

# Current approaches and future directions for rhizosphere microbiome sampling methodologies

**Suzanne Fleishman**, William King, Caylon Yates, Meredith Persico, Jing Guo, Michela Centinari, Terrence Bell, David Eissenstat



Dr. Michela Centinari



Dr. Terry Bell



Dr. David Eissenstat



Dr. William King  
Post-doctoral Scholar



Caylon Yates  
PhD Candidate

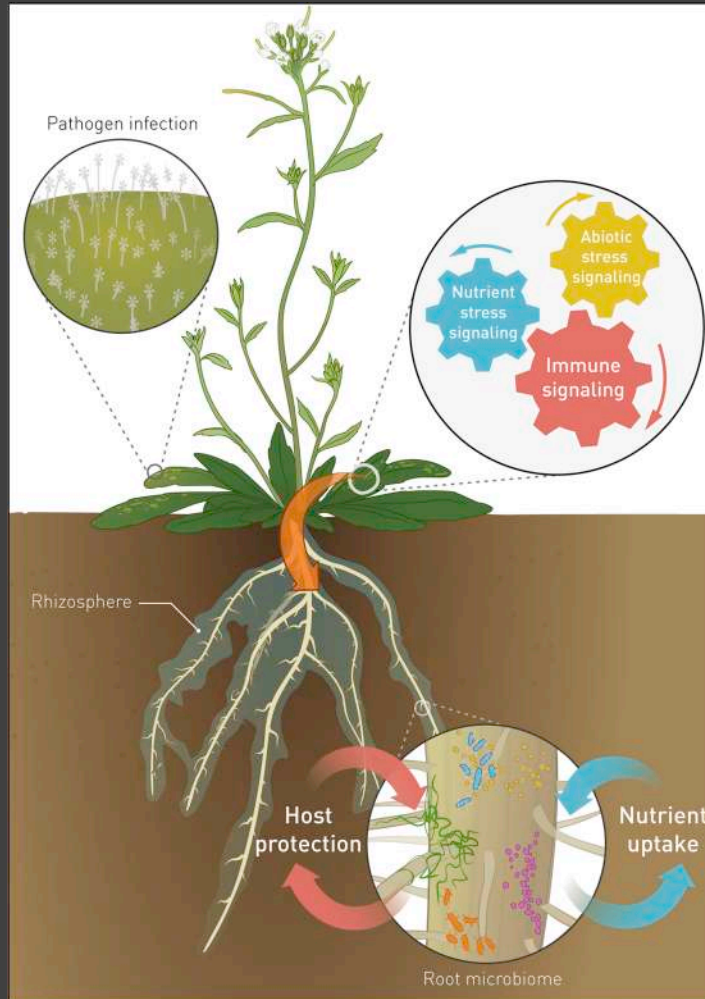


Meredith Persico  
PhD Candidate



Dr. Jing Guo  
Visiting scholar

# Many plant functions are microbially-mediated



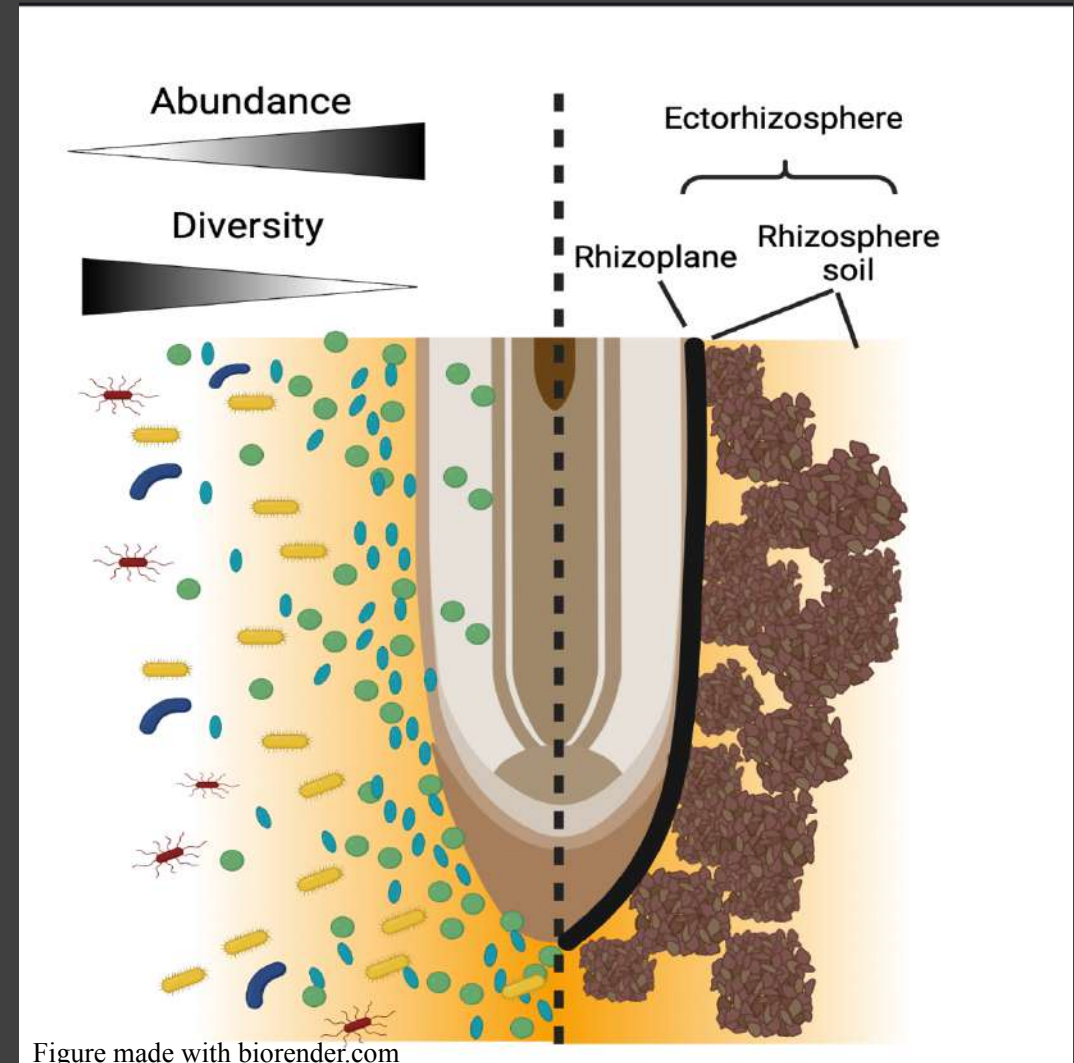
Bakker et al. 2018, Trends in Plant Science

Fleishman et al. 2022, *In Review*

# What exactly is the rhizosphere?

Narrow soil zone directly surrounding the root system and influenced by the root

(Reinhold-Hurek et al., Annu. Rev. Phytopathol. 2015)



# No standard practices for root sampling and rhizosphere removal

- 36% as potted studies
- In field studies, 30% sampled entire root system and 34% from the shallow depths
- 88% on bacteria (16s)
- 33% on fungi (ITS)



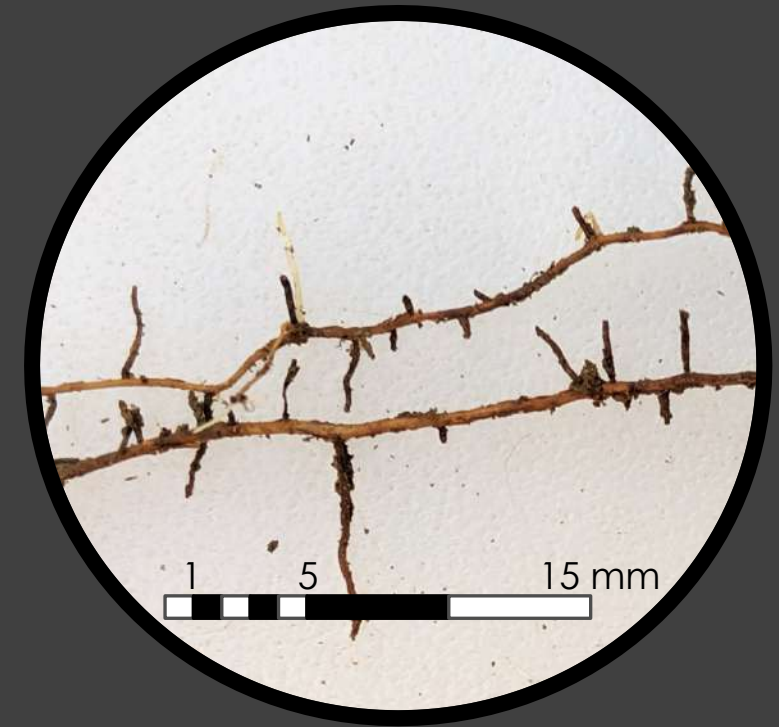
# How does the underlying functional and spatial heterogeneity of roots impact microbial composition?



Absorptive  
vs.  
Transportive



Developmental  
stage



Spatial  
location

Root systems are heterogeneous environments





# Absorptive vs. transportive roots





# Absorptive vs. transportive roots



Root function can be classified by branching order





Root function can be classified by branching order

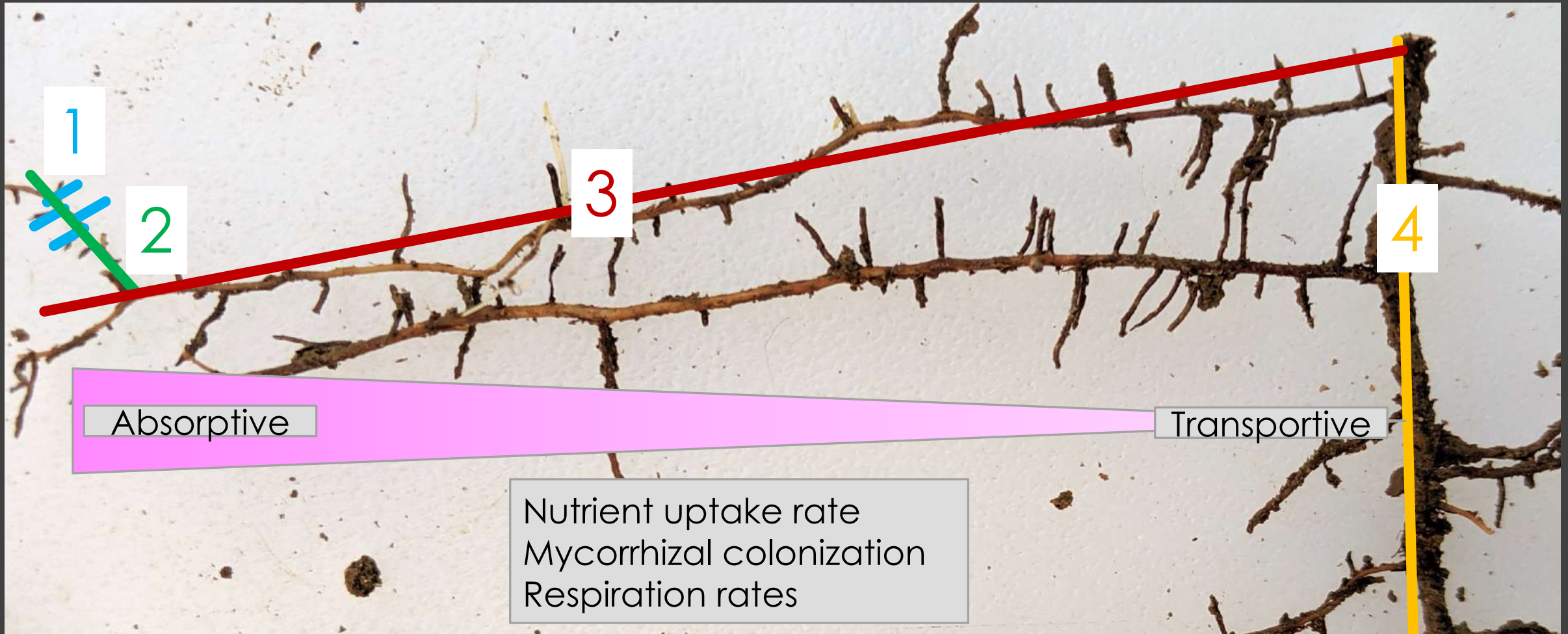




Root function can be classified by branching order



# Root function can be classified by branching order





# Do rhizosphere microbiomes vary with root branching order?

**Common garden**

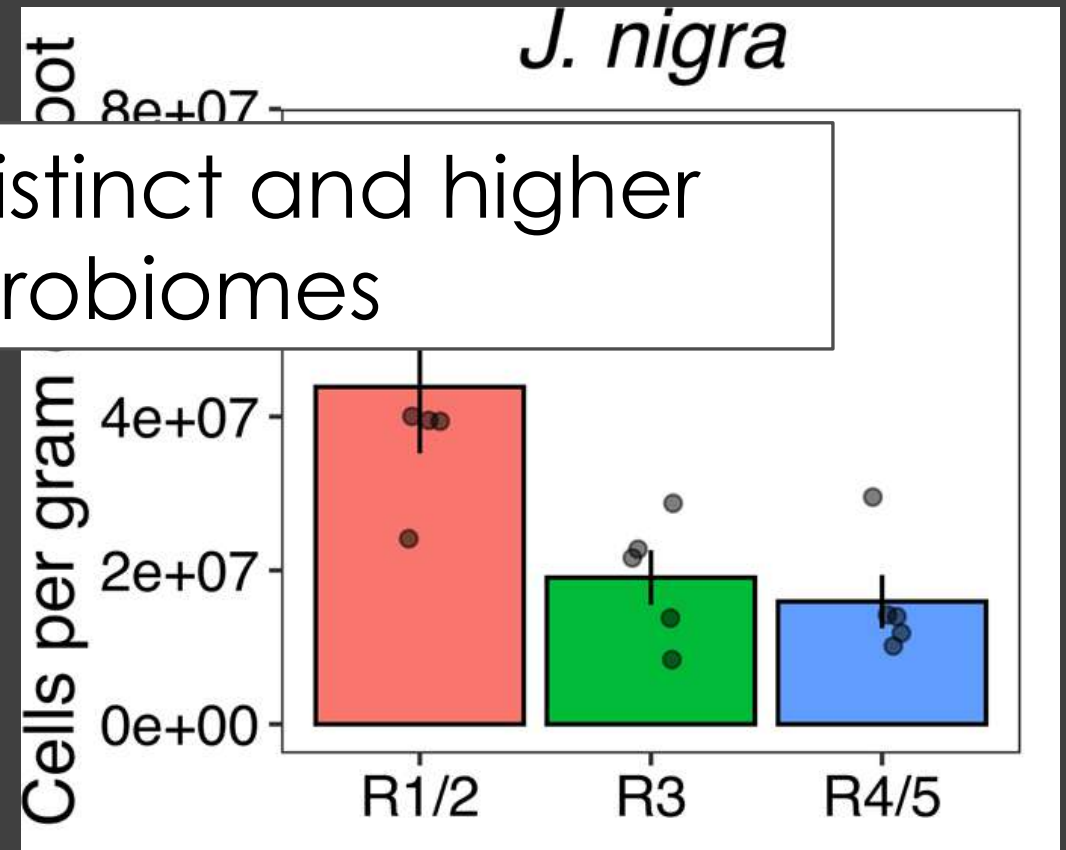
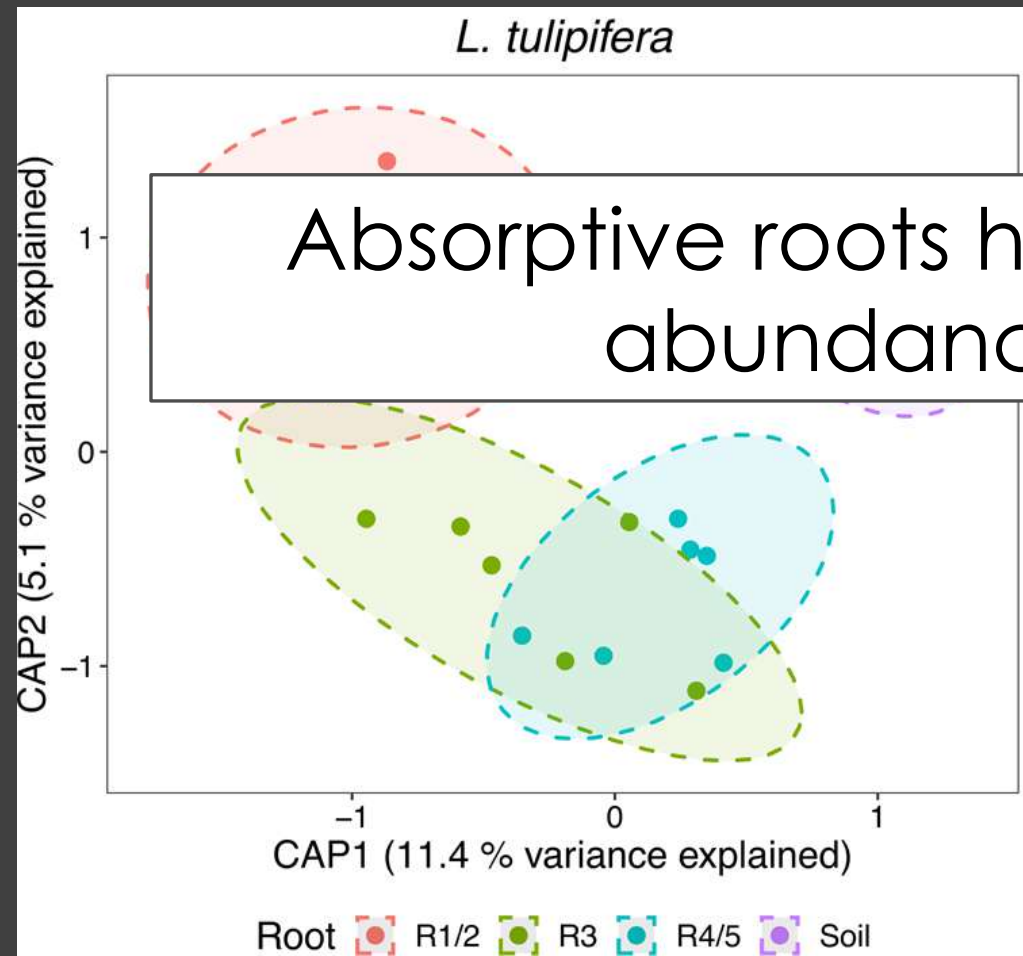


*Liriodendron tulipifera*





# Root branching order determines composition



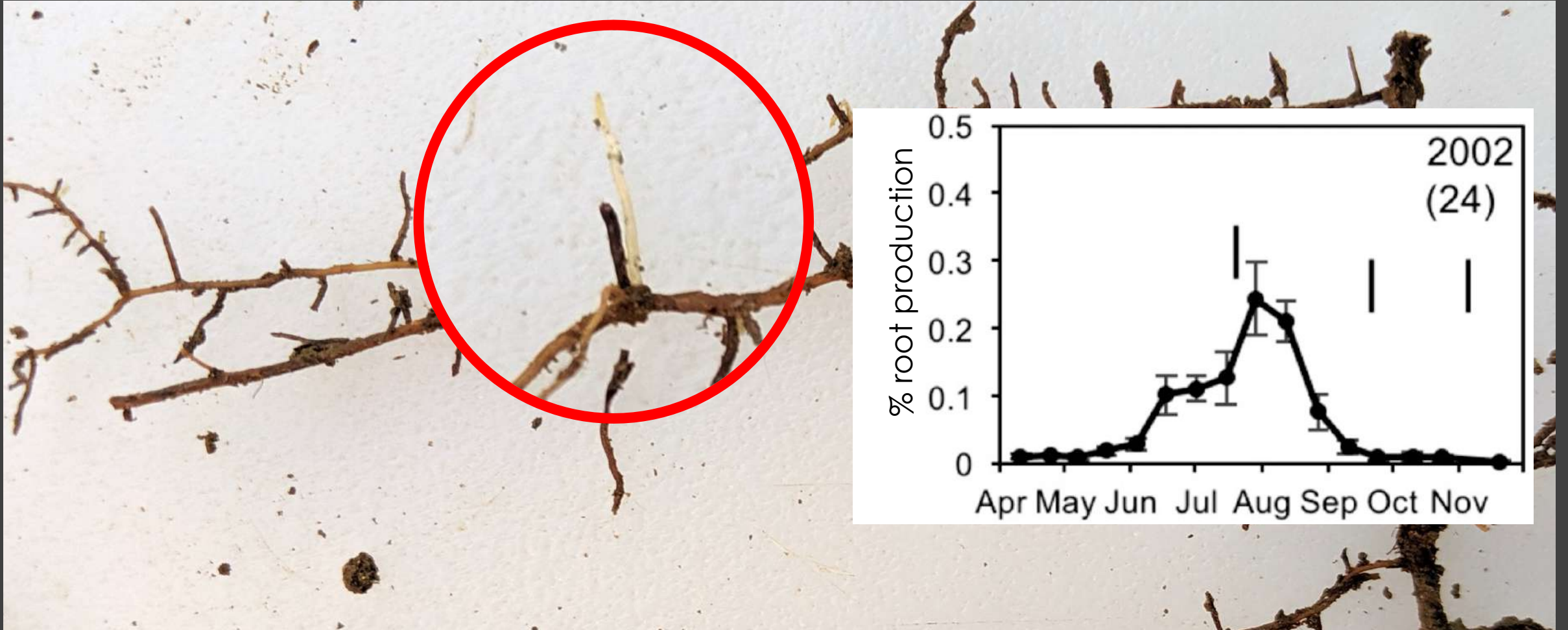
Absorptive roots have distinct and higher abundance microbiomes

\* Pattern existed across all 6 tree species

Absorptive roots differ in developmental stage

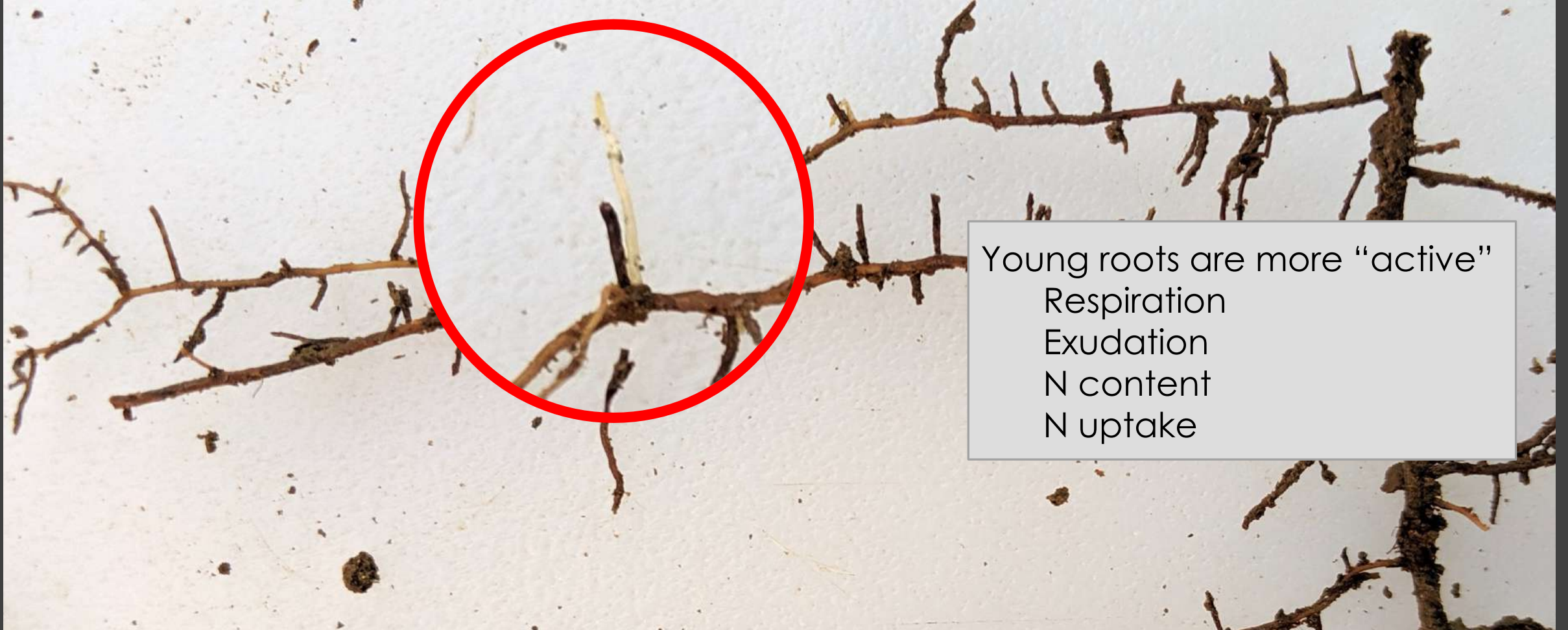


# Absorptive roots differ in developmental stage





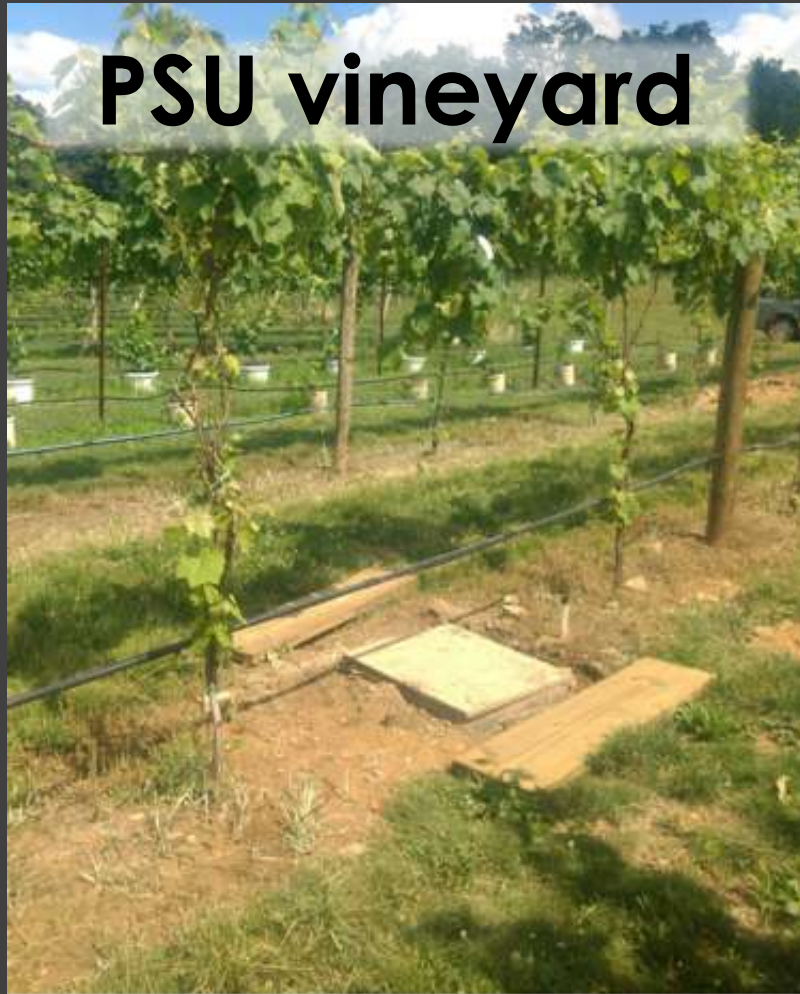
# Absorptive roots differ in developmental stage



Young roots are more “active”  
Respiration  
Exudation  
N content  
N uptake



# Monitoring root age in the field





# Monitoring root age in the field



7 vines  
98 individual roots  
Age: 0.5 – 40 days



Microbial composition minimally differs between young (<11 days) and old (11-44 days) roots

Incredibly high root-to-root variation in rhizosphere bacterial and fungal composition

Does root location explain microbial variation?



Does root location explain microbial variation?





# Mapping a root system



## Variation Partitioning Analysis (VPA)

- Based on constrained ordinations (RDA) to model microbiome response
- Allows for  $R^2$  generation for multivariate explanatory categories
- Allows for comparison of the relative importance of each category

# Mapping a root system



## Variation Partitioning Analysis (VPA)

1. Root traits  
(branching order, age, color)

# Mapping a root system

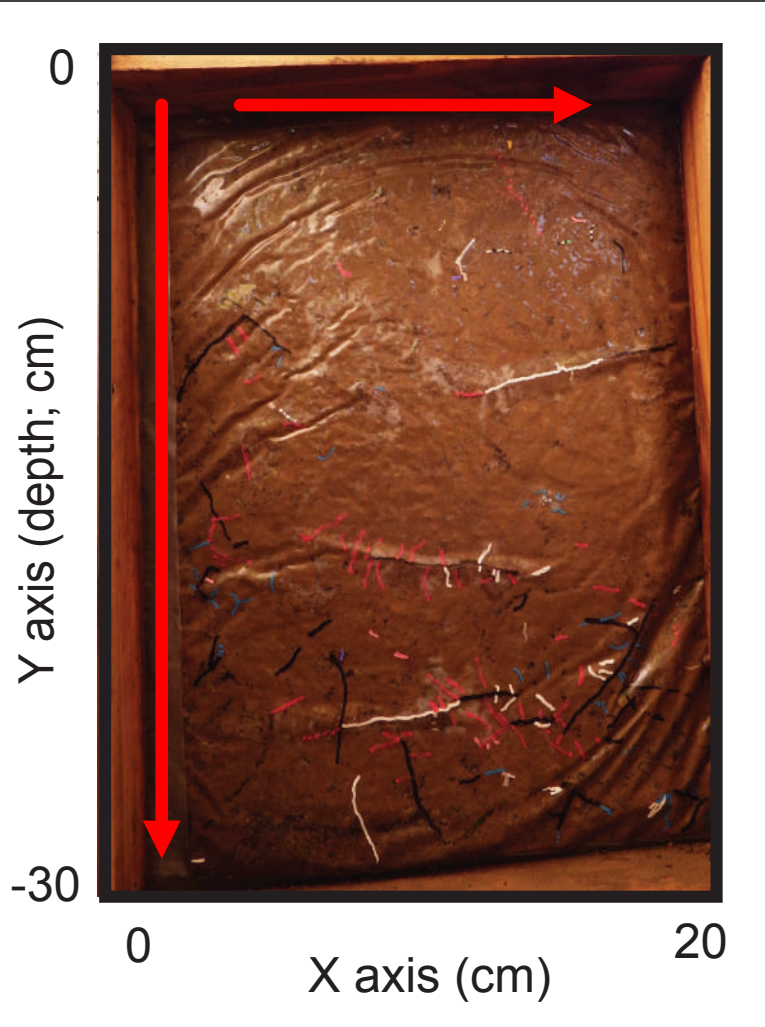


## Variation Partitioning Analysis (VPA)

1. Root traits  
(branching order, age, color)
2. Root cluster



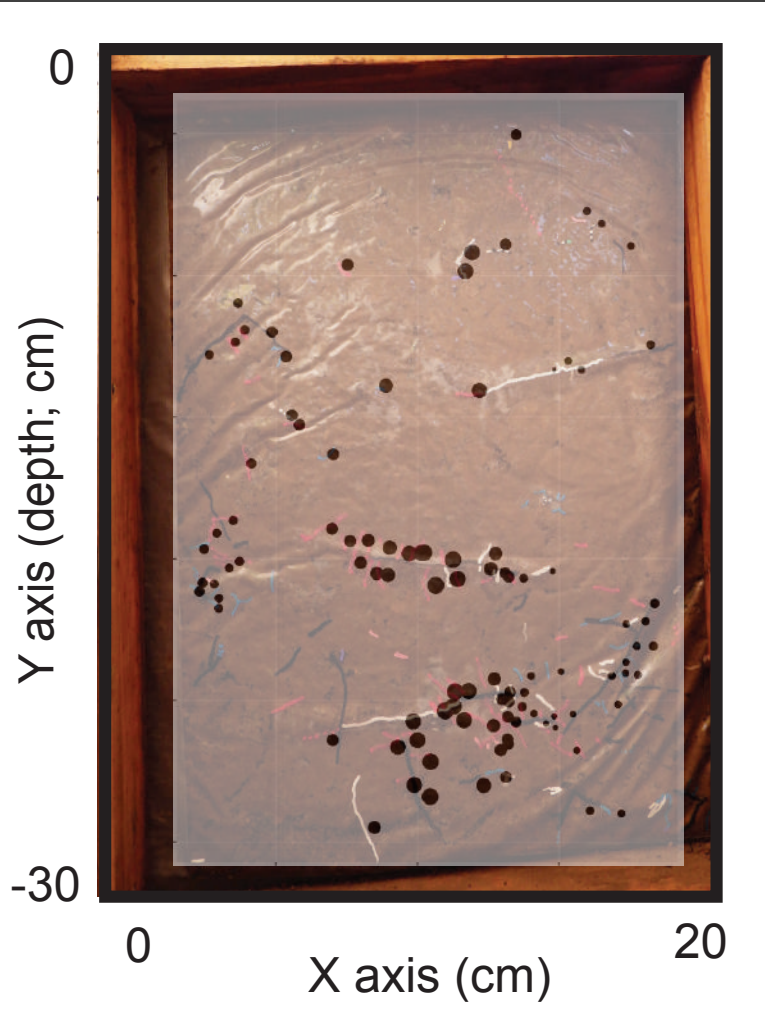
# Mapping a root system



## Variation Partitioning Analysis (VPA)

1. Root traits  
(branching order, age, color)
2. Root cluster
3. Linear spatial trend (X,Y)

# Mapping a root system



## Variation Partitioning Analysis (VPA)

1. Root traits  
(branching order, age, color)
2. Root cluster
3. Linear spatial trend (X,Y)
4. Local groupings (MEMs)

Root traits explain less variation than  
spatially structured factors





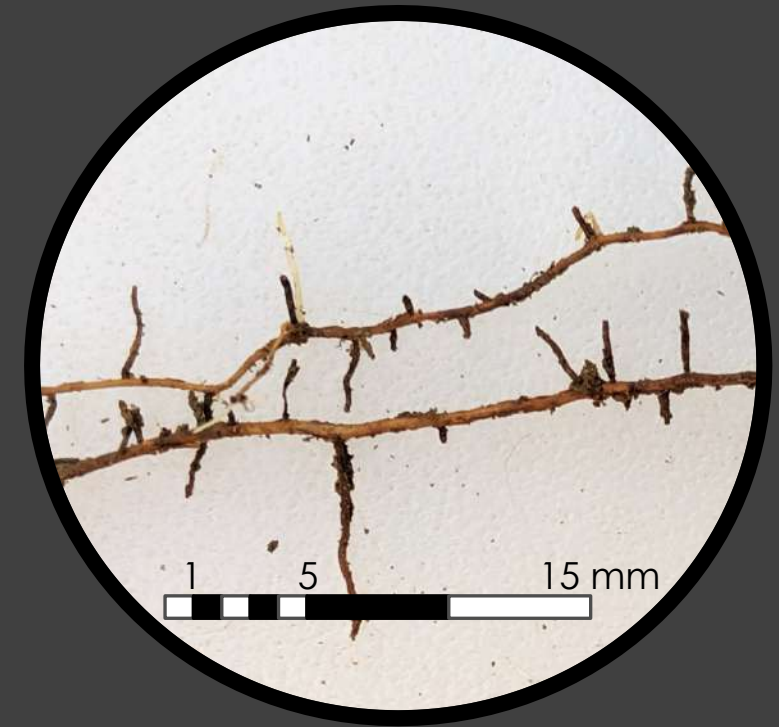
# The underlying functional and spatial heterogeneity of roots impacts microbial composition



Branching  
order



Developmental  
stage



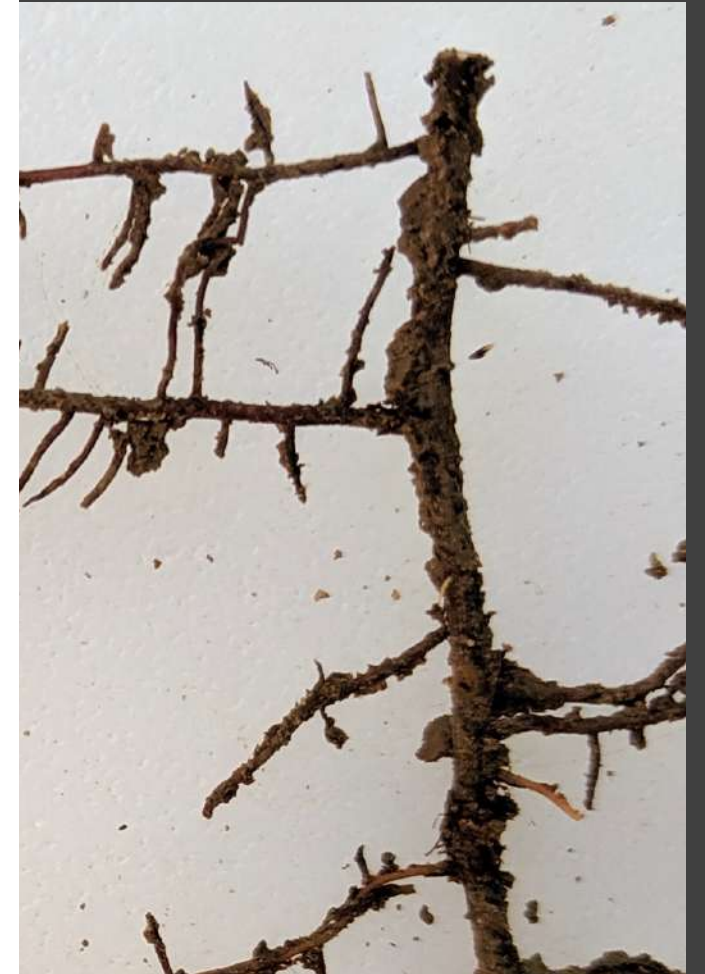
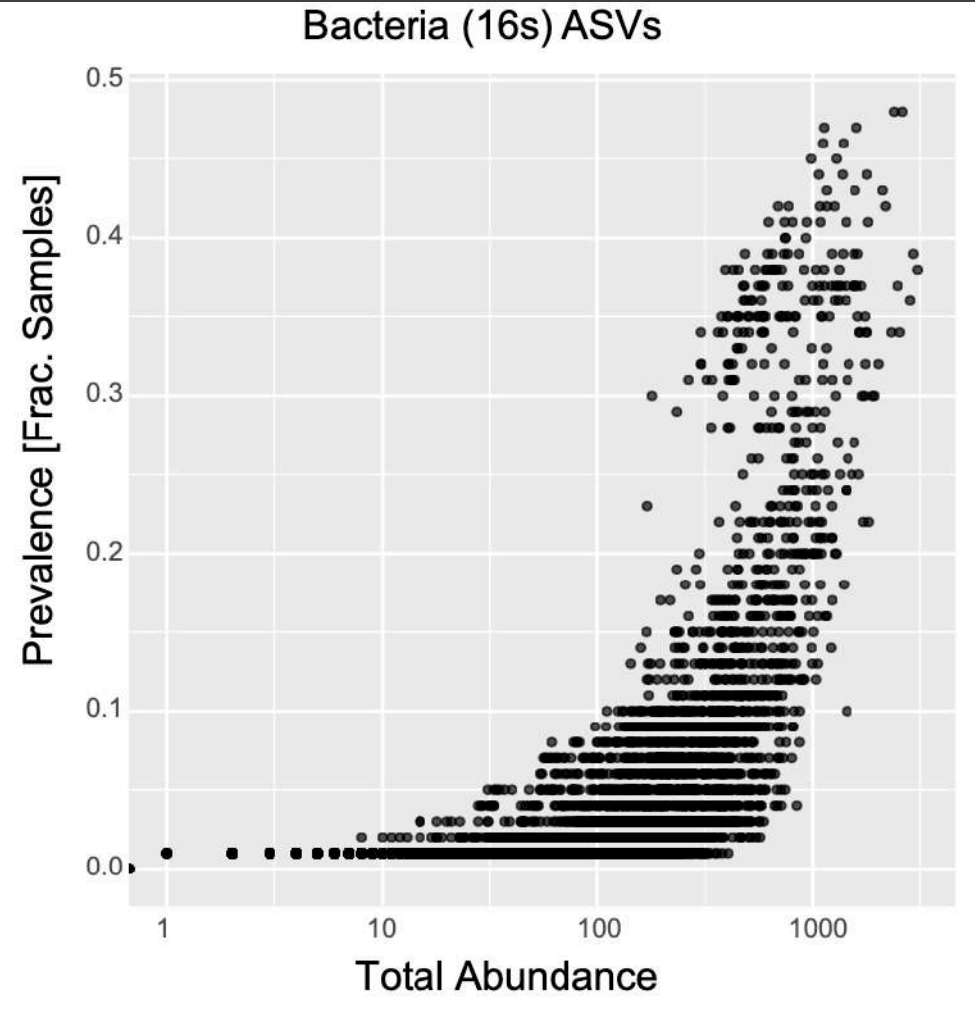
Spatial  
location

# The consequences of pooling samples





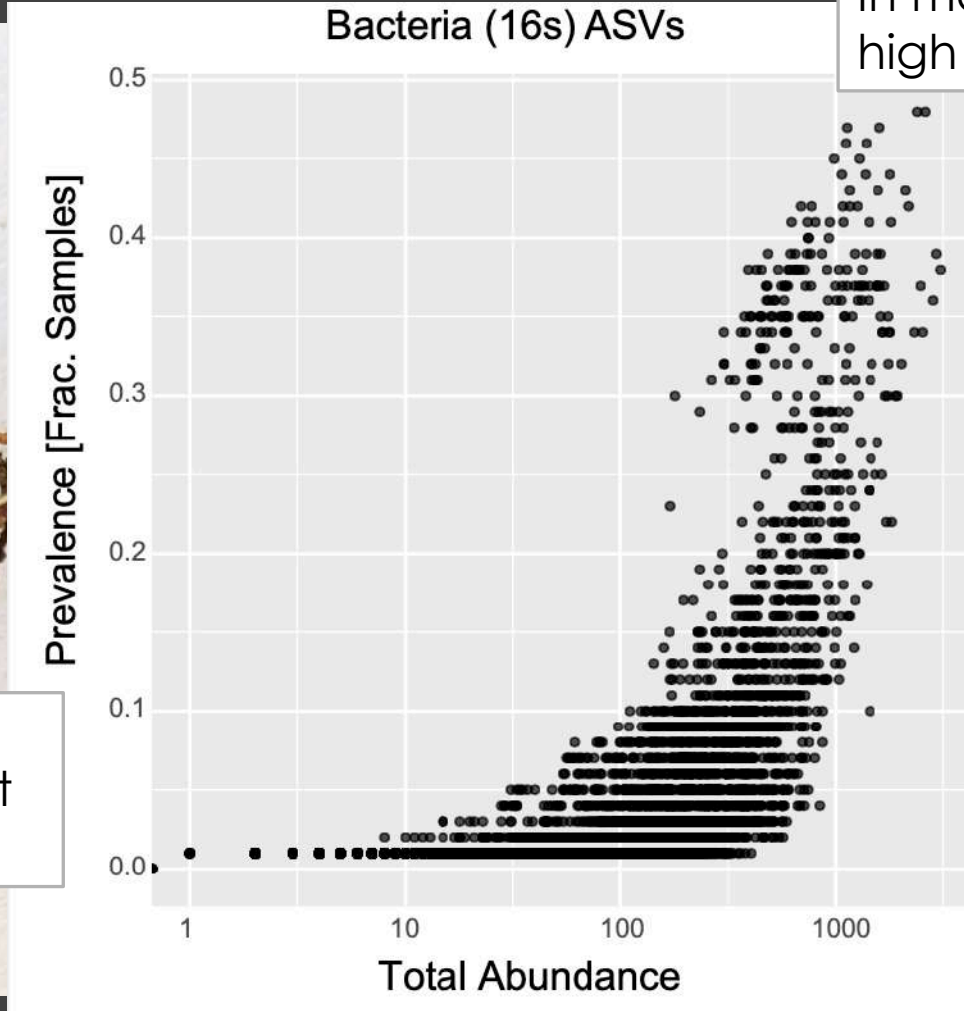
# The consequences of pooling samples



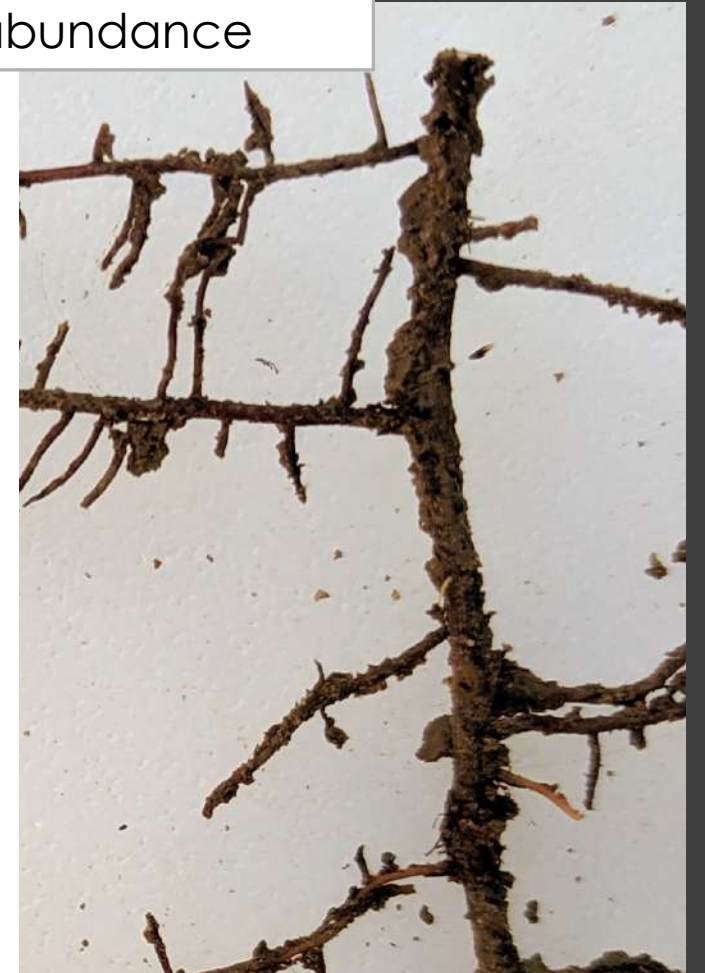
# The consequences of pooling samples



"Rare"  
In few samples at  
low abundance

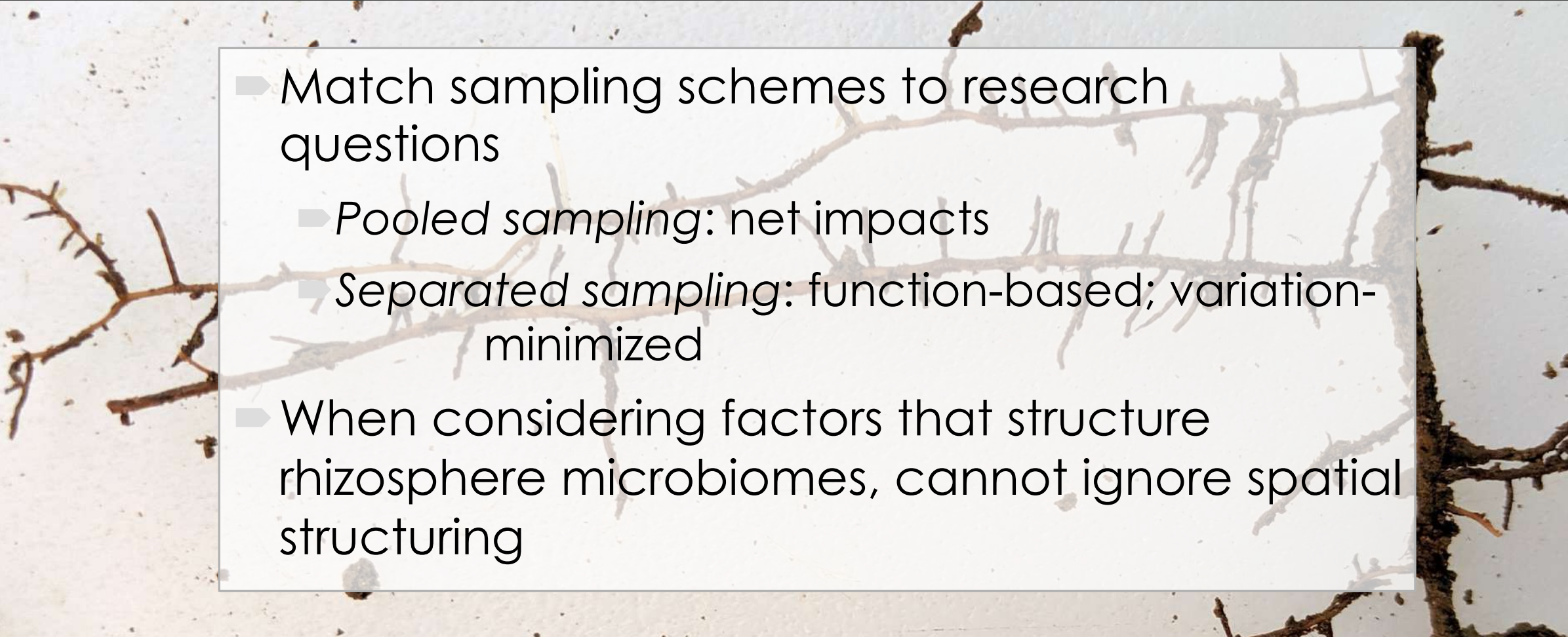


"Core"  
In many samples at  
high abundance





# How can functionally explicit sampling advance rhizosphere microbial research?

- 
- The background of the slide is a photograph showing a network of brown, fibrous plant roots extending horizontally and vertically across a light-colored, granular soil surface. The roots are thin and delicate, with some branching visible. The soil has a slightly uneven texture with small clumps and particles.
- Match sampling schemes to research questions
    - *Pooled sampling*: net impacts
    - *Separated sampling*: function-based; variation-minimized
  - When considering factors that structure rhizosphere microbiomes, cannot ignore spatial structuring



Applying these lessons in vineyards

# Applying these lessons in vineyards



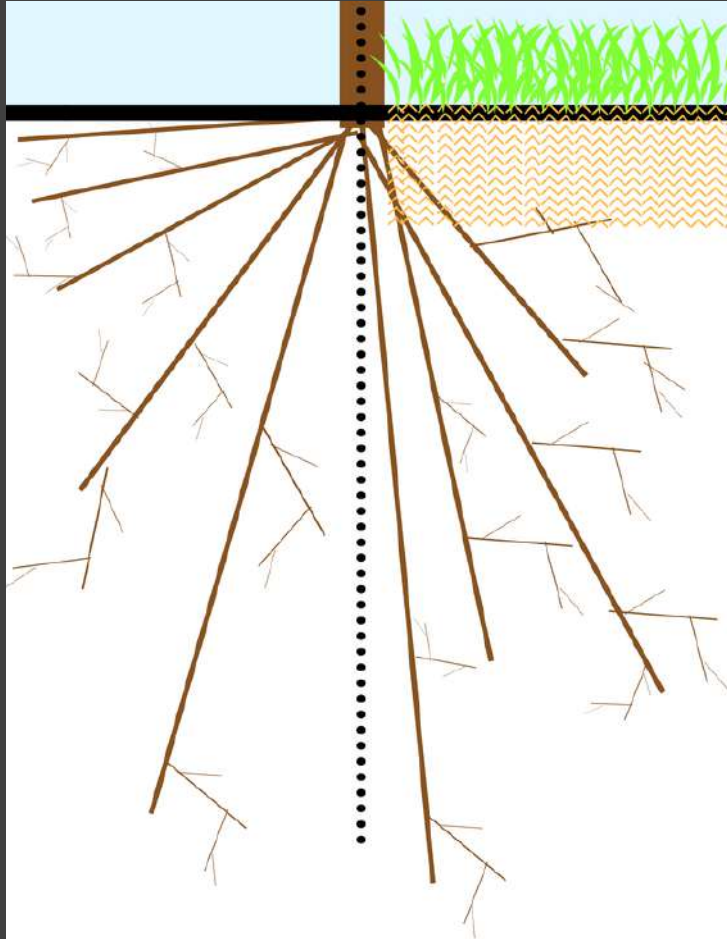
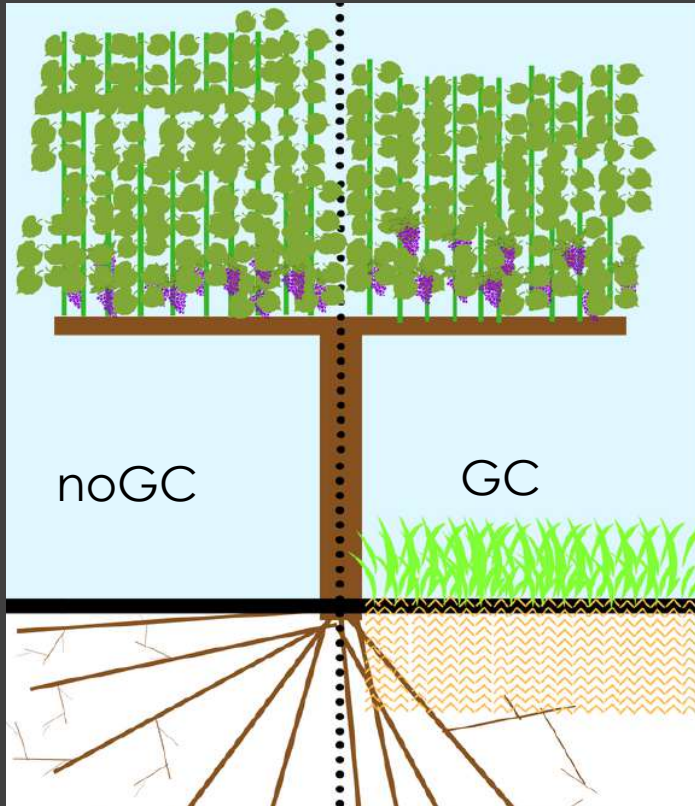


# Applying these lessons in vineyards





# Grapevines shift roots deeper in response to groundcovers



Spatial heterogeneity  
across depth

- Soil moisture ↑
- Nutrient availability ↓
- Carbon content ↓
- Microbial biomass & diversity ↓

# Study Site (2020)

- PSU vineyard at Rock Springs
- Noiret grapevines (*Vitis* hybrid)
- Groundcover treatment(fall 2016)
  - **noGC** (herbicide)
  - **GC** (Red Fescue; *Festuca rubra*)
- 32, 1-m deep root boxes





# Competition varies with depth



High competition  
0-30 cm

Moderate competition  
31-60 cm

Indirect competition  
61-100 cm



# Root distributions and sampling



High competition  
0-30 cm

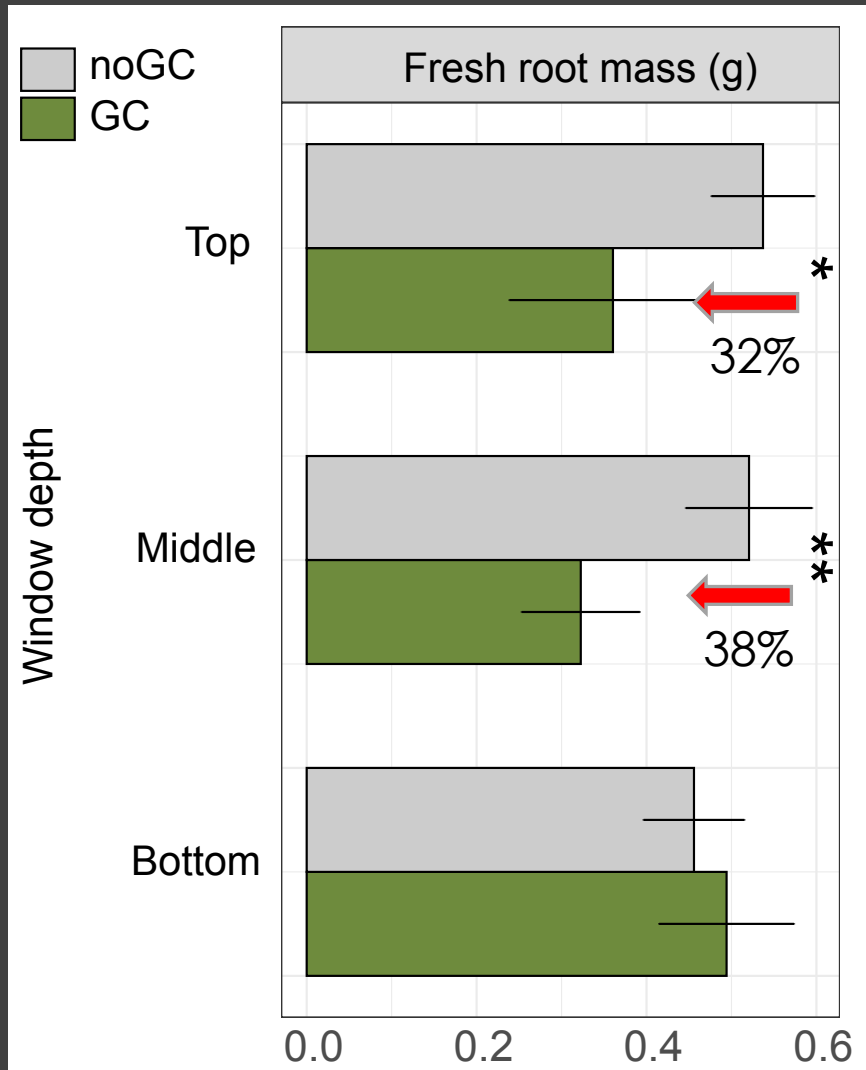
Moderate competition  
31-60 cm

Indirect competition  
61-100 cm



- Absorptive root sampling during peak root growth (July 13-17, 2020)
- Fresh root weight (g)
- Counts of new white roots at each depth
- Rhizosphere removed by vortexing in PBS buffer in the field
- Concurrent study on root transcriptomics and metabolomics

# Root systems shift deeper under GC



\* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$

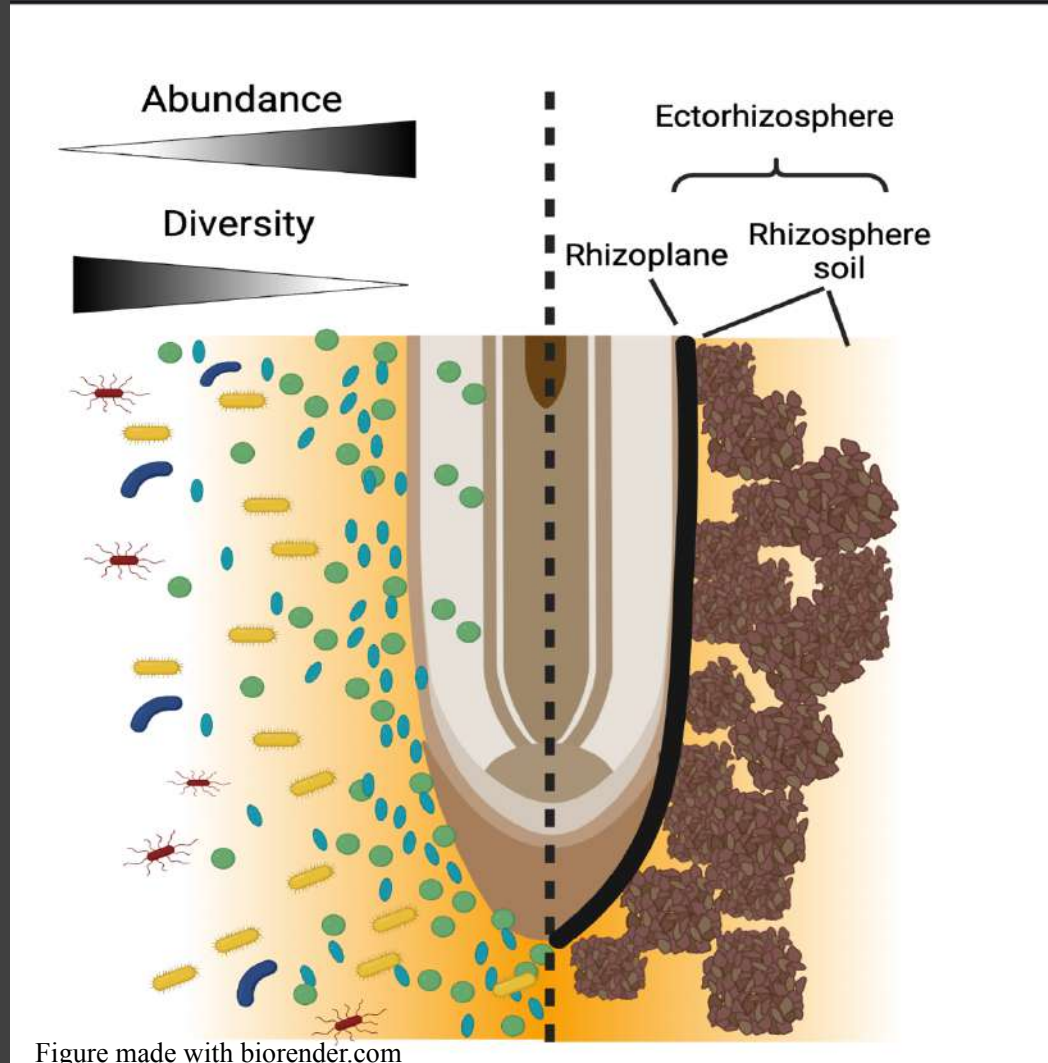
Fleishman et al. 2022, *In Prep*

Shifts beyond the groundcover rooting zone are closely related to the proportion of new roots at depth

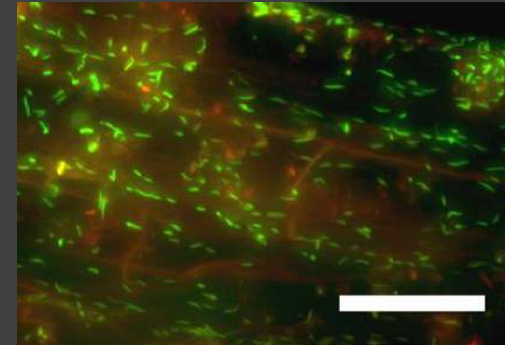
Common practices of shallow soil sampling and pooled root sampling may have obscured findings



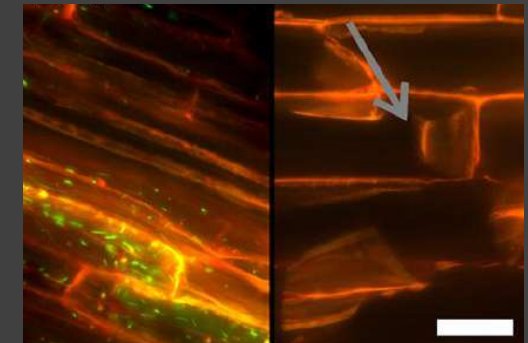
# Does the method of removing the rhizosphere matter?



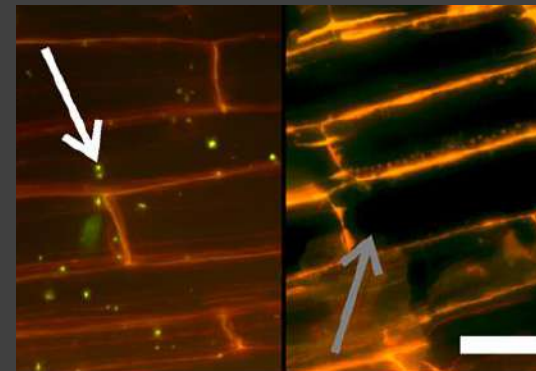
Untreated



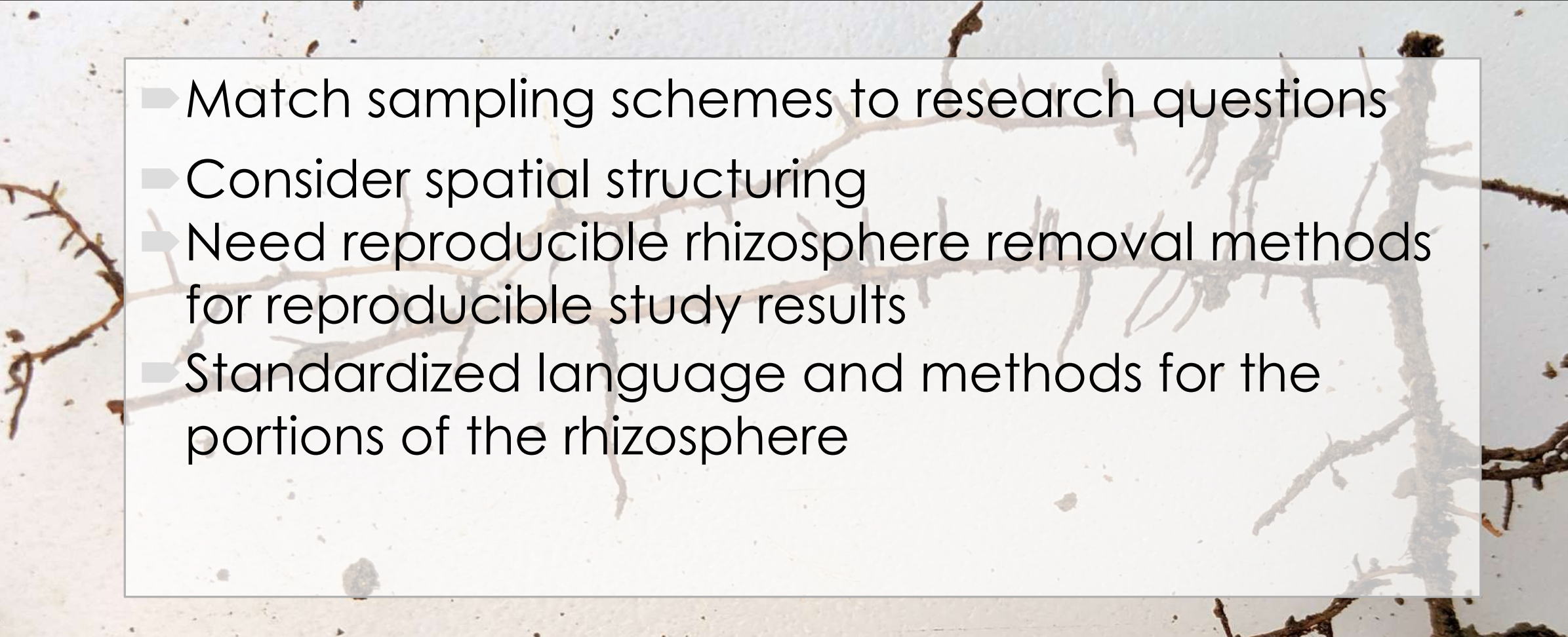
Low intensity sonication



High intensity sonication



# How can we improve rhizosphere methodologies?

- 
- A photograph showing a network of brown plant roots extending horizontally and vertically through a light-colored, granular soil. The roots are thin and branching, with some small root hairs visible. The soil has a slightly uneven texture with some small dark spots.
- Match sampling schemes to research questions
  - Consider spatial structuring
  - Need reproducible rhizosphere removal methods for reproducible study results
  - Standardized language and methods for the portions of the rhizosphere



# Acknowledgements

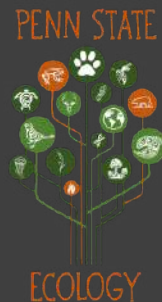
Centinari Viticulture Lab

Bell Microbiome Manipulation Lab

Eissenstat Root Ecology Lab

Don Smith, Jeremy Harper, and Hort Farm Staff

Hayden Bock, Jing Guo, Ryan Trexler



**Suzanne Fleishman, [suf44@psu.edu](mailto:suf44@psu.edu)**

**[centinarilab.squarespace.com/publications](https://centinarilab.squarespace.com/publications)**

King, W.L., et al., 2021. The hierarchy of root branching order determines bacterial composition, microbial carrying capacity and microbial filtering. Commun Biol 4, 483. <https://doi.org/10.1038/s42003-021-01988-4>



# Penn State Microbiome Center

- Established in 2016
  - 80 Research Groups from 10 different colleges and institutes (arts and anthropology to zygotes and zoology; human, animal, plant, environmental health)
  - 40 weekly seminars and 2-3 workshops per semester
  - Kick Start Program for beginners
  - Data Analysis Working Group (DAWG) led by students and postdocs
  - Industrial programming and partnerships (contact [CaroleeBull@psu.edu](mailto:CaroleeBull@psu.edu) for information)
  - Formal Microbiome Sciences Dual Title being proposed
  - NEW SENIOR LEVEL DIRECTOR being recruited
- 
- Visit our website: [microbiome.psu.edu](http://microbiome.psu.edu)
  - Follow us on Twitter @PSUmBiome

