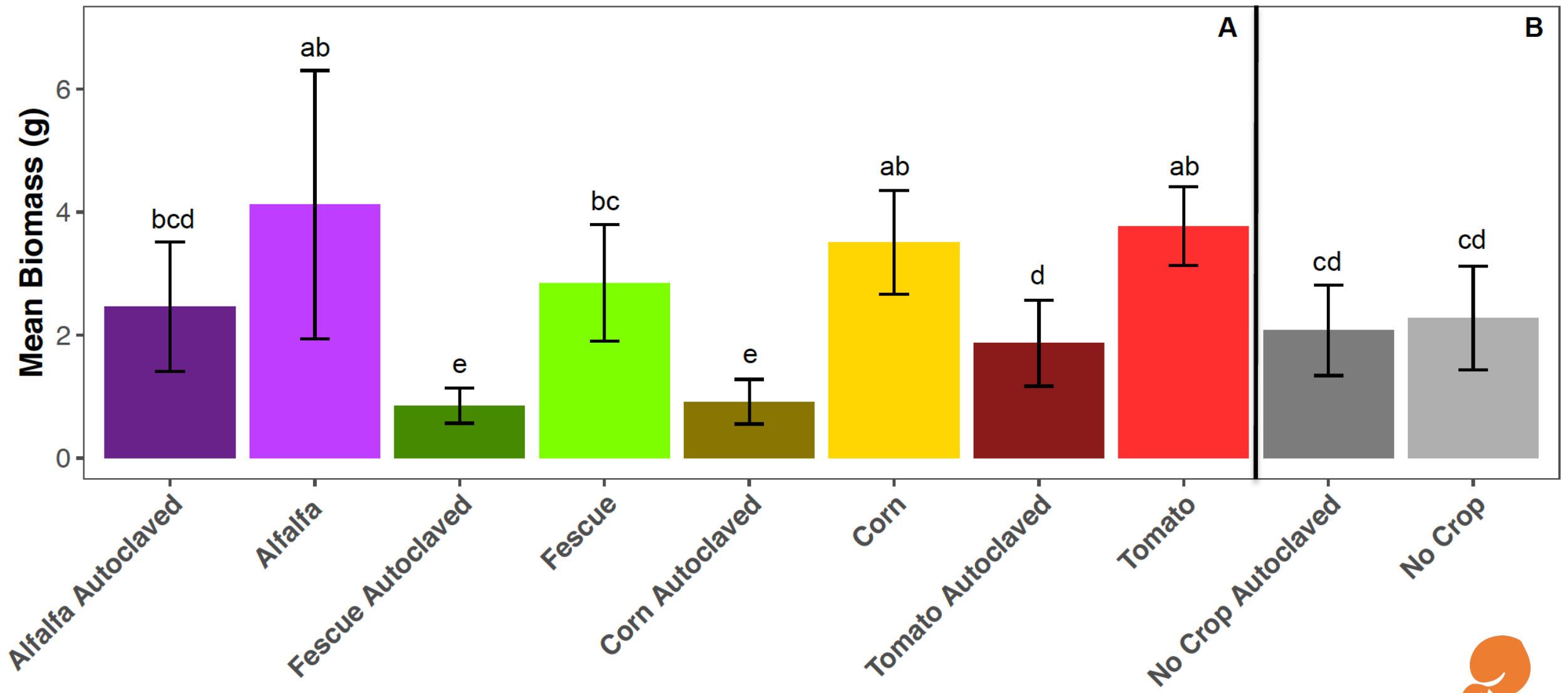
Autoclaved and Alfalfa Cover  
Crop



Non-Autoclaved and Alfalfa Cover  
Crop





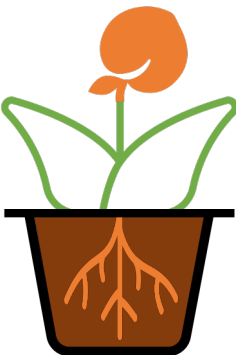
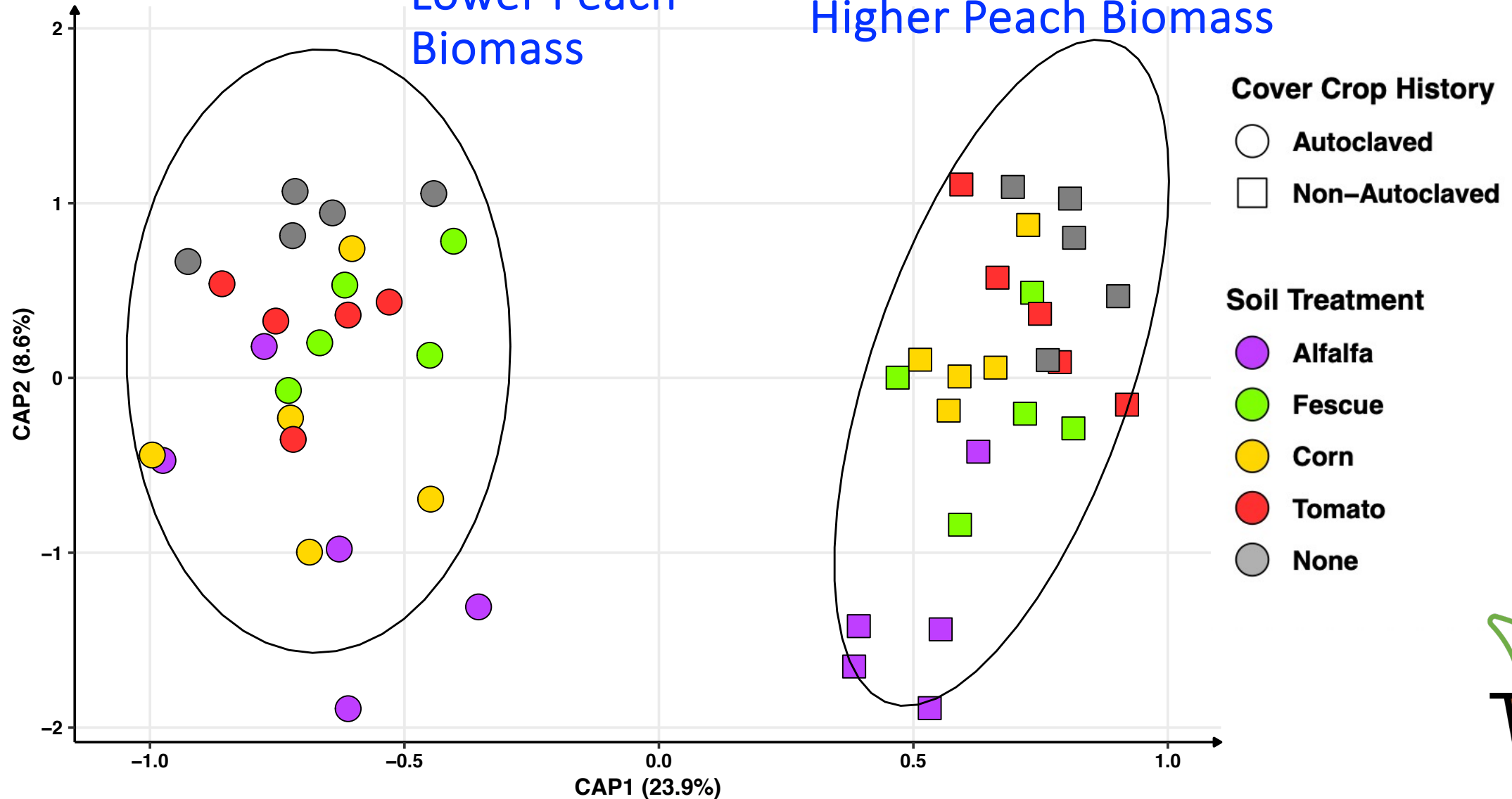




# Peach bulk soil bacterial microbiomes (22 weeks)

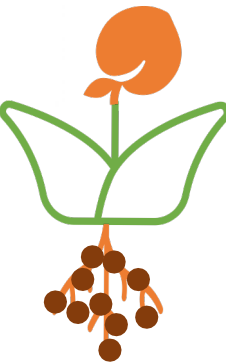
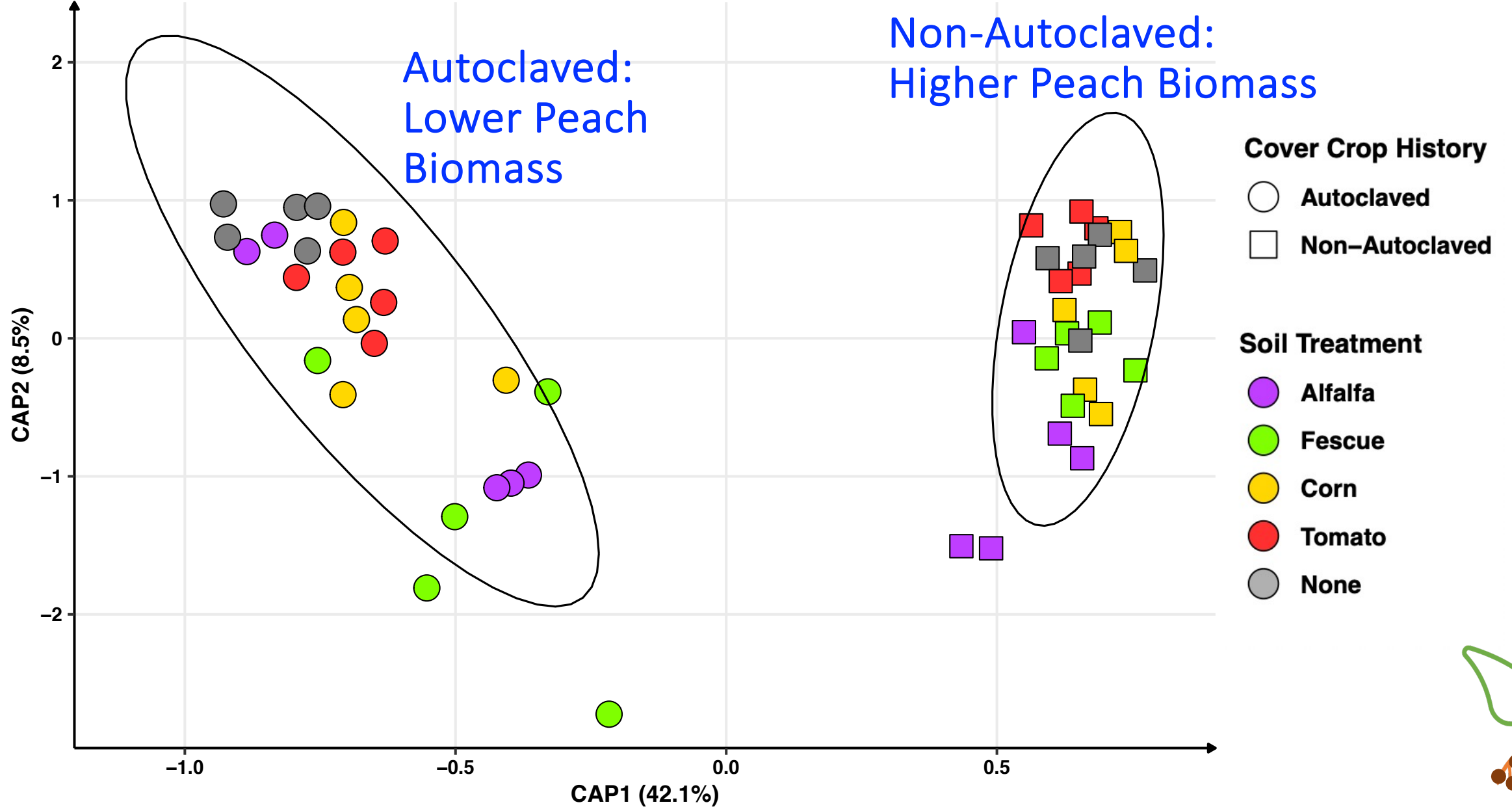
Autoclaved:  
Lower Peach  
Biomass

Non-Autoclaved:  
Higher Peach  
Biomass





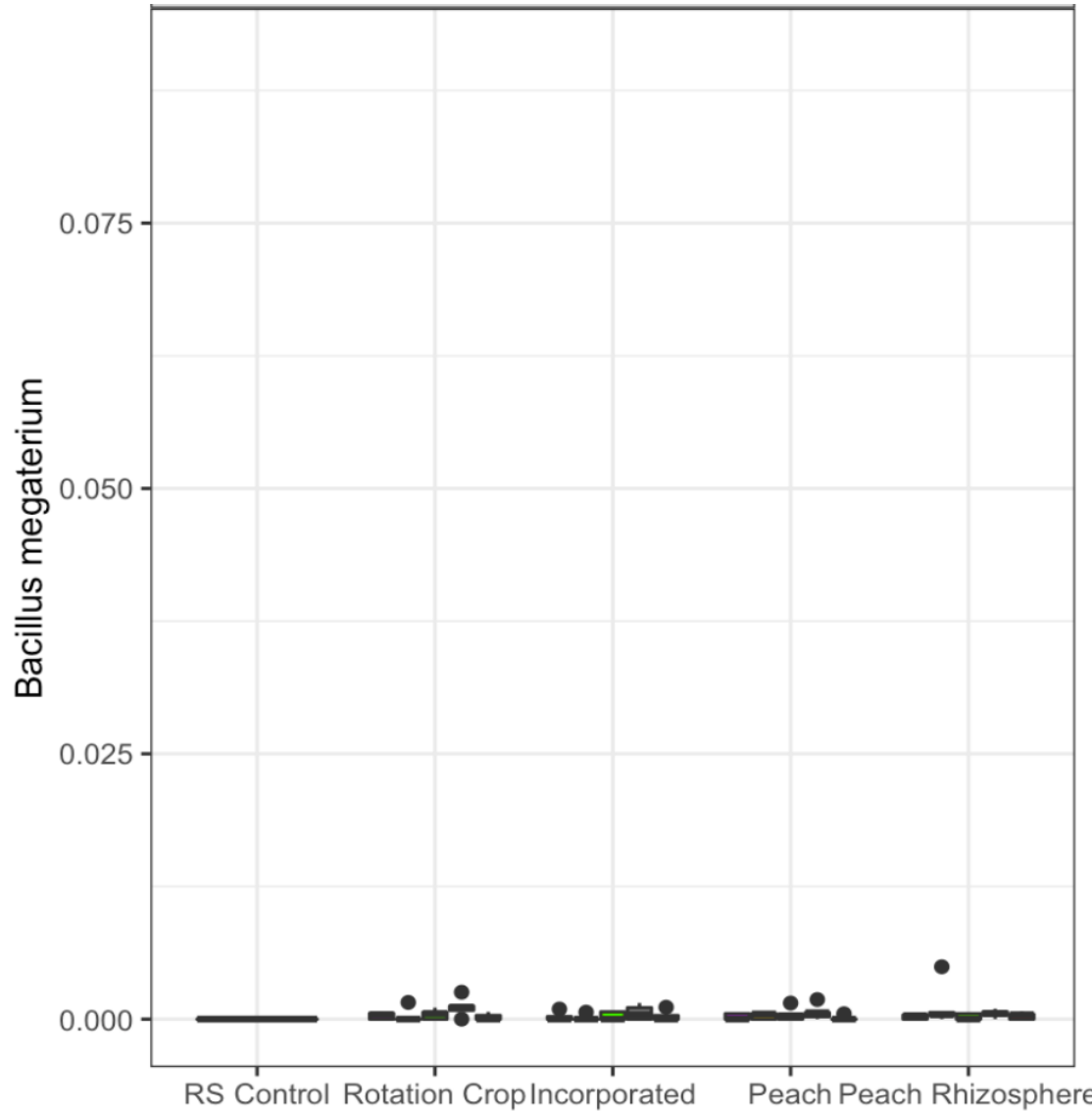
# Peach rhizosphere bacterial microbiomes (22 weeks)



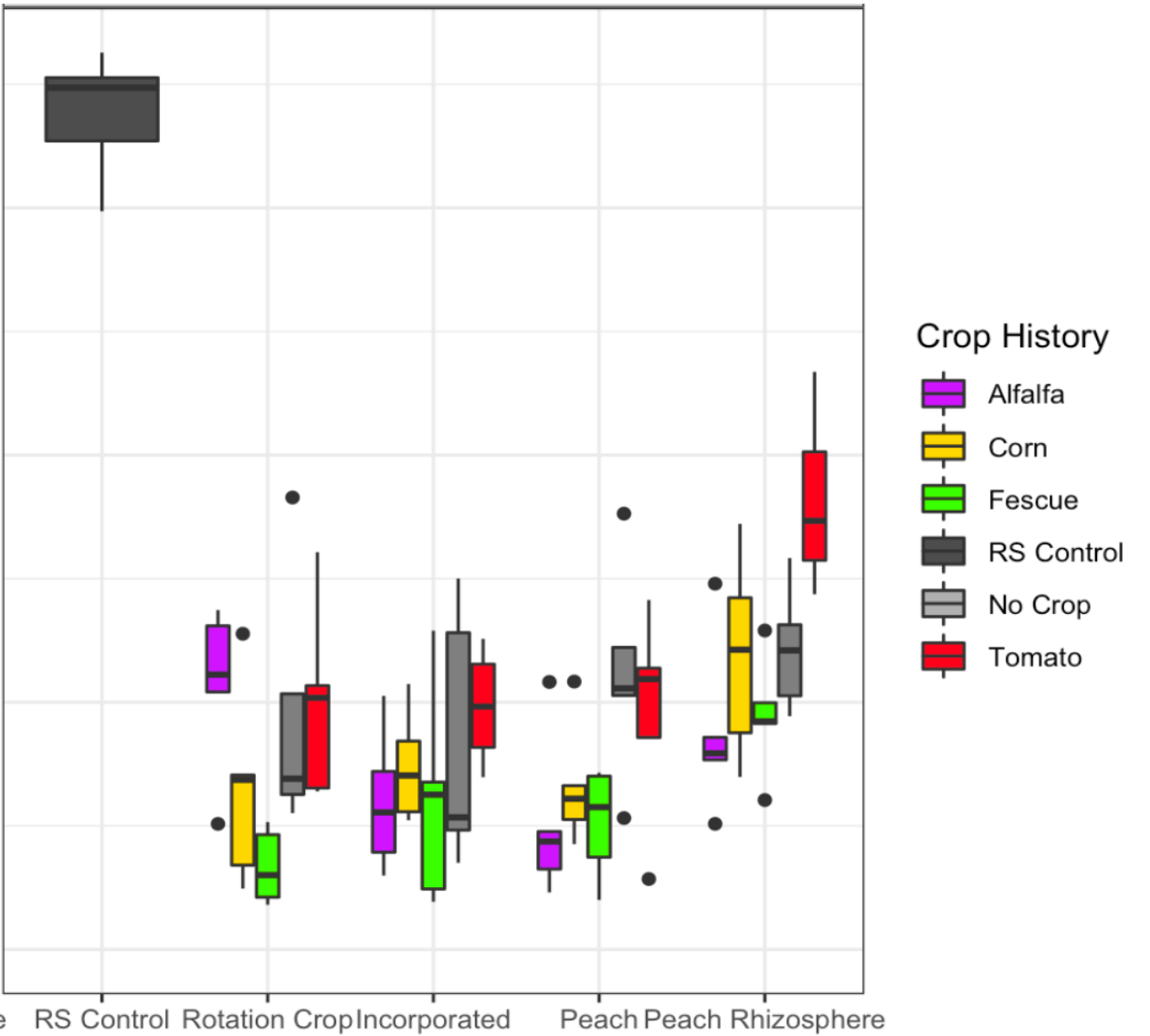


# *Bacillus megaterium*

## Autoclaved

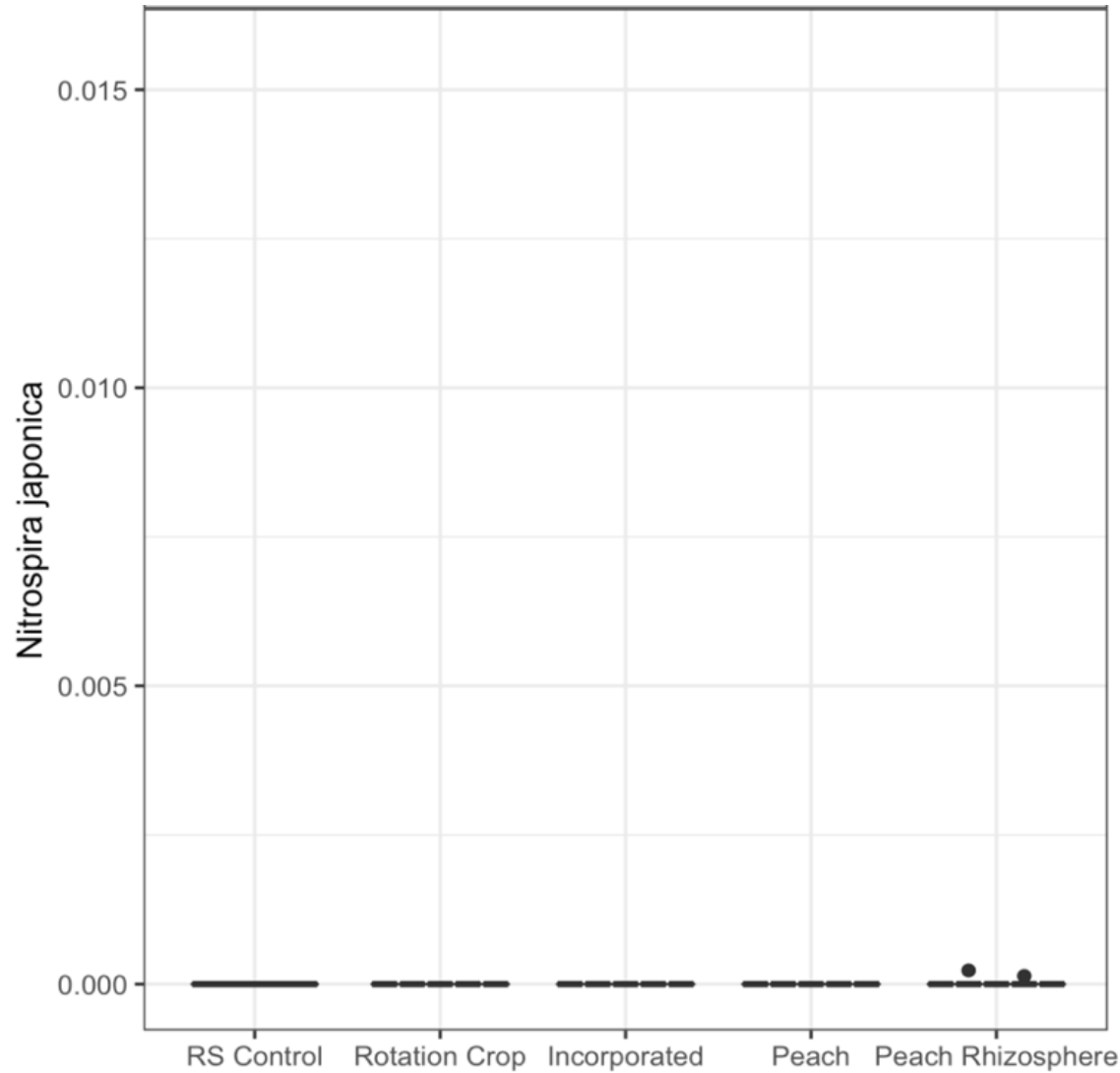


## Non-Autoclaved

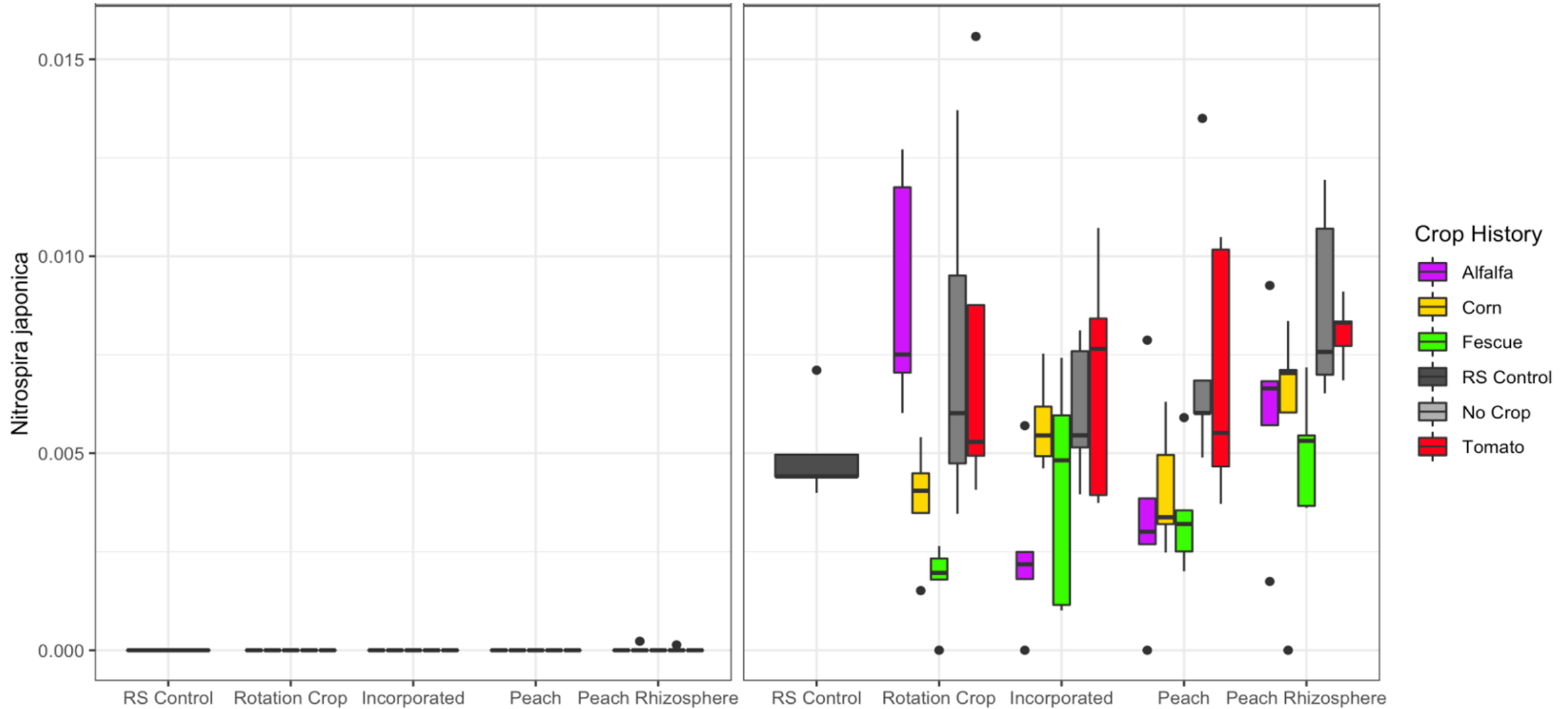


# *Nitrospira japonica*

## Autoclaved



## Non-Autoclaved

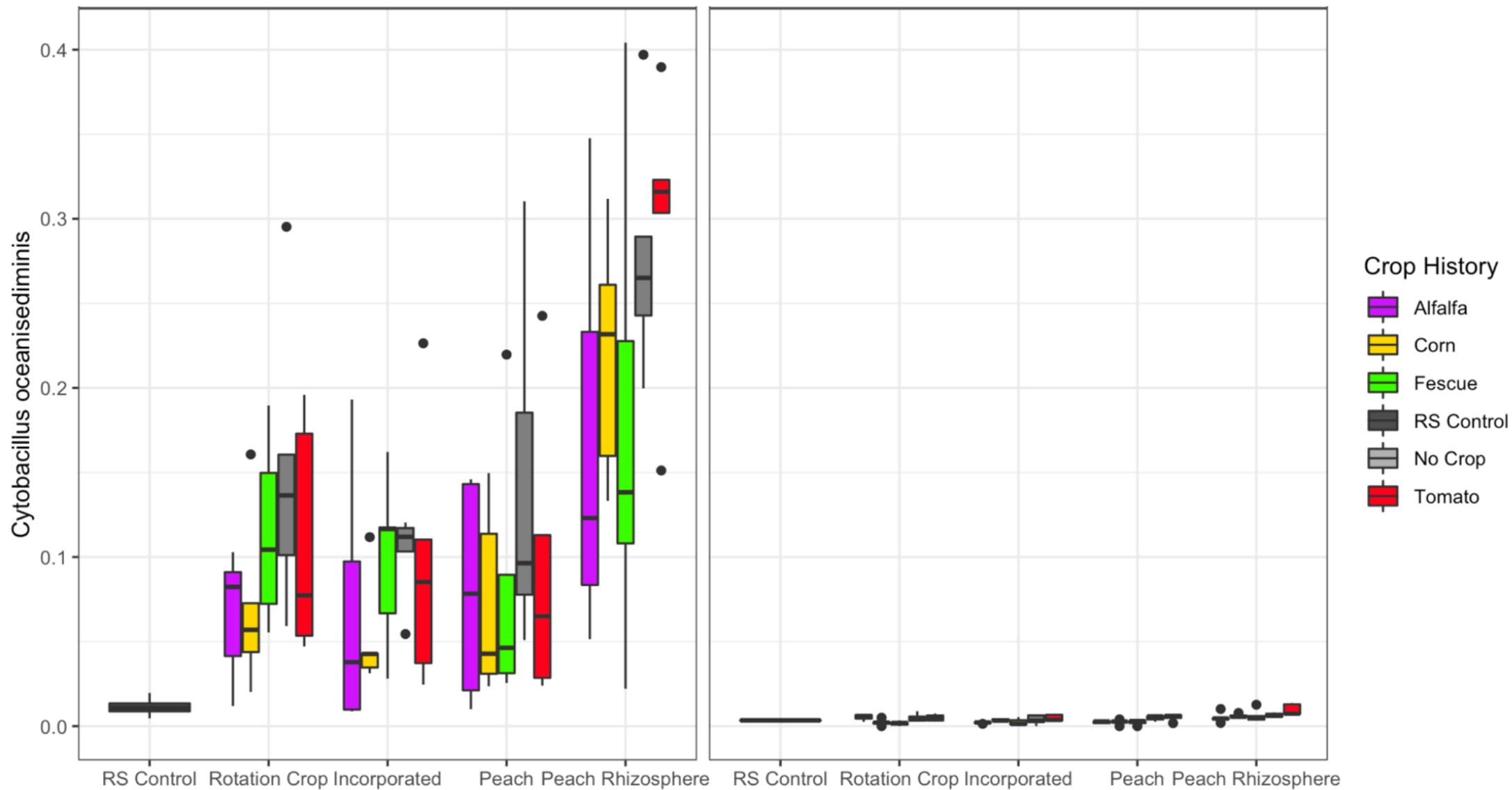




# *Cytobacillus oceanisediminis*

## Autoclaved

## Non-Autoclaved



# Summary

---

**Cover crop biomass was higher in autoclaved soils**

**Peach tree biomass was higher in non-autoclaved soils**

**Non-/autoclaved no plant controls show no biomass difference**

**Autoclaved and non-autoclaved bacterial microbiomes remained separated**

**Using genetically different crops worked- not all CC were equal**

**Beneficial bacteria were lost due to autoclaving soils: moderate disinfection technique**

***Paenibacillus castaneae* and *Bellilinea caldifistulae*: Beneficial bacterial species that were cultivated exclusively in the peach rhizosphere of non-autoclaved soils with a cover crop history**

**Only see part of the picture**



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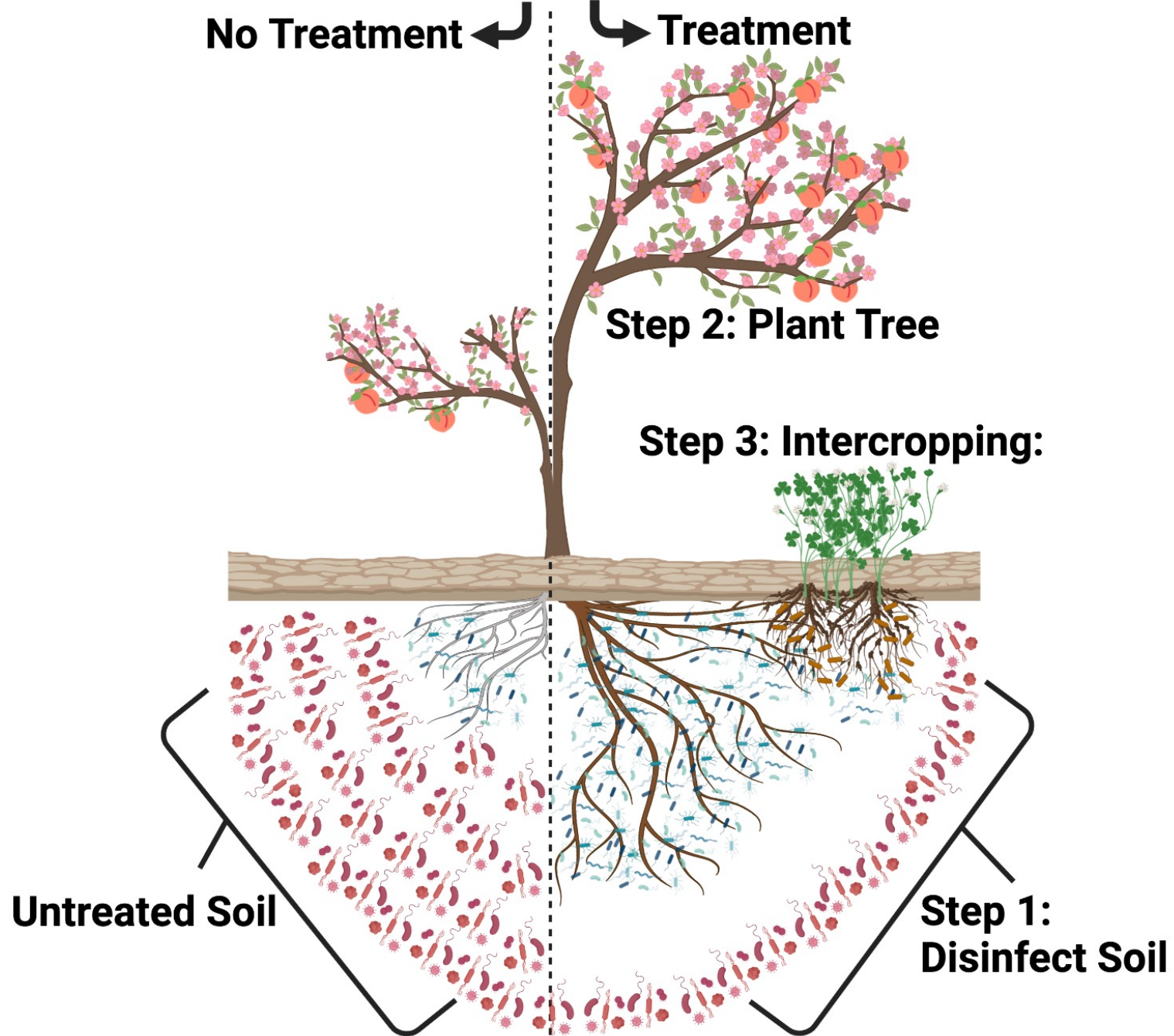
**Only see part of the picture**



# Conceptual Replant Syndrome Solution

No Treatment ←

→ Treatment

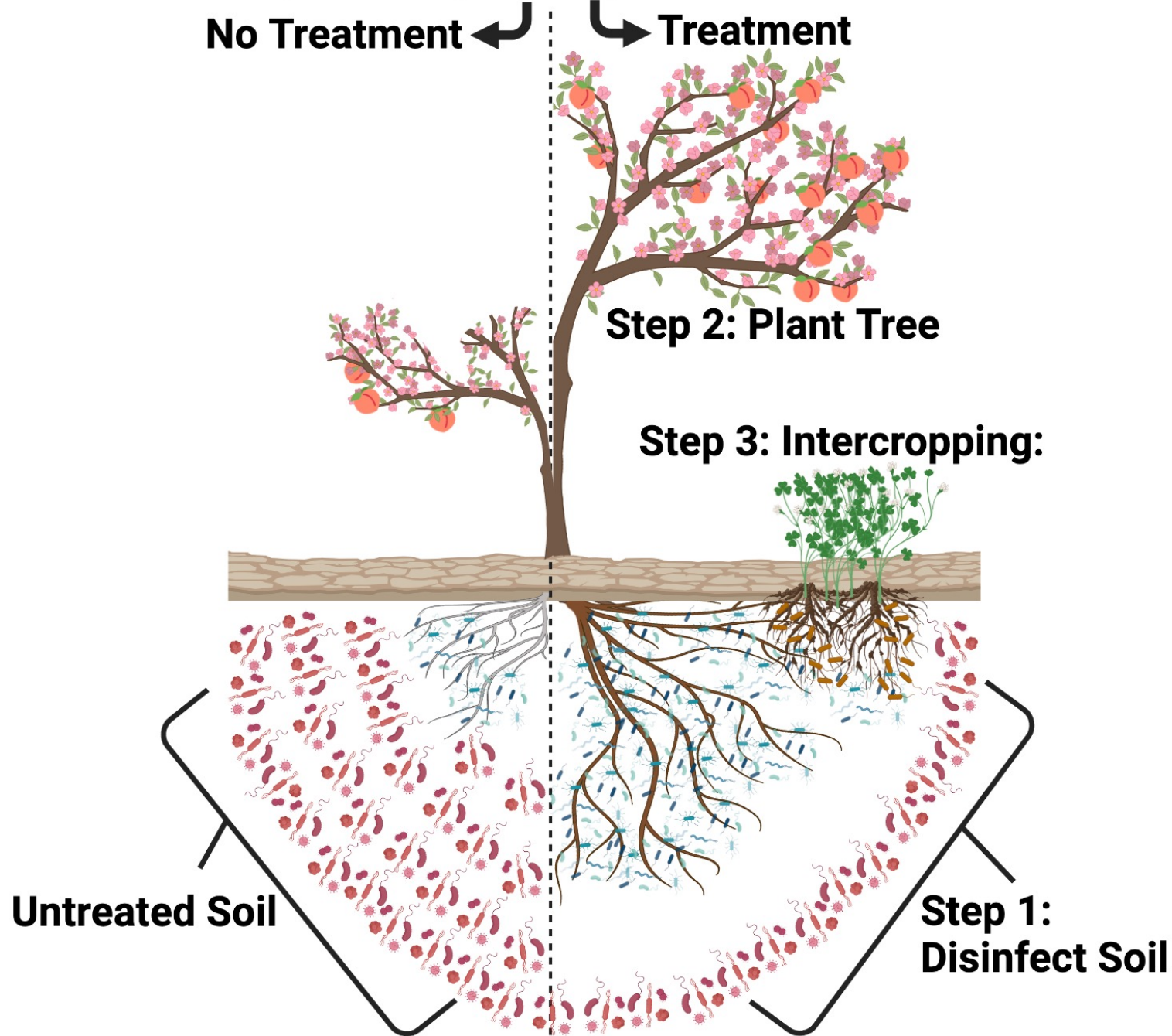




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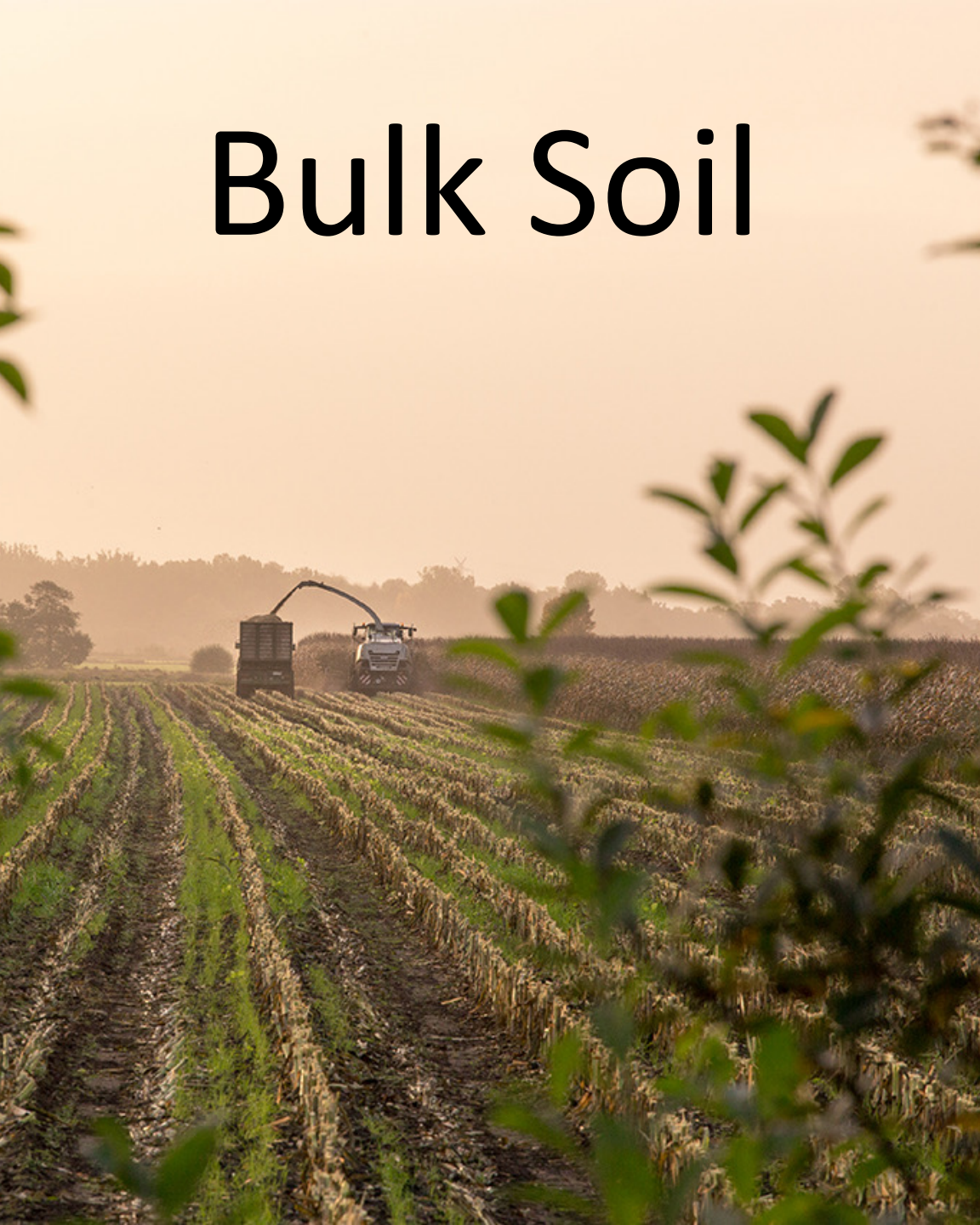
No Treatment ←

→ Treatment





# Bulk Soil



# Bulk Soil









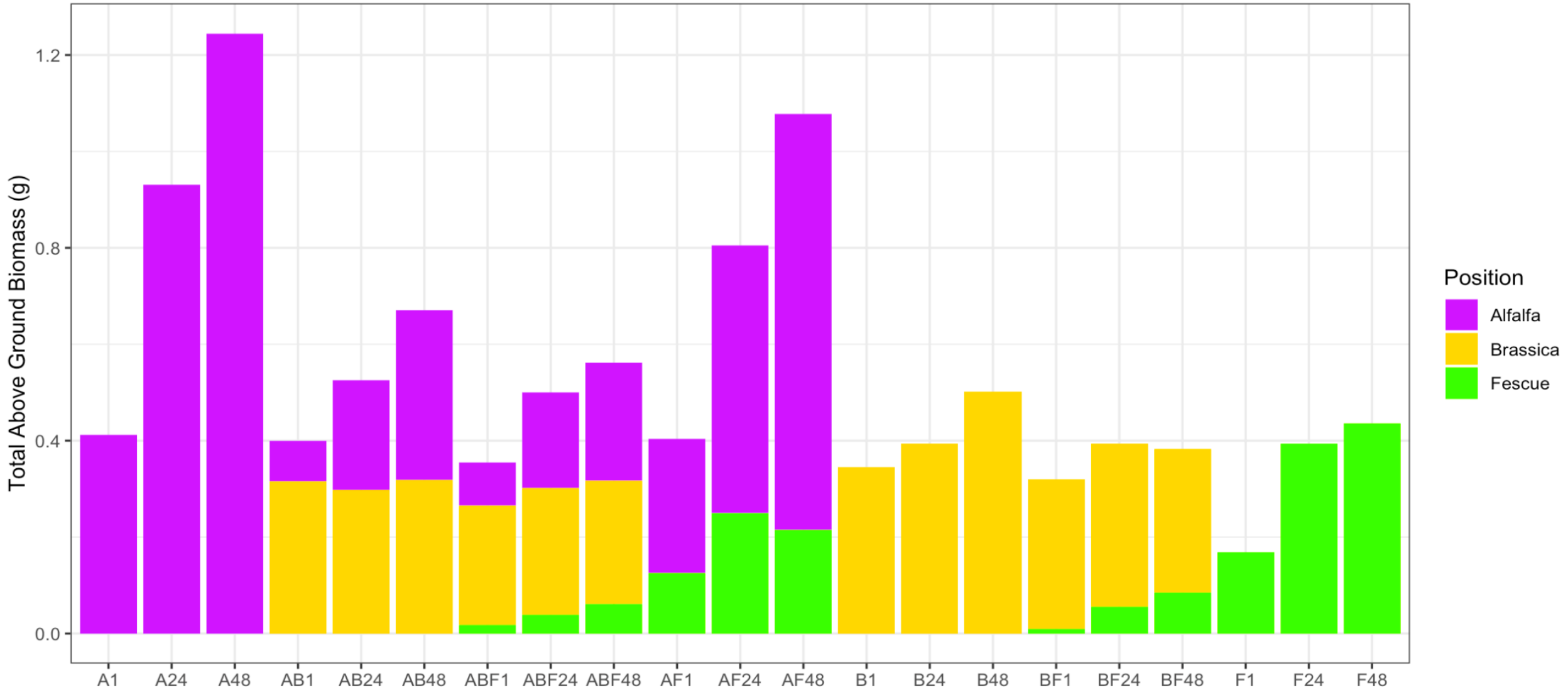
# Microcosm Competition Experiment Set Up

- 7 diversity treatments (1. alfalfa, 2. brassica, 3. fescue, 4. alfalfa-brassica, 5. alfalfa-fescue, 6. brassica-fescue, 7. alfalfa-brassica-fescue)
- 3 density treatments (low: 1-3 plants, medium: 24 plants, and high: 48 plants)
- 21 treatments total

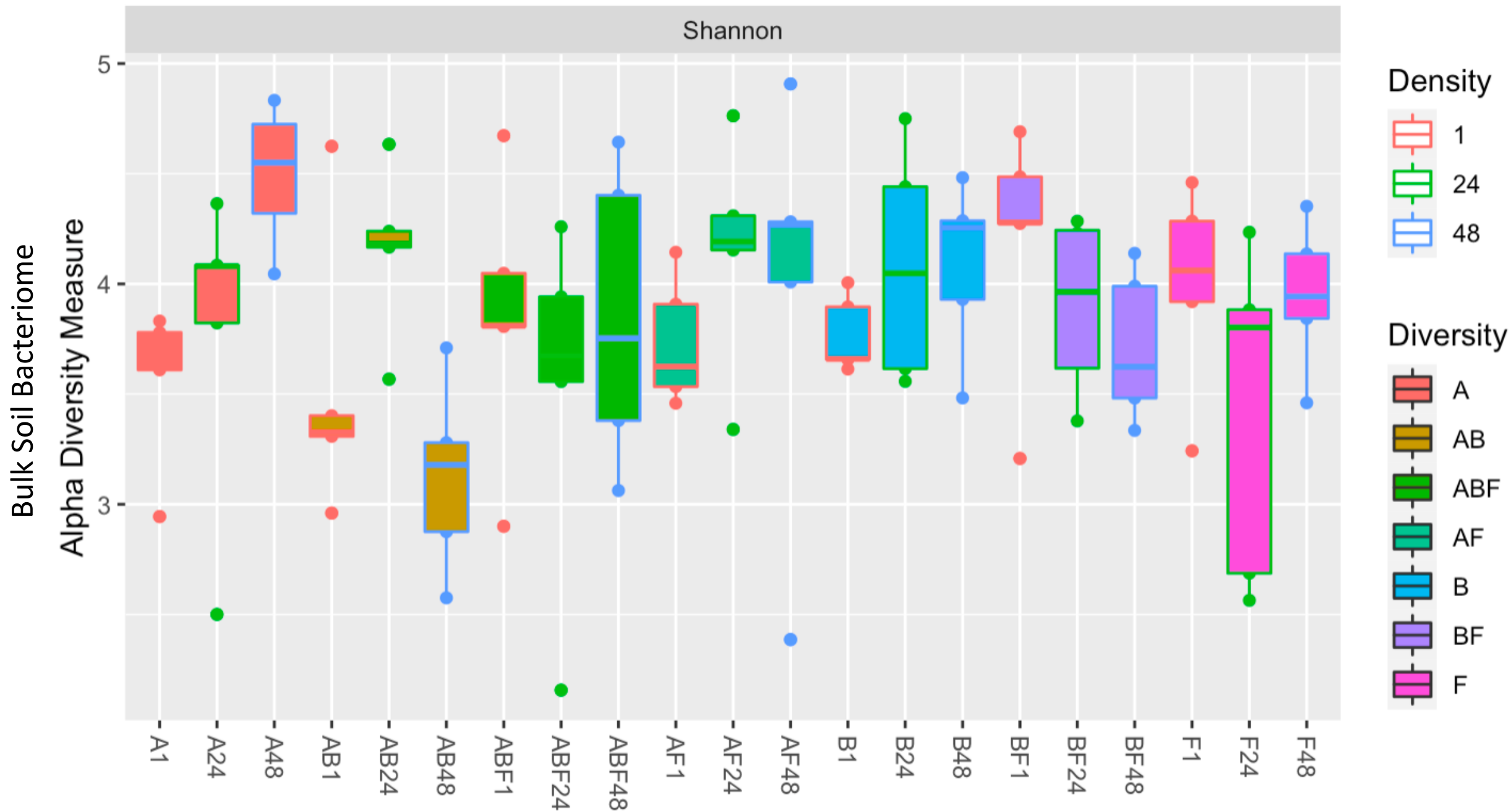
	Monoculture	Polyculture
Low Density (1-3 Plants per microcosm)	1. Alfalfa (1 Plant) 2. Brassica (1 Plant) 3. Fescue (1 Plant)	10. Alfalfa (1 Plant) and Brassica (1 Plant) 11. Alfalfa (1 Plant) and Fescue (1 Plant) 12. Brassica (1 Plant) and Fescue (1 Plant) 13. Alfalfa (1 Plant), Brassica (1 Plant), and Fescue (1 Plant)
Medium Density (24 Plants per microcosm)	4. Alfalfa (24 Plants) 5. Brassica (24 Plants) 6. Fescue (24 Plants)	14. Alfalfa (12 Plants) and Brassica (12 Plants) 15. Alfalfa (12 Plants) and Fescue (12 Plants) 16. Brassica (12 Plants) and Fescue (12 Plants) 17. Alfalfa (8 Plants), Brassica (8 Plants), and Fescue (8 Plants)
High Density (48 Plants per microcosm)	7. Alfalfa (48 Plants) 8. Brassica (48 Plants) 9. Fescue (48 Plants)	18. Alfalfa (24 Plants) and Brassica (24 Plants) 19. Alfalfa (24 Plants) and Fescue (24 Plants) 20. Brassica (24 Plants) and Fescue (24 Plants) 21. Alfalfa (16 Plants), Brassica (16 Plants), and Fescue (16 Plants)



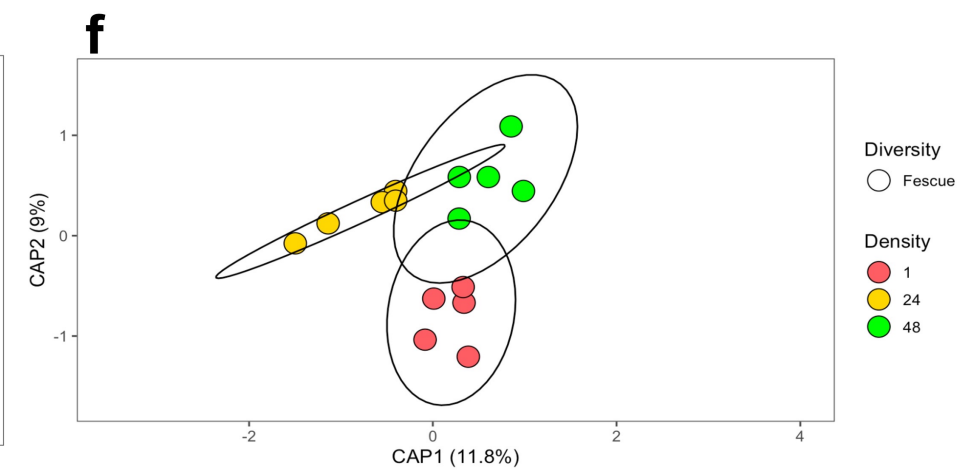
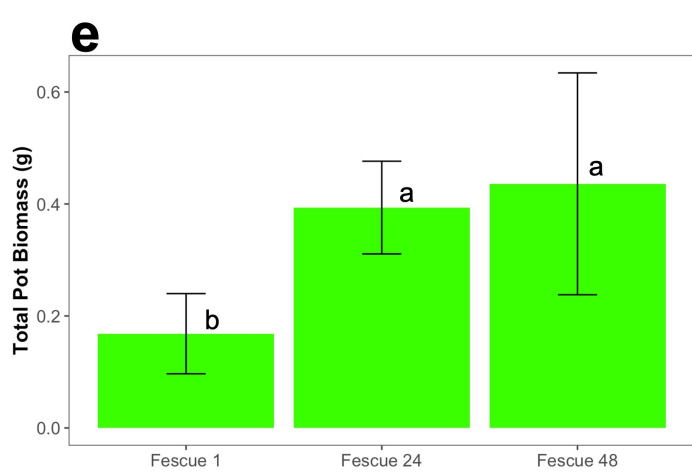
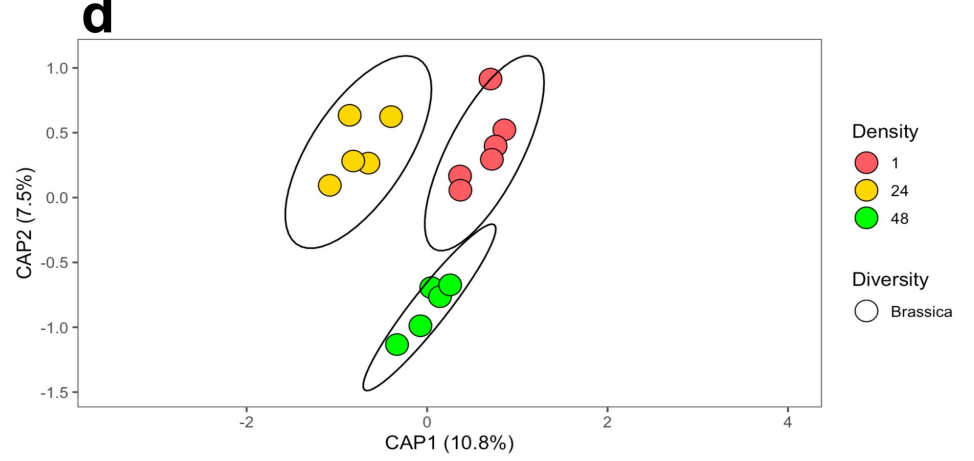
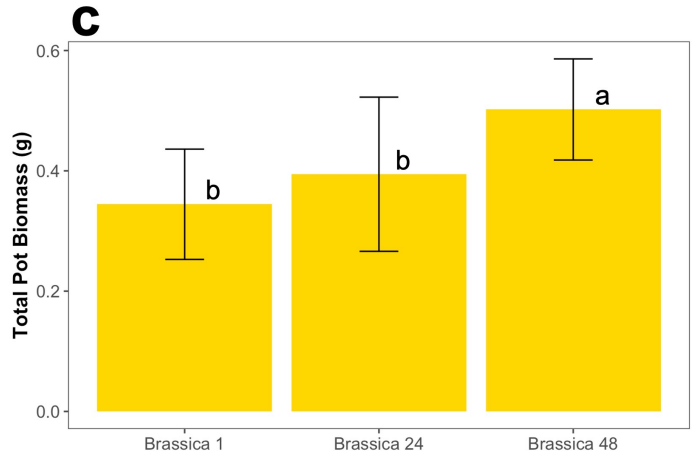
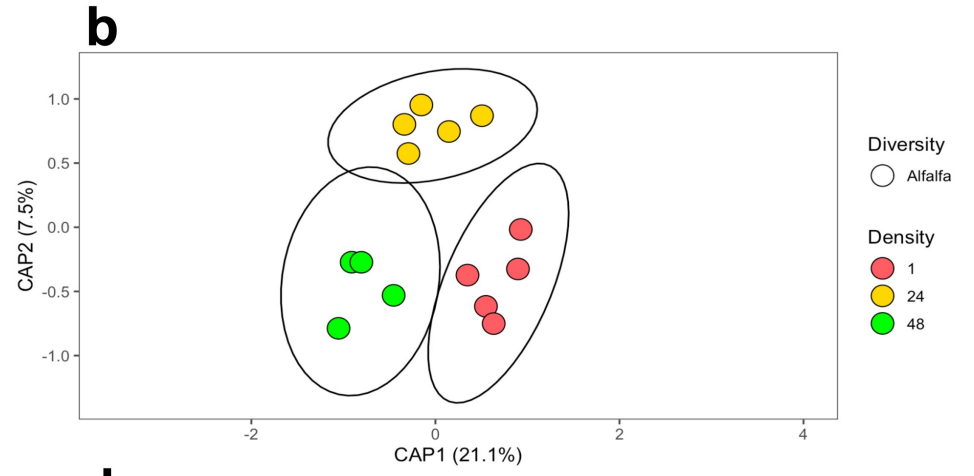
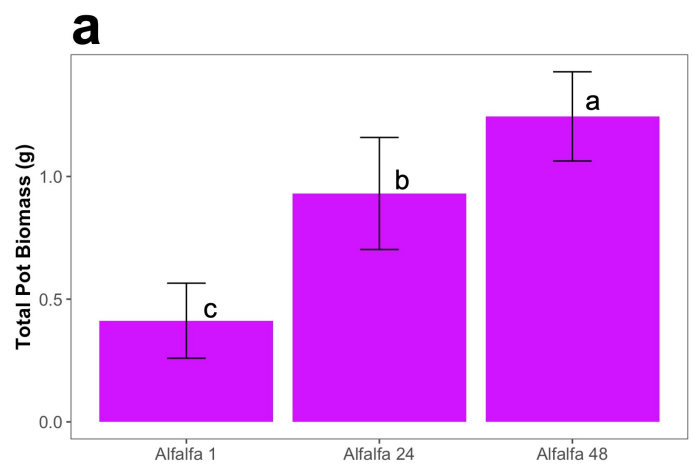
Above Ground Cover Crop Biomass

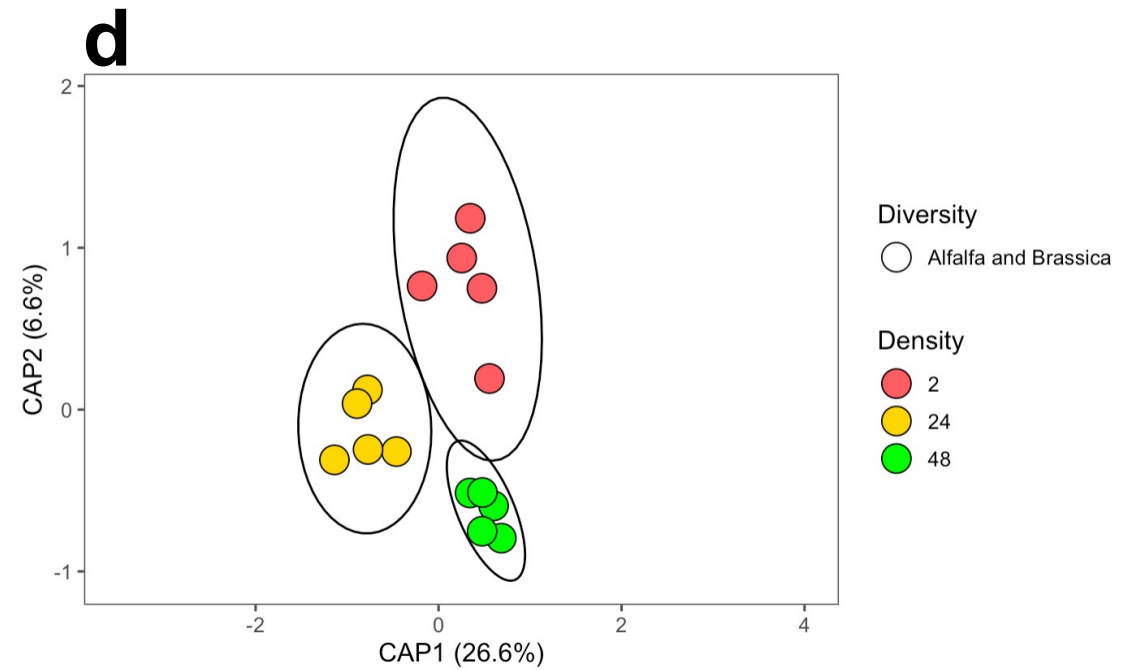
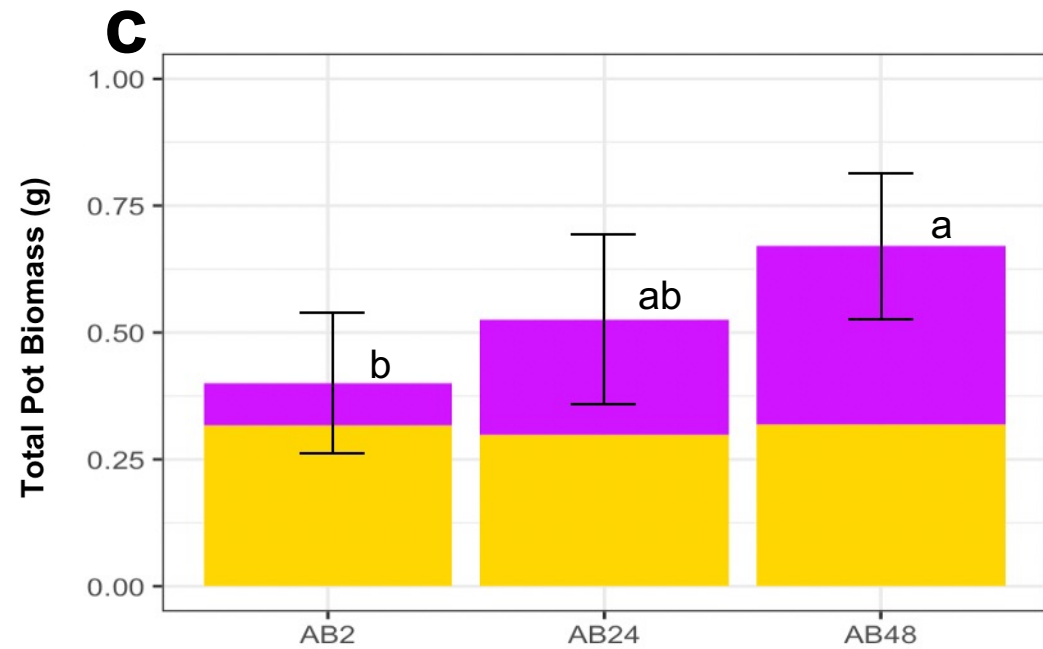
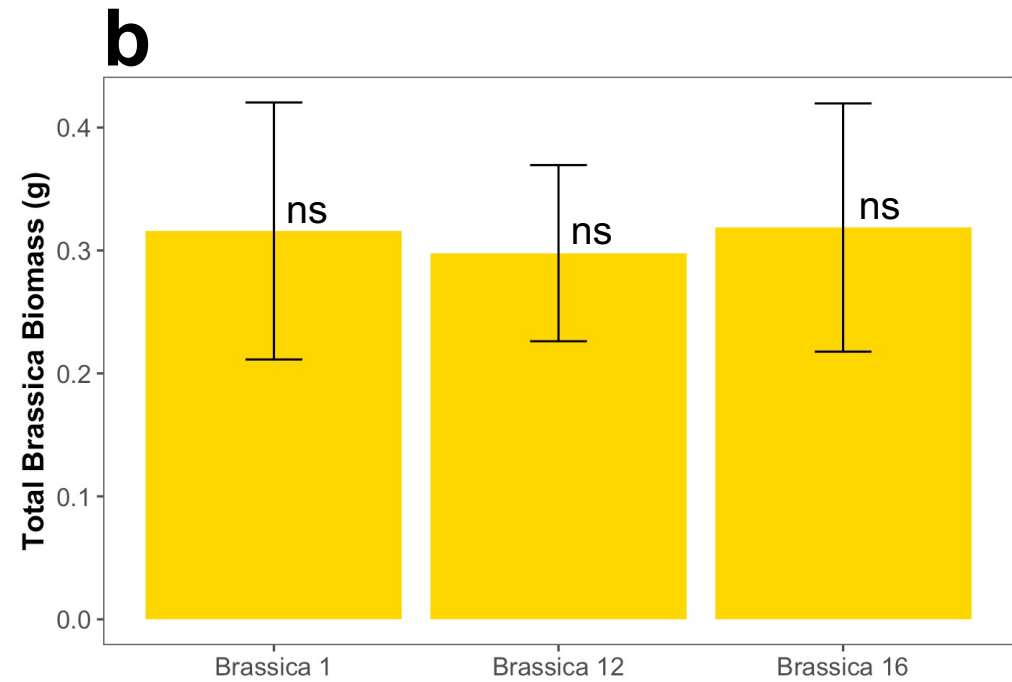
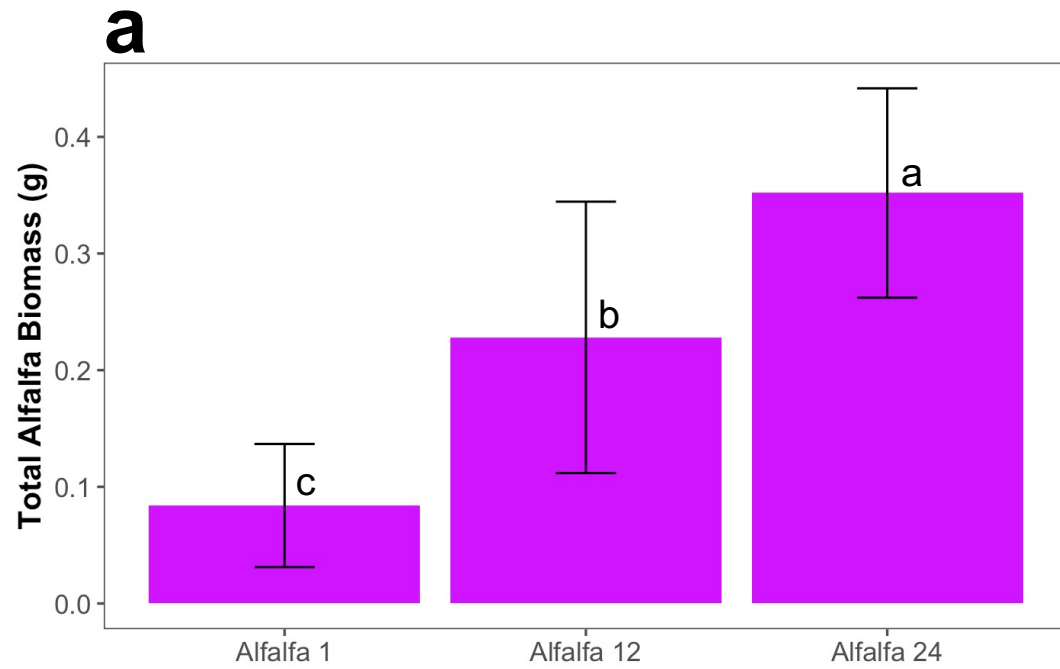


# Shannon Index by Density and Diversity

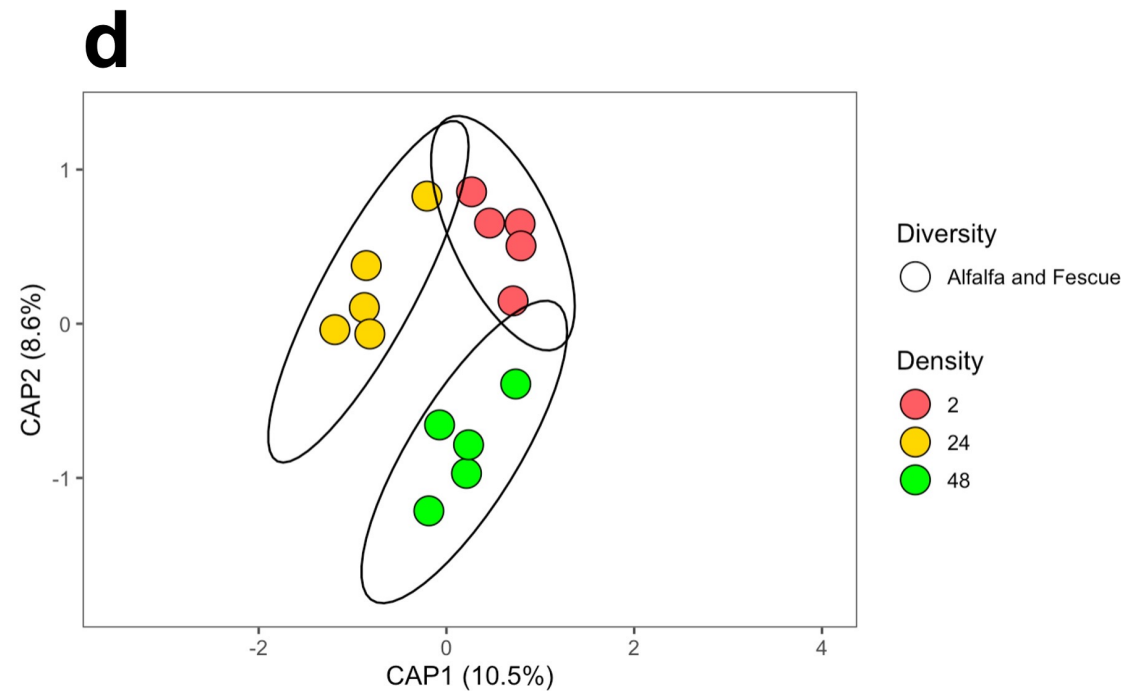
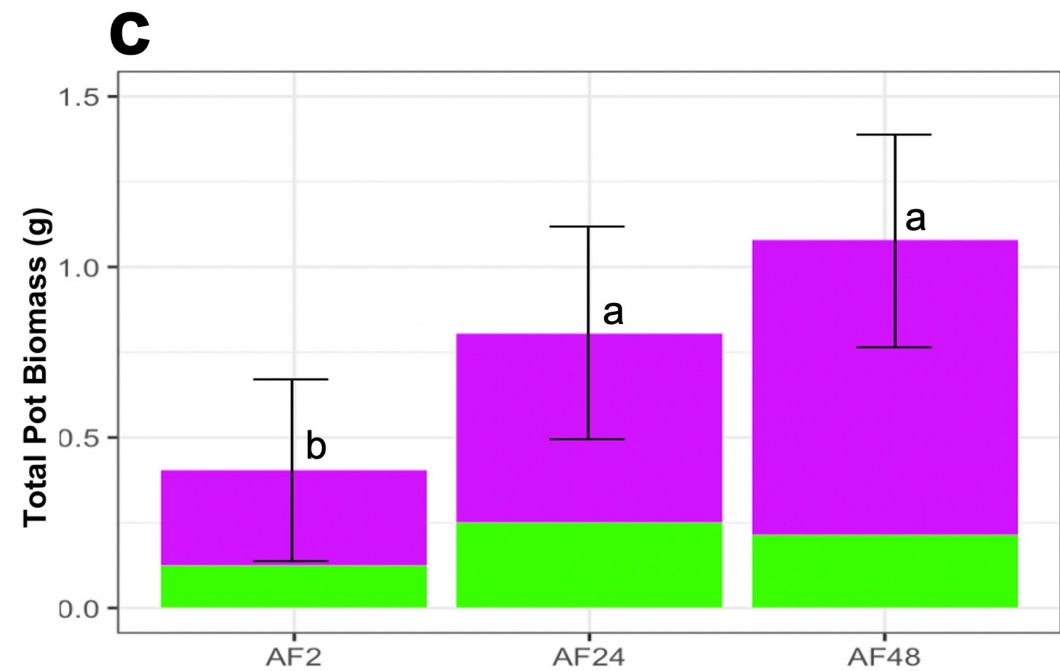
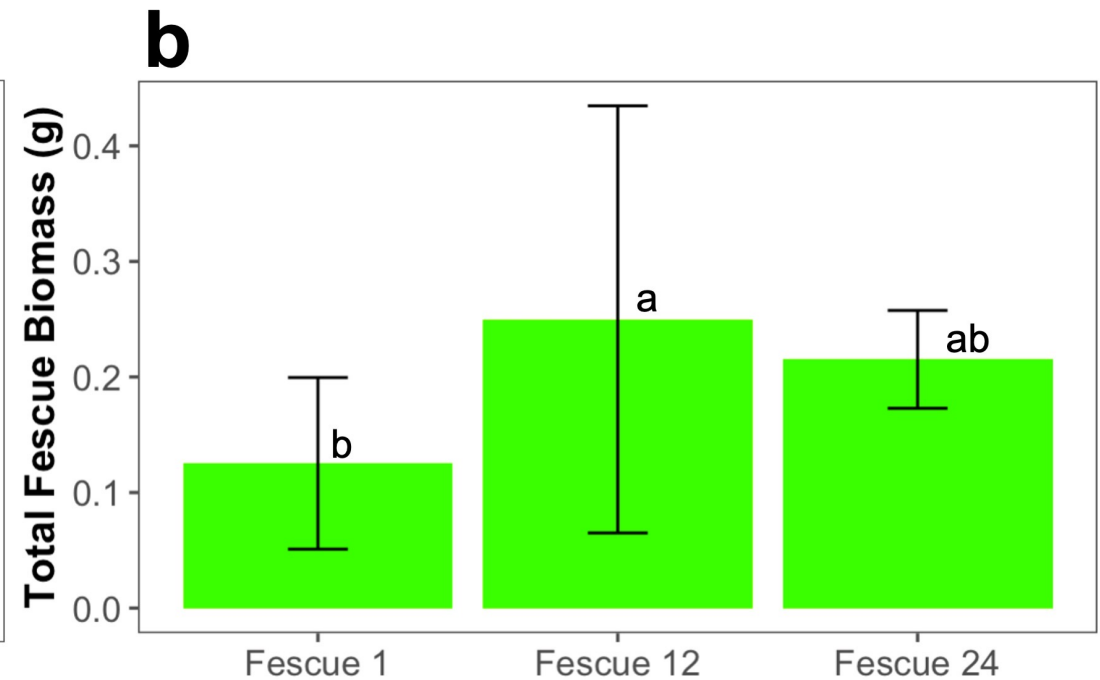
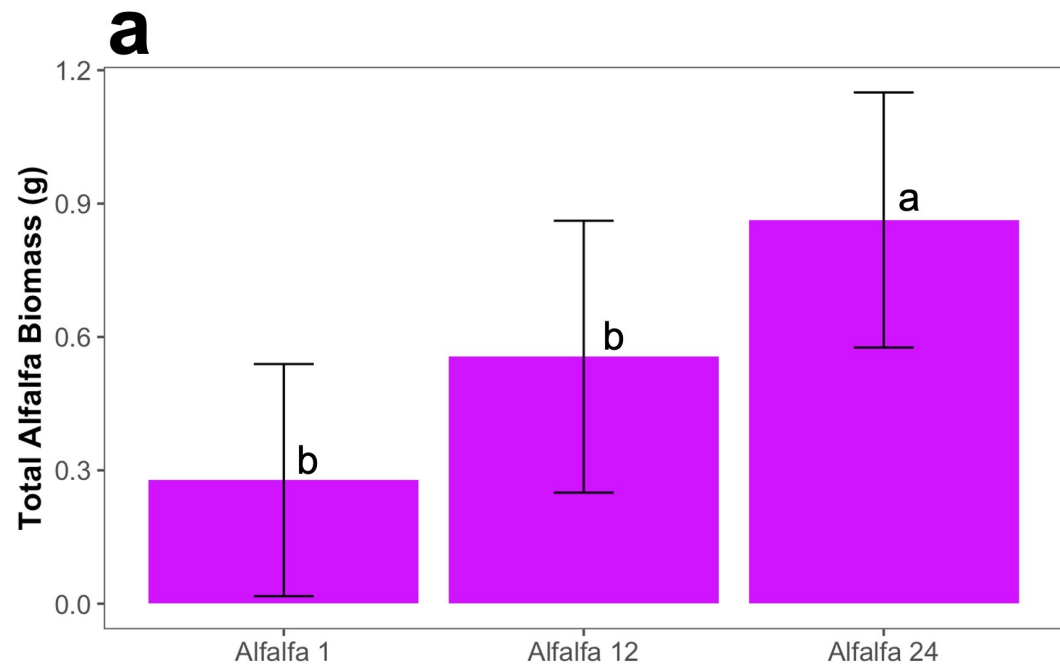


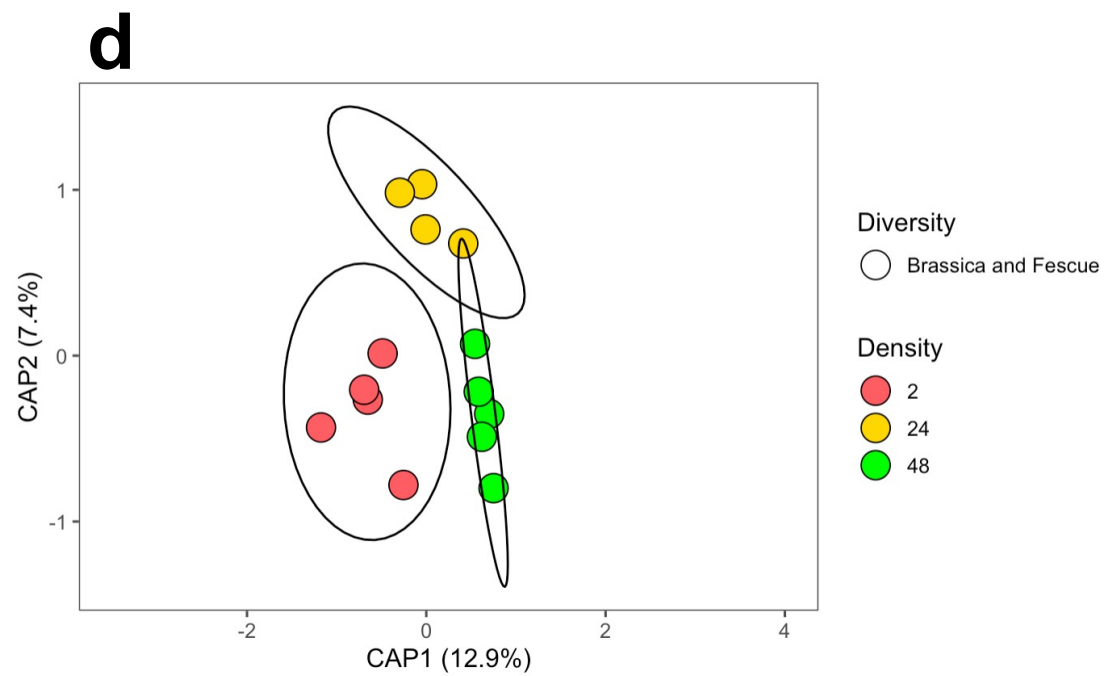
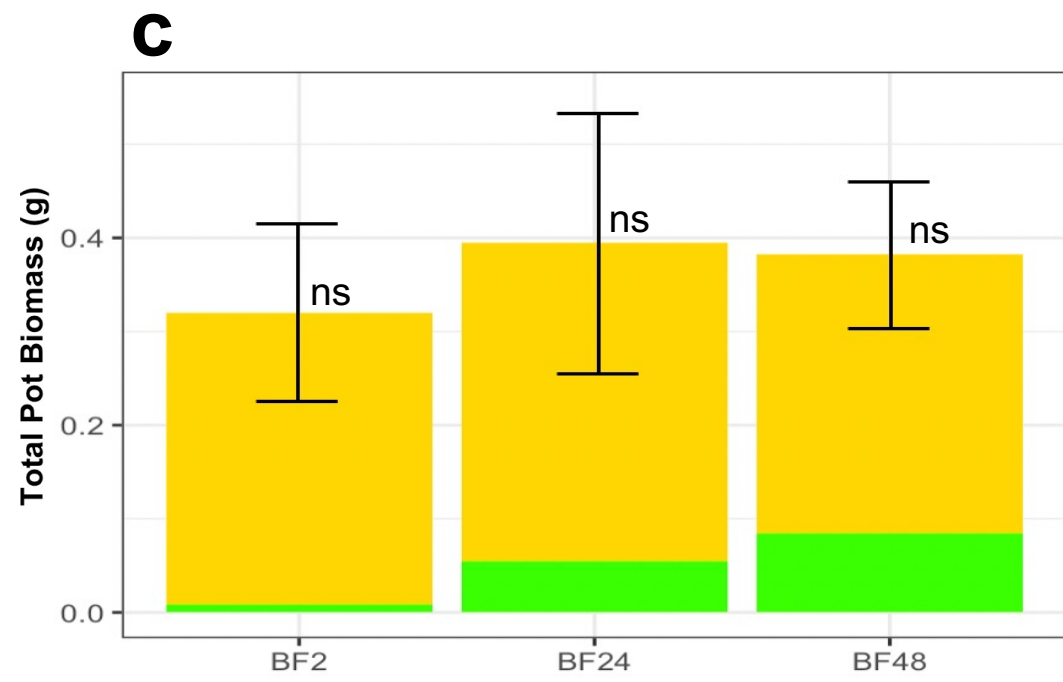
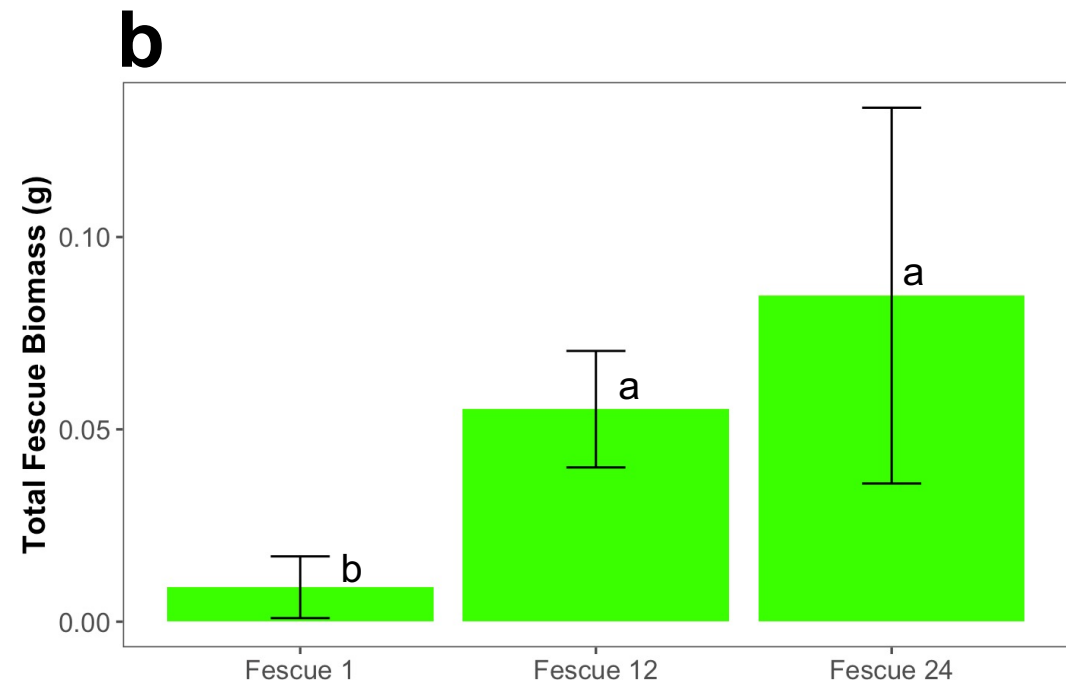
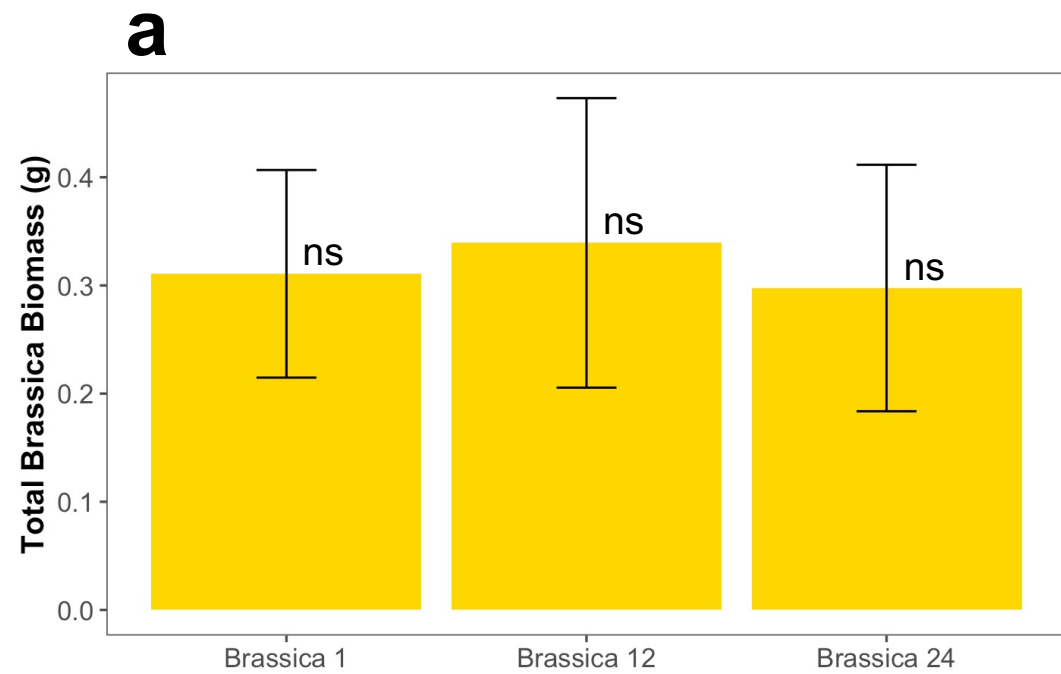




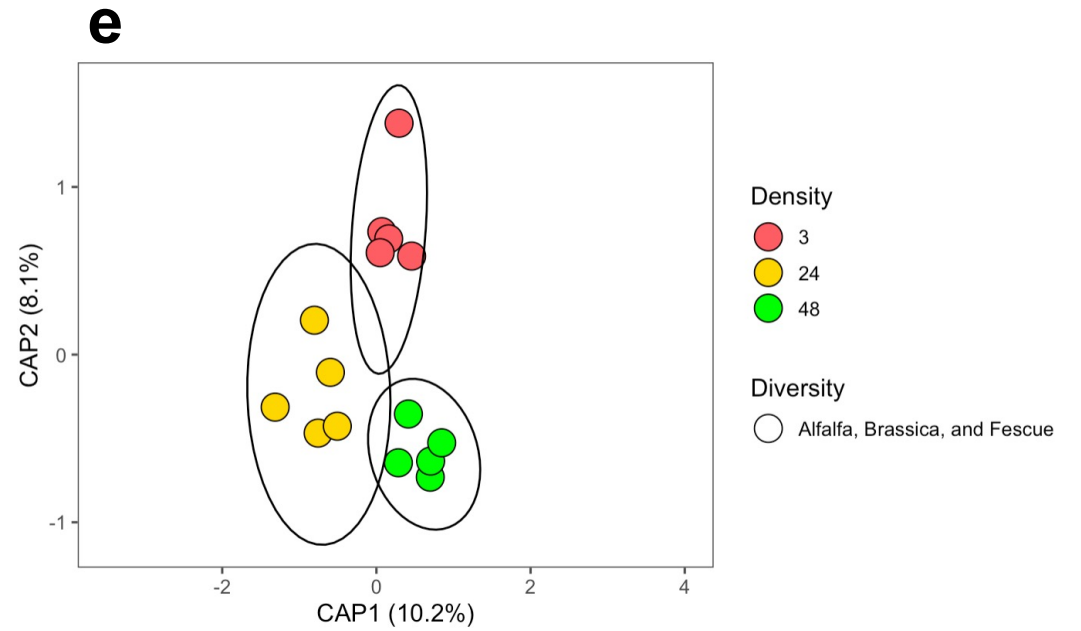
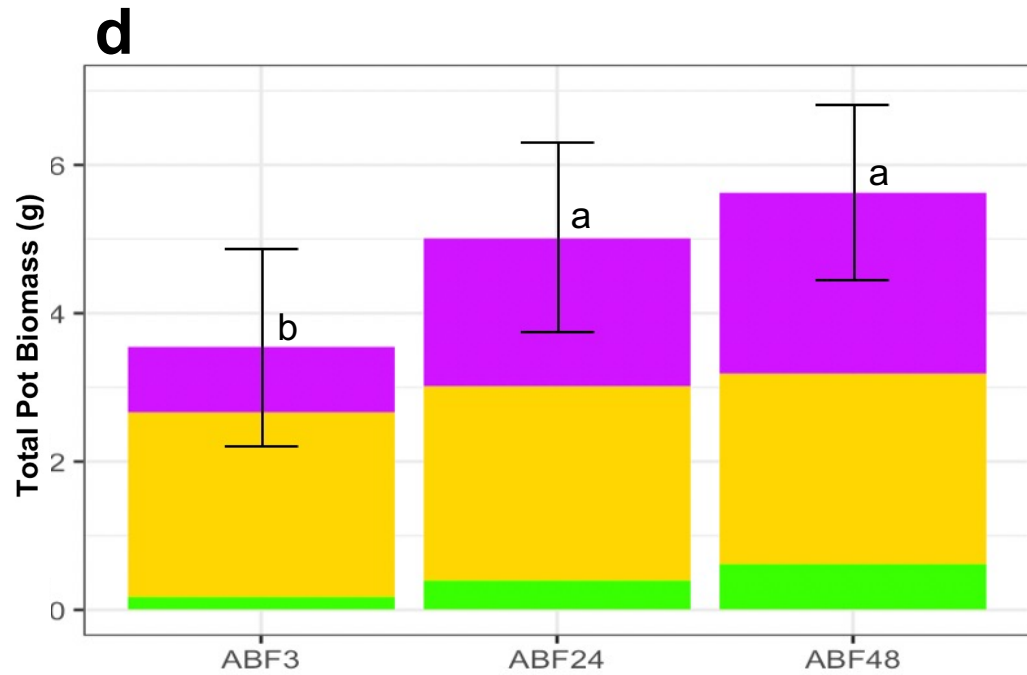
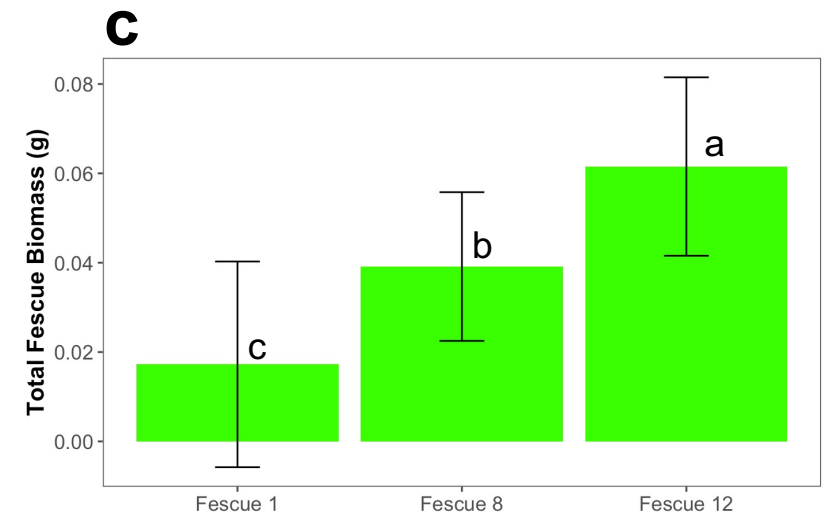
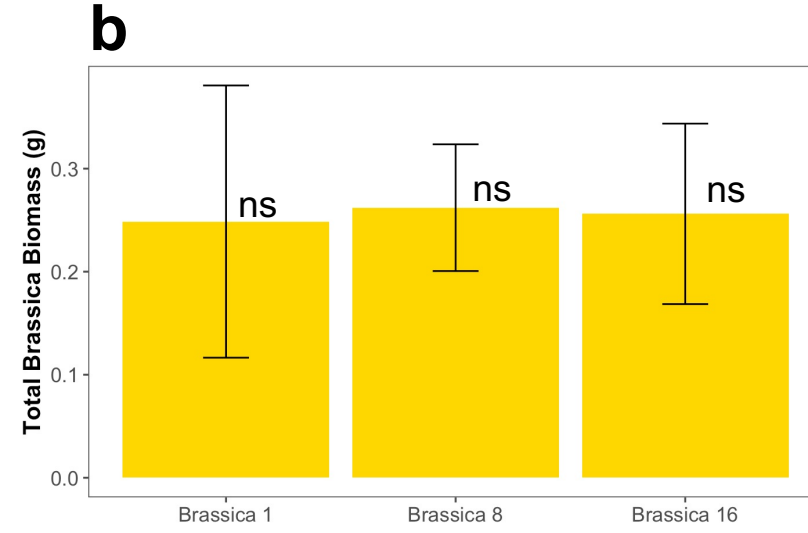
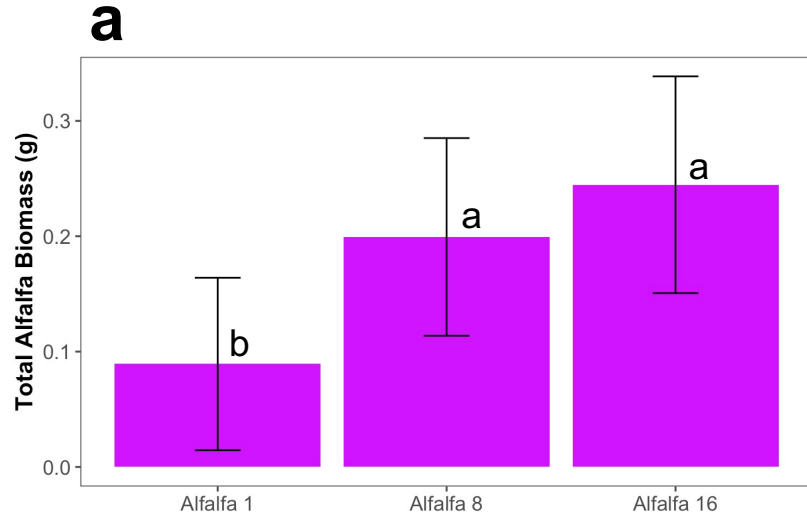












# Overall Summary for the Bulk Soil Bacteriome

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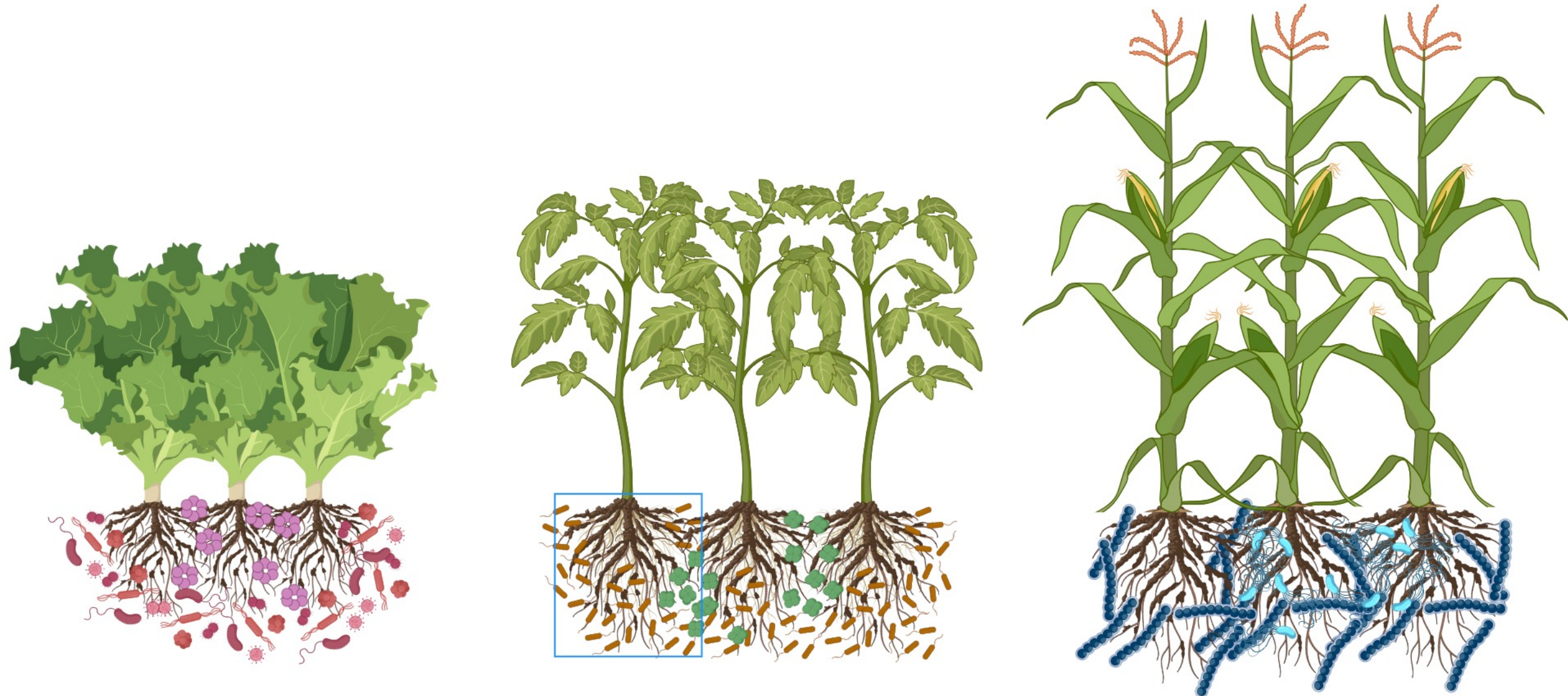
- Bulk Soil of Alfalfa 48 exhibited the least dispersion, suggesting that the surrounding bulk soil bacteriome can progress towards a tailored microbiome for alfalfa as intraspecific competition increases
- Trade off: Different plant combinations enriched different (beneficial) bacteria
  - Even in high plant competition plant mixtures and densities
- *Azospirillum* spp. enriched in alfalfa and brassica monocultures
  - Not enriched in alfalfa-brassica bulk soil
  - *Pseudarthrobacter phenanthrenivorans*- phytohormone producer



# Plant Neighbor and the Rhizosphere



# Plant Neighbor and the Rhizosphere: Density





# Plant Neighbor and the Rhizosphere: Density & Diversity

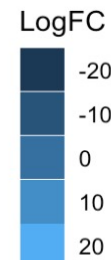
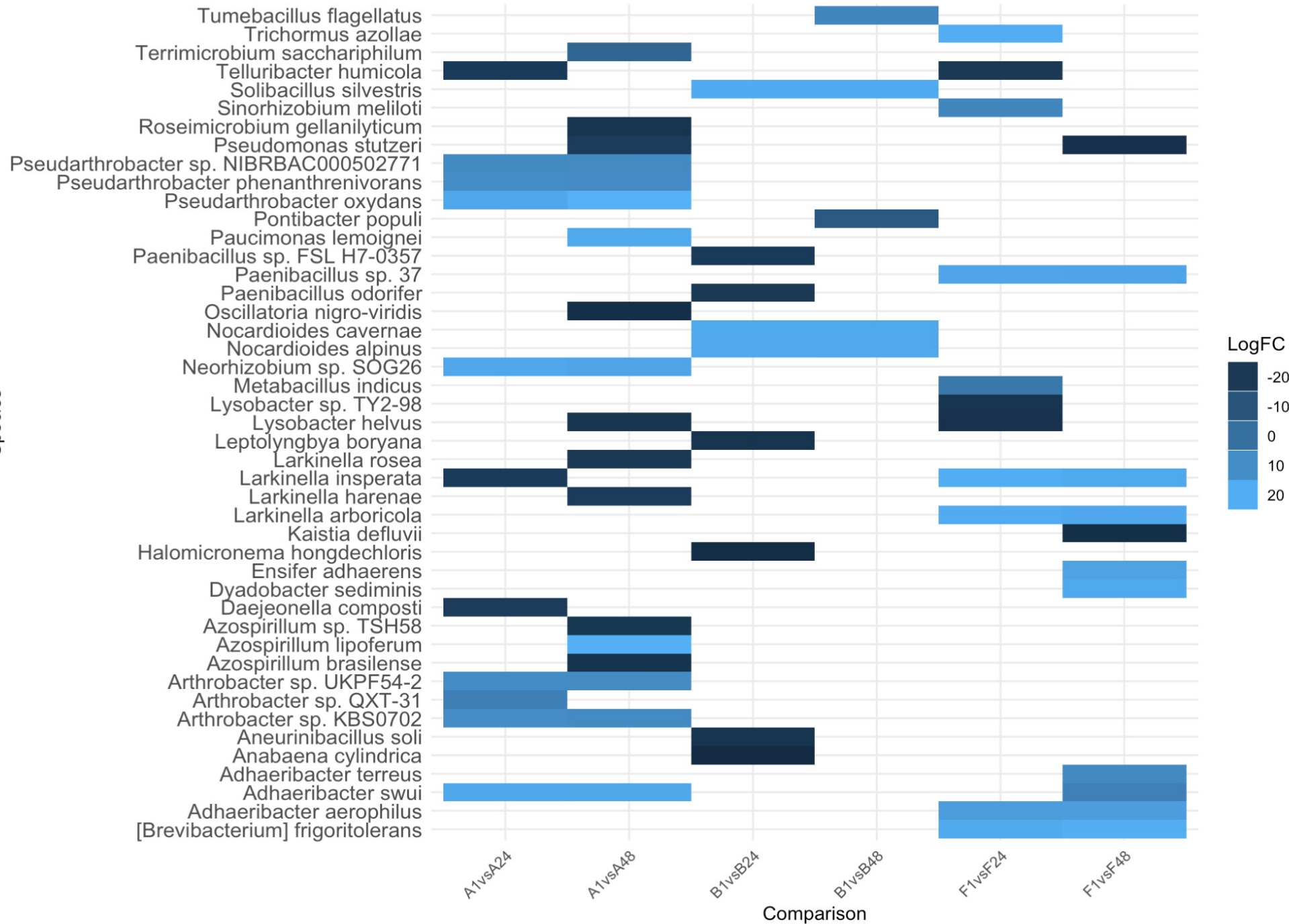


# Rhizosphere: Monoculture



Species

## Differential Abundance for Monoculture



Comparison



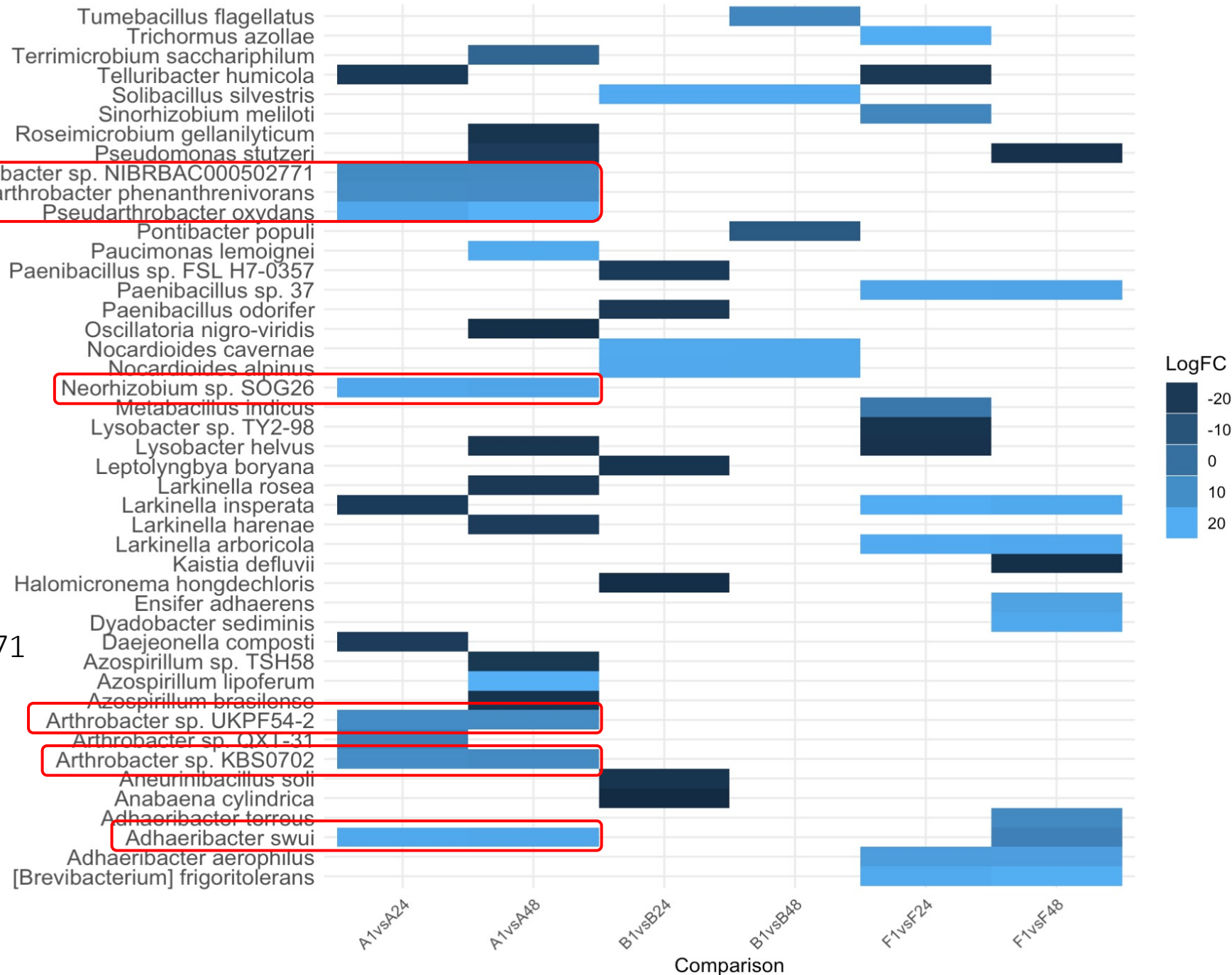
# Rhizosphere: Monoculture Alfalfa



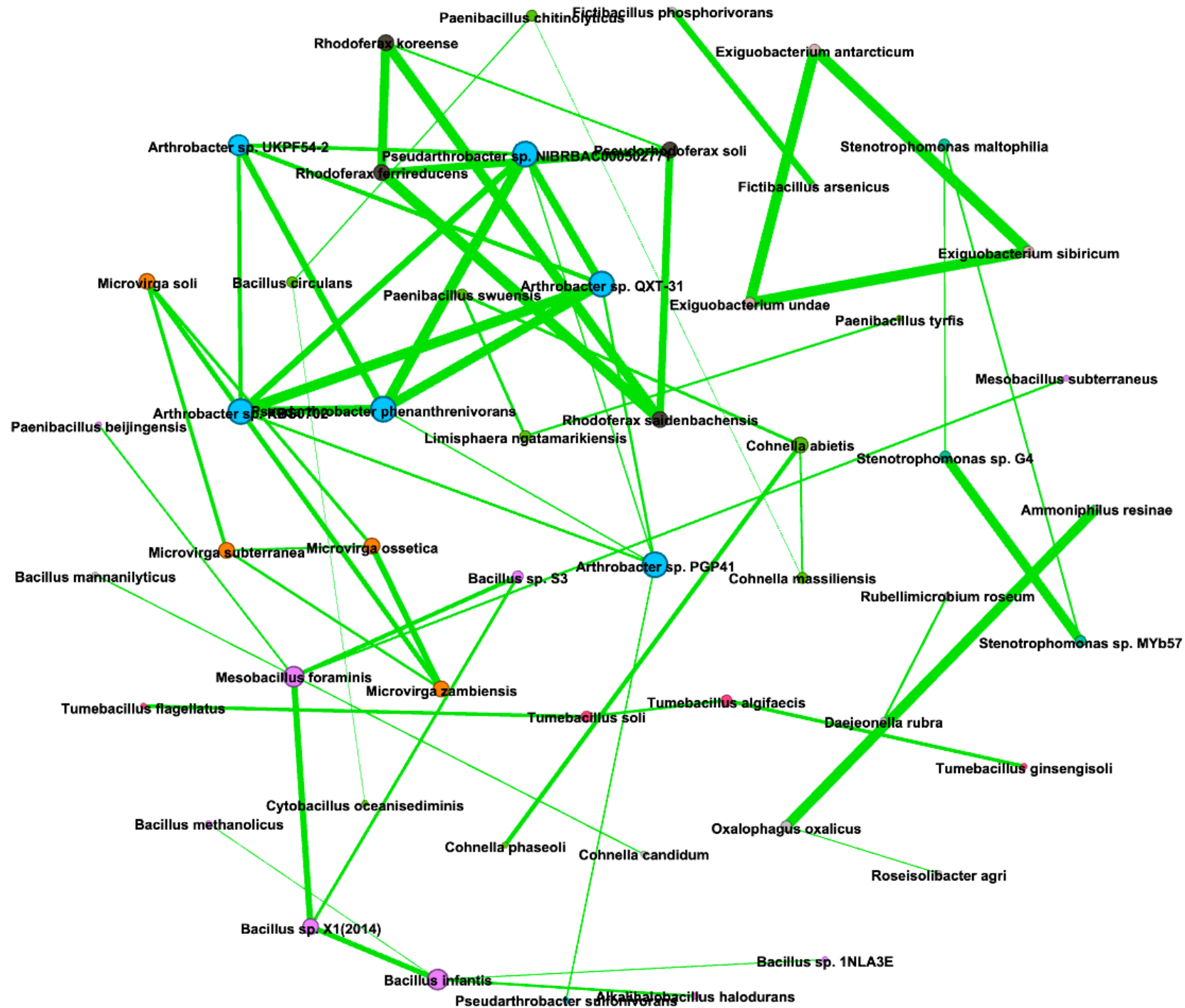
Differential Abundance:  
*Pseudarthrobacter* sp. NIBRBAC000502771  
*Pseudarthrobacter phenanthrenivorans*  
*Pseudarthrobacter oxydans*  
*Neorhizobium* sp. SOG26  
*Arthrobacter* sp. UKPF54-2  
*Arthrobacter* sp. KBS0702  
*Adhaeribacter swiu*

Species

## Differential Abundance for Monoculture

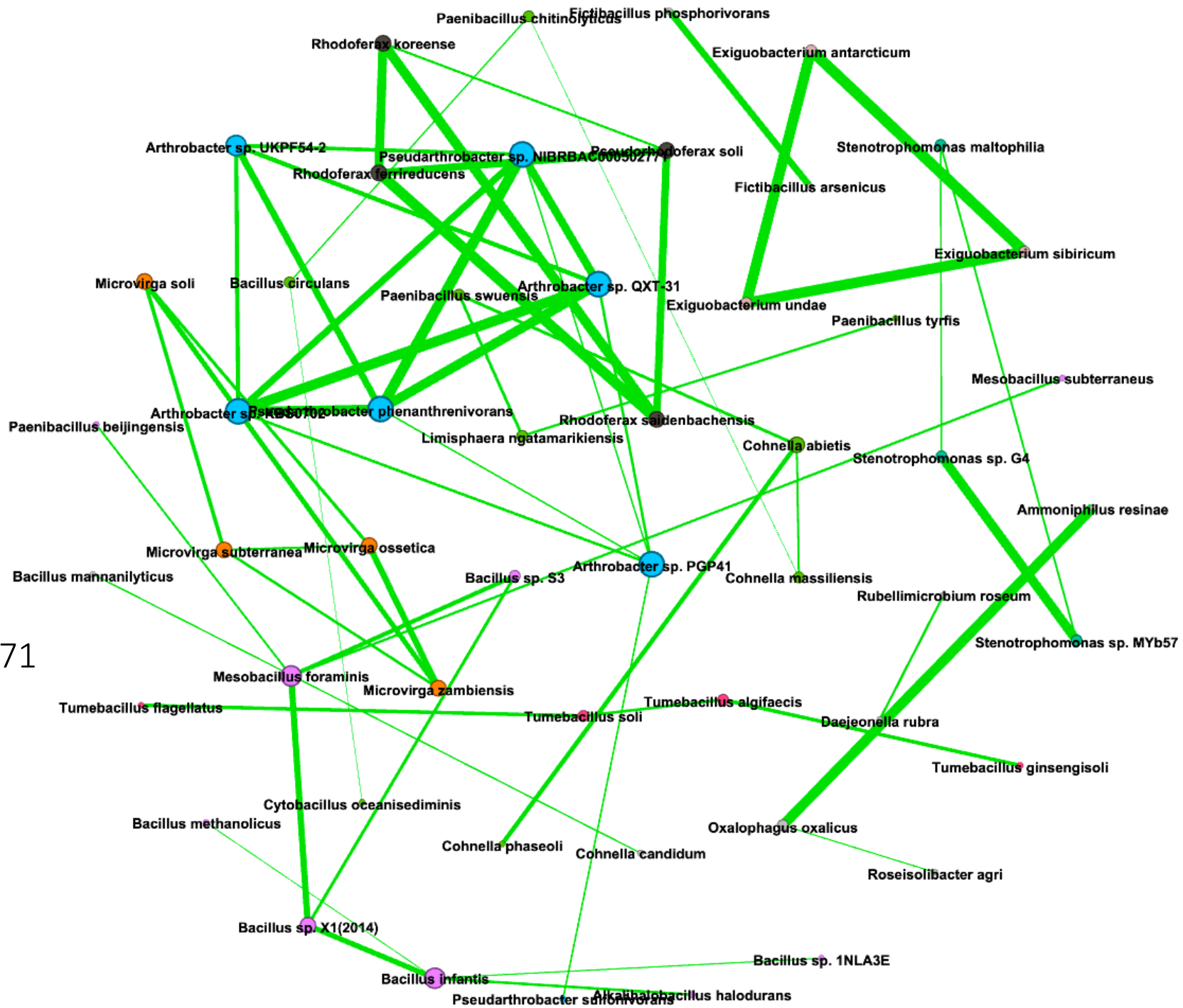


# Alfalfa Rhizosphere Monoculture



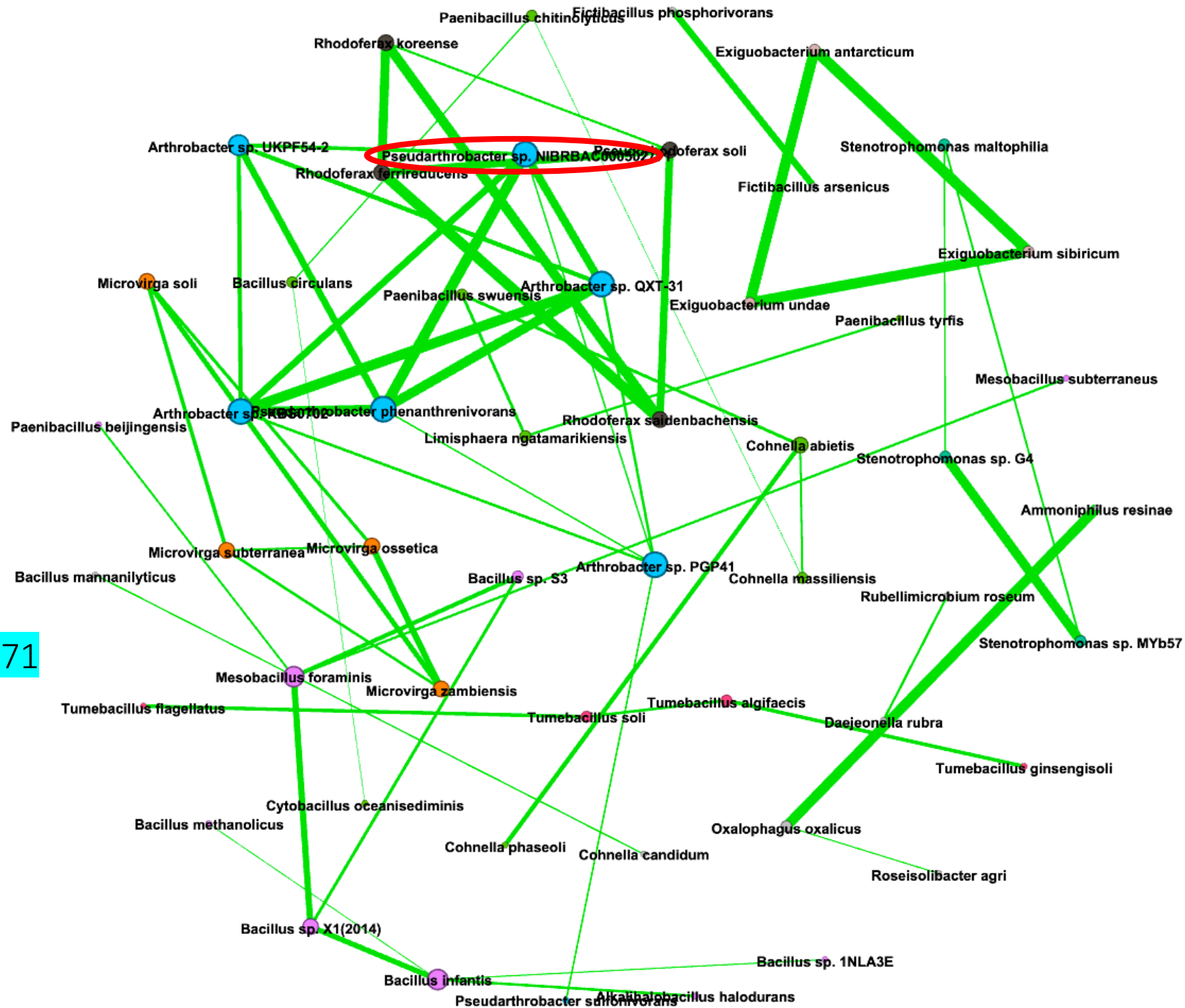


# Alfalfa Rhizosphere Monoculture



Differential Abundance:  
*Pseudarthrobacter* sp. NIBRBAC000502771  
*Pseudarthrobacter phenanthrenivorans*  
*Pseudarthrobacter oxydans*  
*Neorhizobium* sp. SOG26  
*Arthrobacter* sp. UKPF54-2  
*Arthrobacter* sp. KBS0702  
*Adhaeribacter swiu*

# Alfalfa Rhizosphere Monoculture



Differential Abundance:

*Pseudarthrobacter sp. NIBRBAC000502771*

*Pseudarthrobacter phenanthrenivorans*

*Pseudarthrobacter oxydans*

*Neorhizobium sp. SOG26*

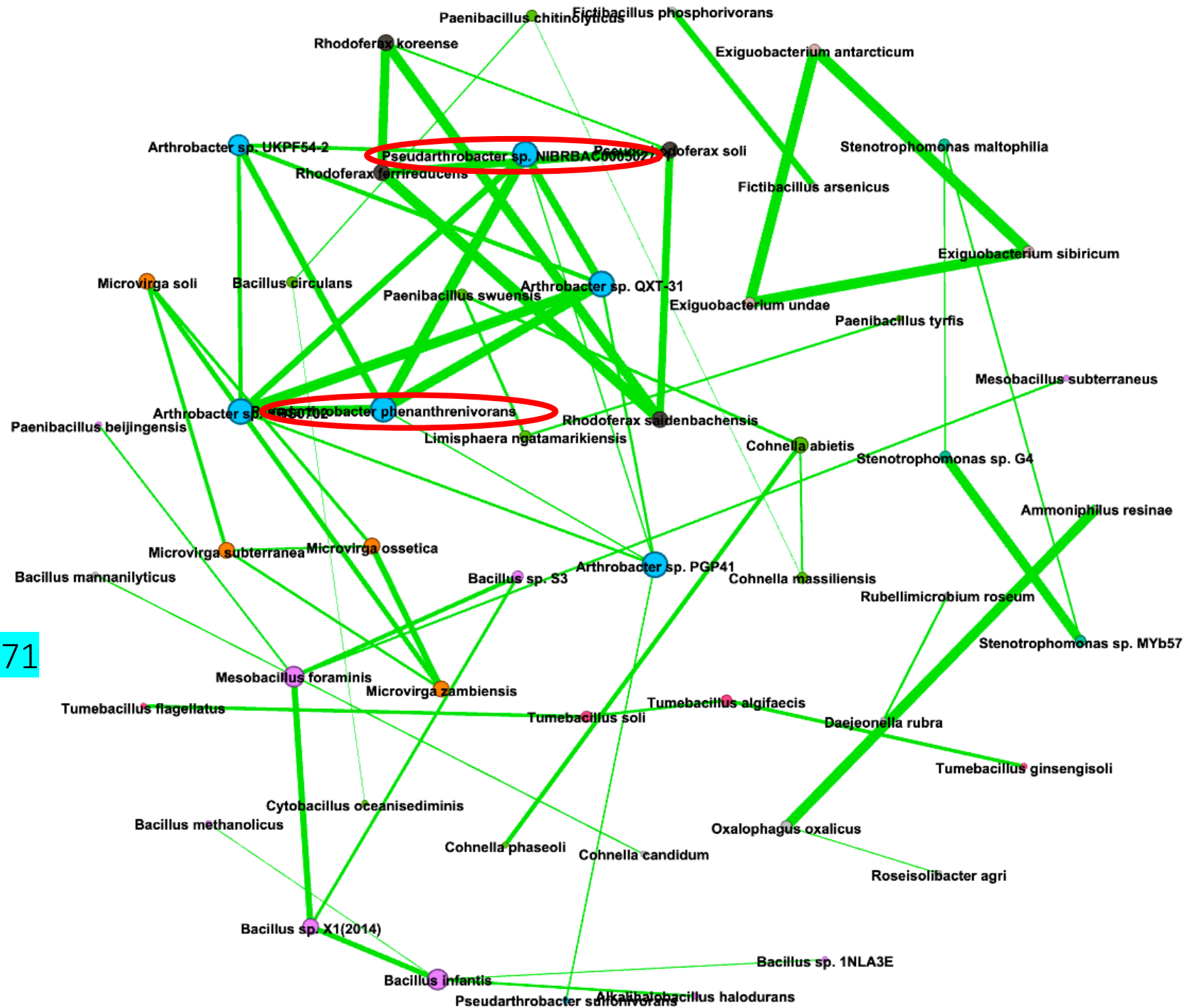
*Arthrobacter sp. UKPF54-2*

*Arthrobacter sp. KBS0702*

*Adhaeribacter swiu*



# Alfalfa Rhizosphere Monoculture



Differential Abundance:

*Pseudarthrobacter* sp. NIBRBAC000502771

*Pseudarthrobacter phenanthrenivorans*

*Pseudarthrobacter oxydans*

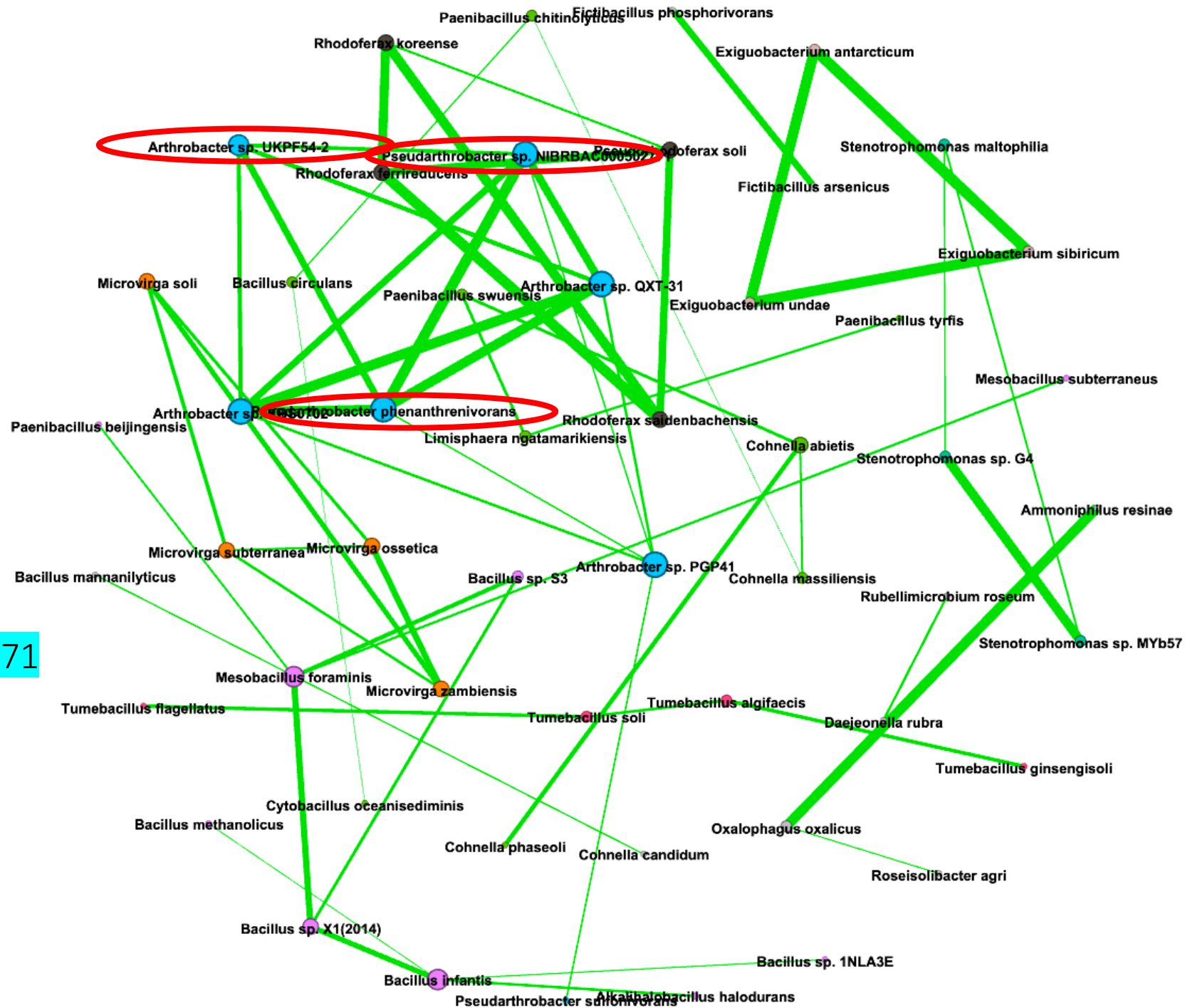
*Neorhizobium* sp. SOG26

*Arthrobacter* sp. UKPF54-2

*Arthrobacter* sp. KBS0702

*Adhaeribacter swiu*

# Alfalfa Rhizosphere Monoculture



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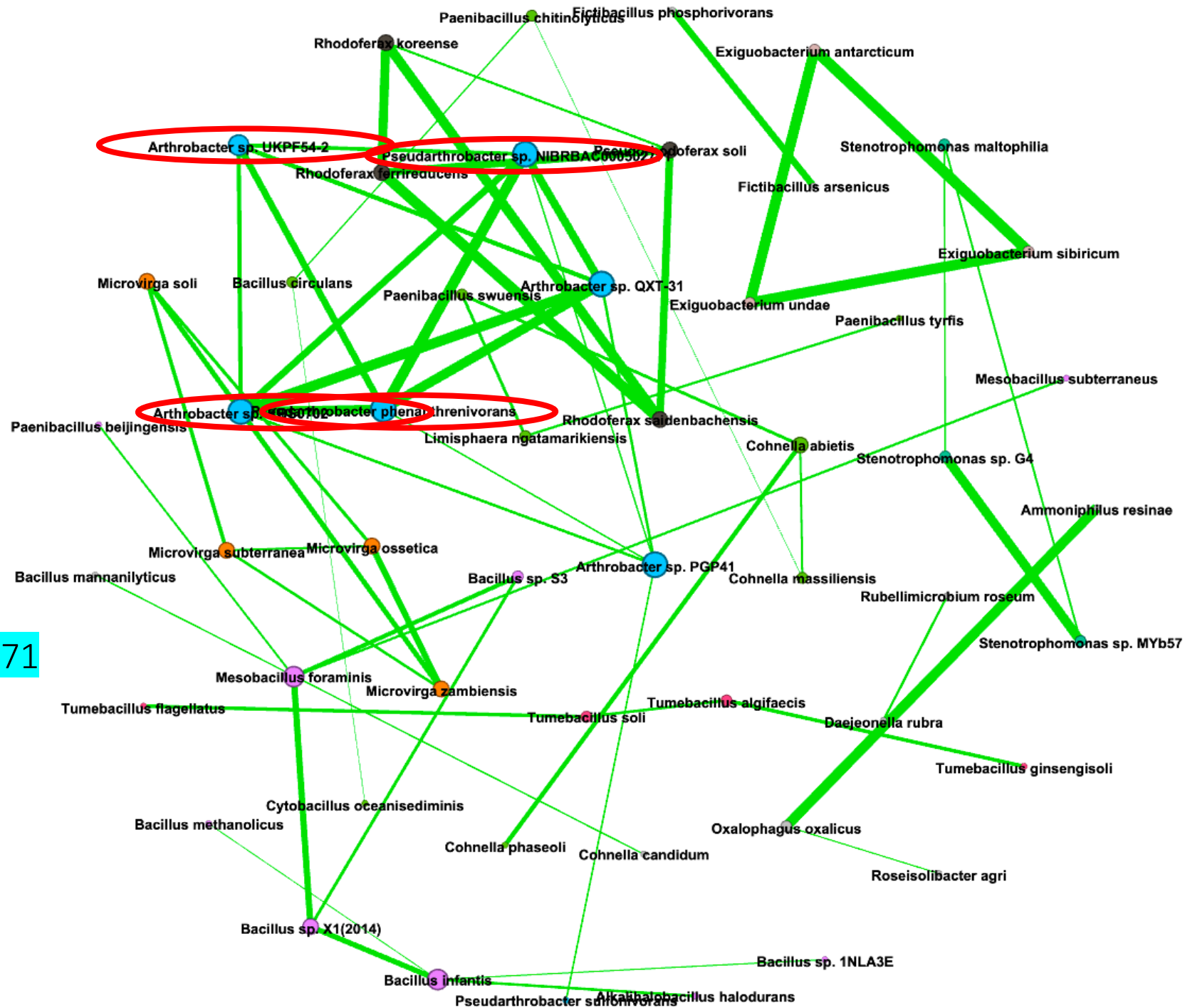
*Arthrobacter* sp. UKPF54-2

*Arthrobacter* sp. KBS0702

*Adhaeribacter swiu*



# Alfalfa Rhizosphere Monoculture



Differential Abundance:

*Pseudarthrobacter* sp. NIBRBAC000502771

*Pseudarthrobacter phenanthrenivorans*

*Pseudarthrobacter oxydans*

*Neorhizobium* sp. SOG26

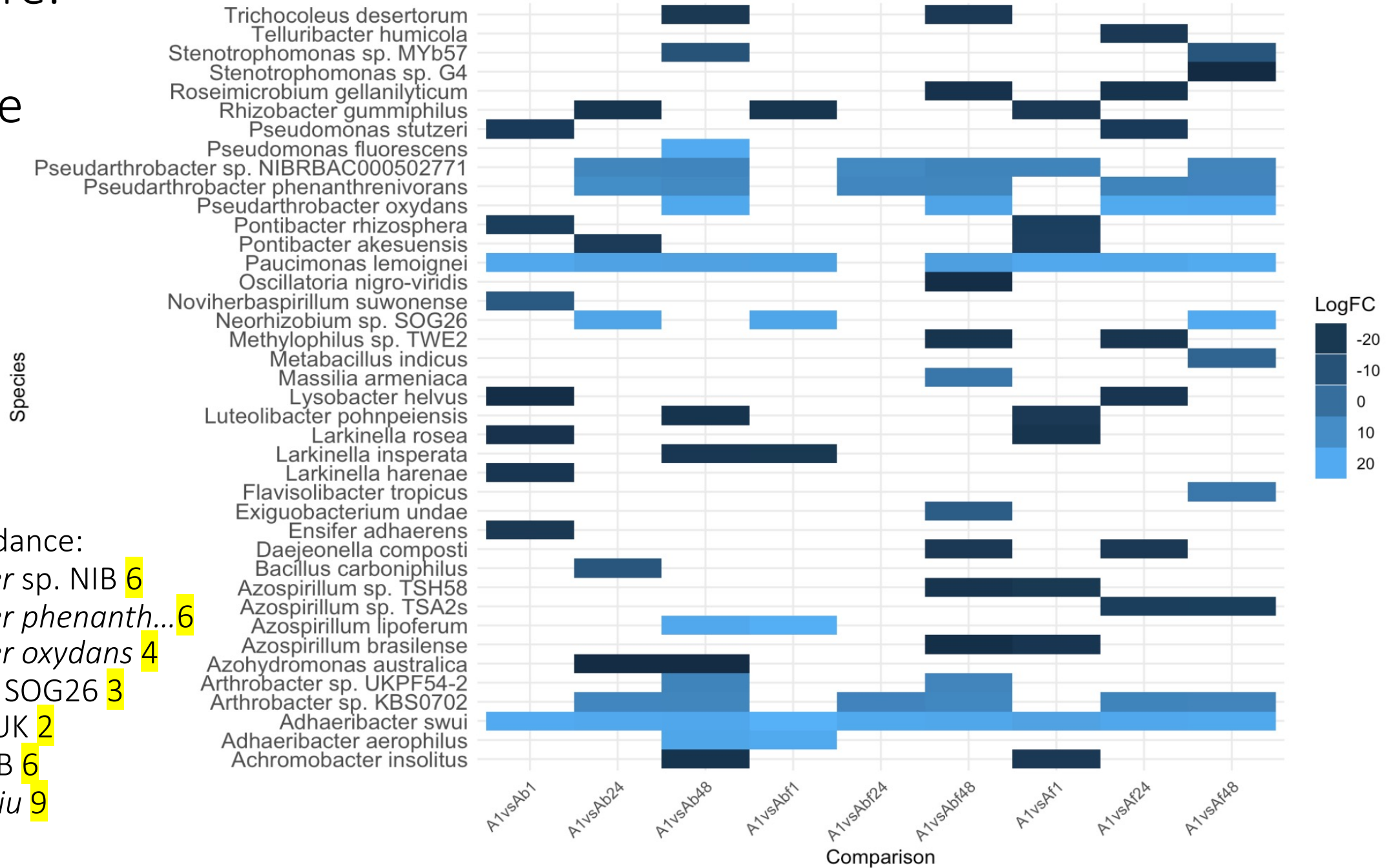
*Arthrobacter* sp. UKPF54-2

*Arthrobacter* sp. KBS0702

*Adhaeribacter swiu*

# Rhizosphere: Alfalfa Polyculture

## Differential Abundance for Alfalfa Polyculture



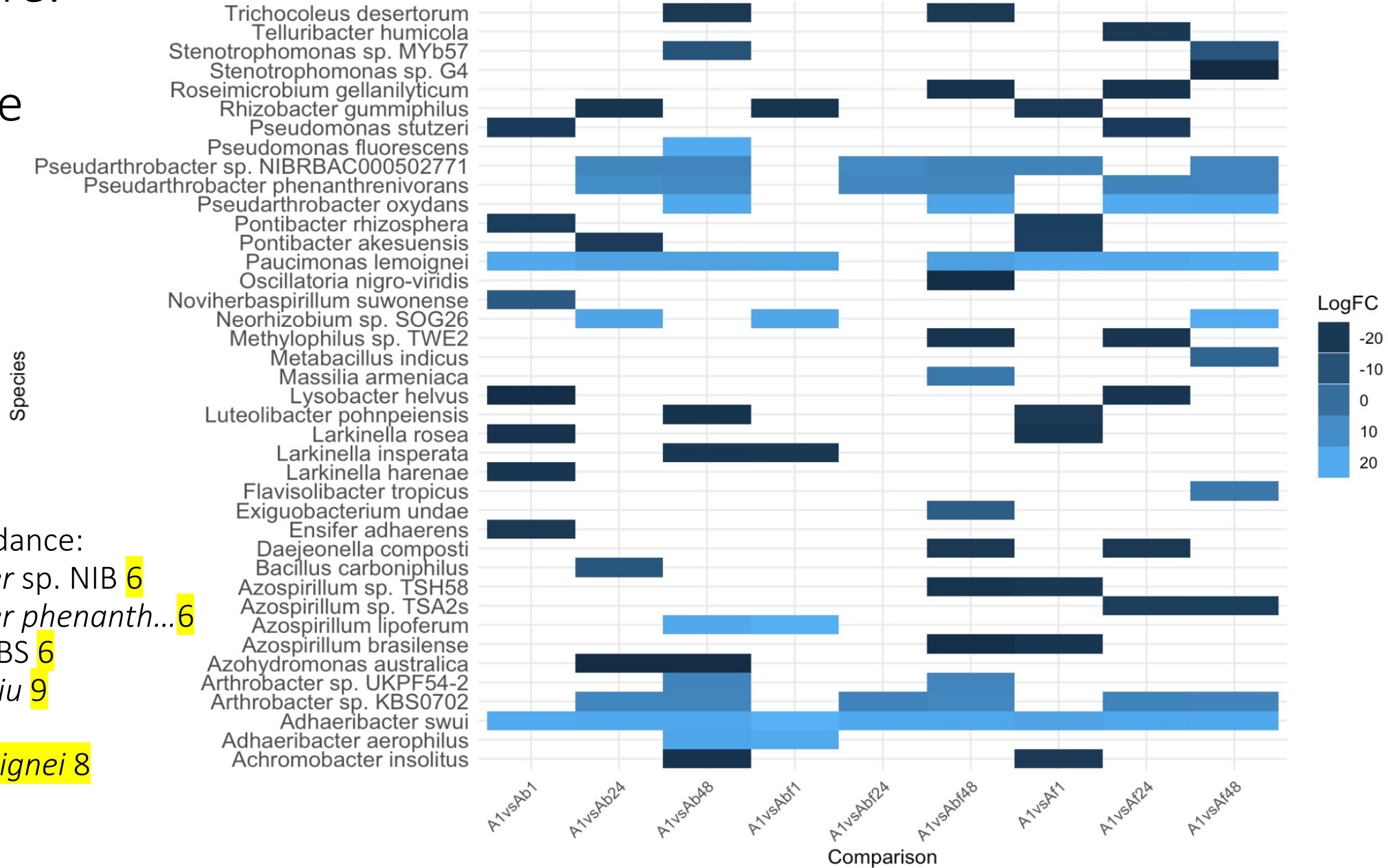
### Differential Abundance:

- Pseudarthrobacter* sp. NIB 6
- Pseudarthrobacter phenanth...* 6
- Pseudarthrobacter oxydans* 4
- Neorhizobium* sp. SOG26 3
- Arthrobacter* sp. UK 2
- Arthrobacter* sp. KB 6
- Adhaeribacter swiu* 9



# Rhizosphere: Alfalfa Polyculture

Differential Abundance for Alfalfa Polyculture



Differential Abundance:

*Pseudarthrobacter* sp. NIB 6

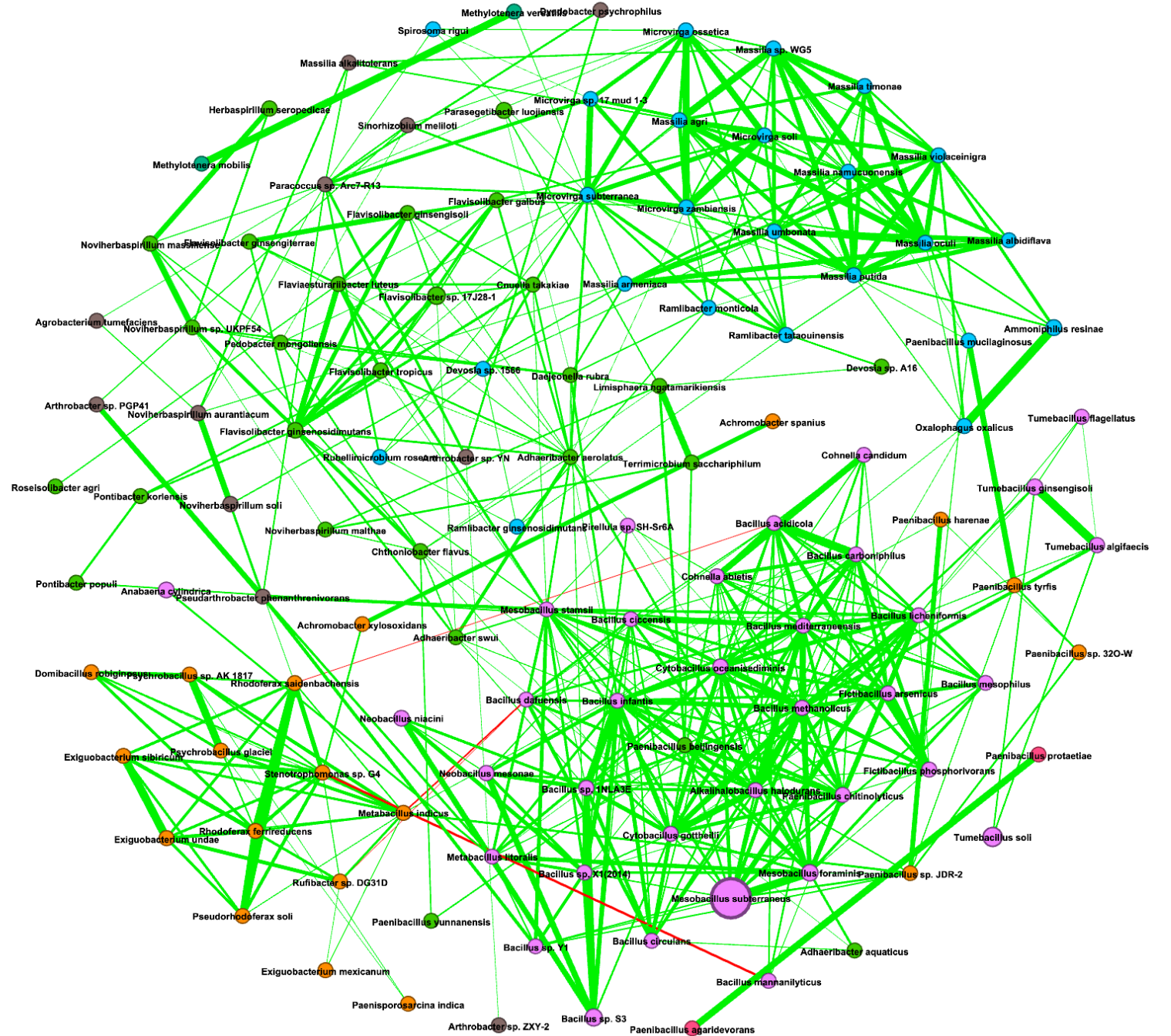
*Pseudarthrobacter phenanth...* 6

*Arthobacter* sp. KBS 6

*Adhaeribacter swiu* 9

*Paucimonas lemoignei* 8

# Alfalfa Rhizosphere Polyculture



Differentially Abundant:

*Pseudarthrobacter* sp. NIBRBAC000502771

*Pseudarthrobacter phenanthrenivorans*

*Pseudarthrobacter oxydans*

*Neorhizobium* sp. SOG26

*Arthrobacter* sp. UKPF54-2

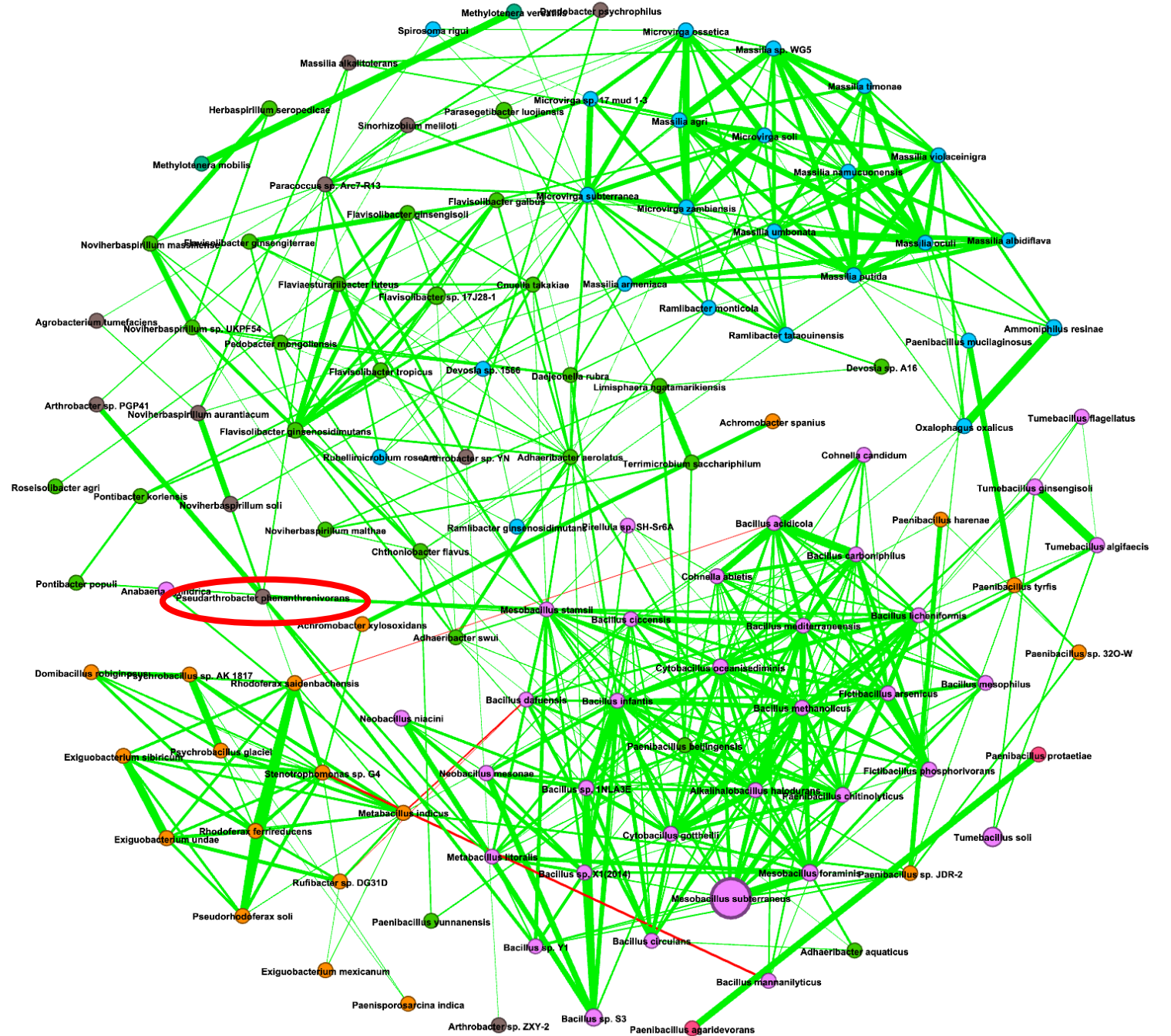
*Arthrobacter* sp. KBS0702

*Adhaeribacter swiu*

*Paucimonas lemoignei*



# Alfalfa Rhizosphere Polyculture



Differential Abundance:

*Pseudarthrobacter* sp. NIBRBAC000502771

*Pseudarthrobacter phenanthrenivorans*

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*Neorhizobium* sp. SOG26

*Arthrobacter* sp. UKPF54-2

*Arthrobacter* sp. KBS0702

*Adhaeribacter swiu*

*Paucimonas lemoignei*



## Future Works

- Polyculture Networks
- Hub Species functionality using Picrust2



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