

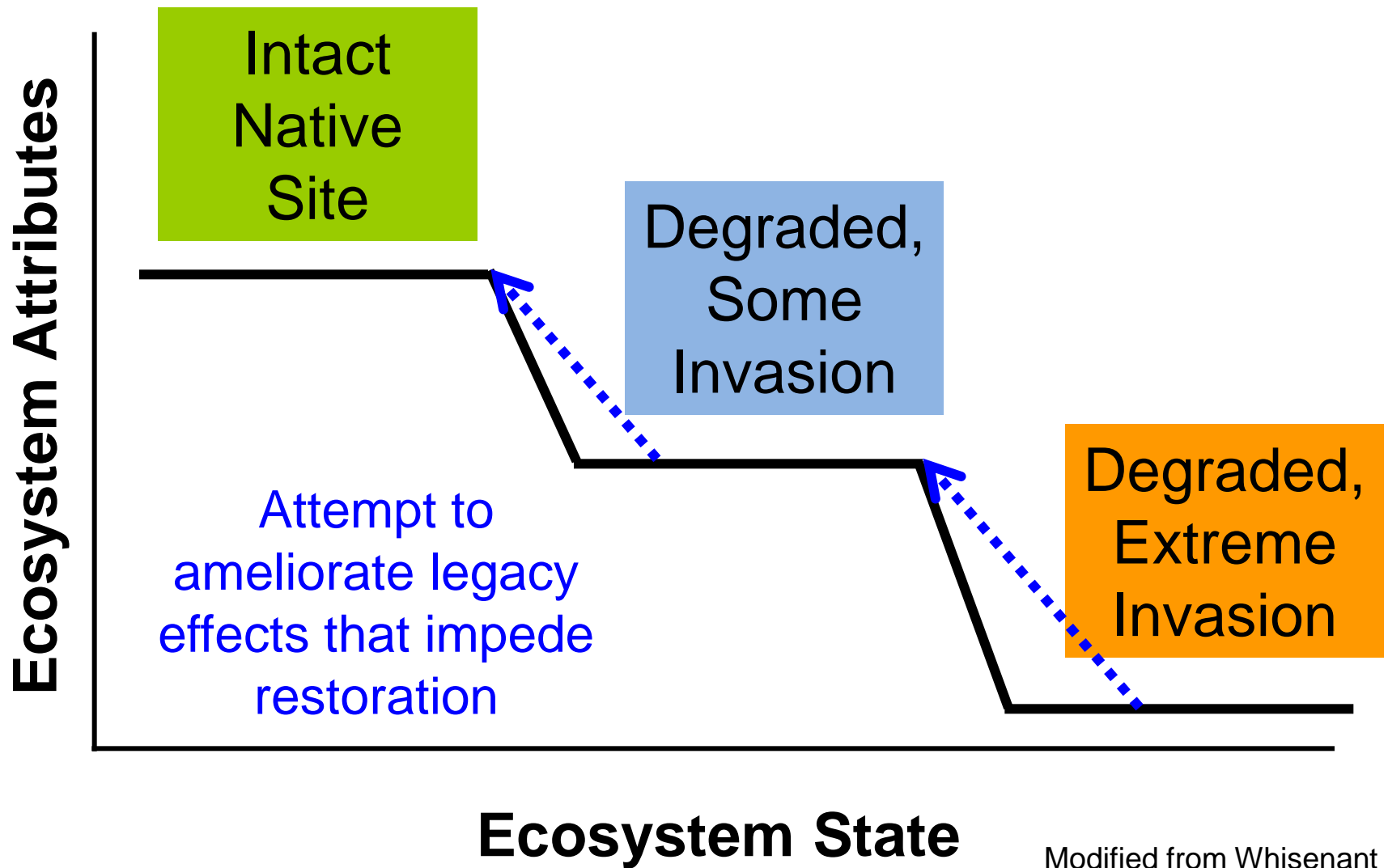
# Understanding and mitigating soil legacies to improve restoration success

Christine V. Hawkes

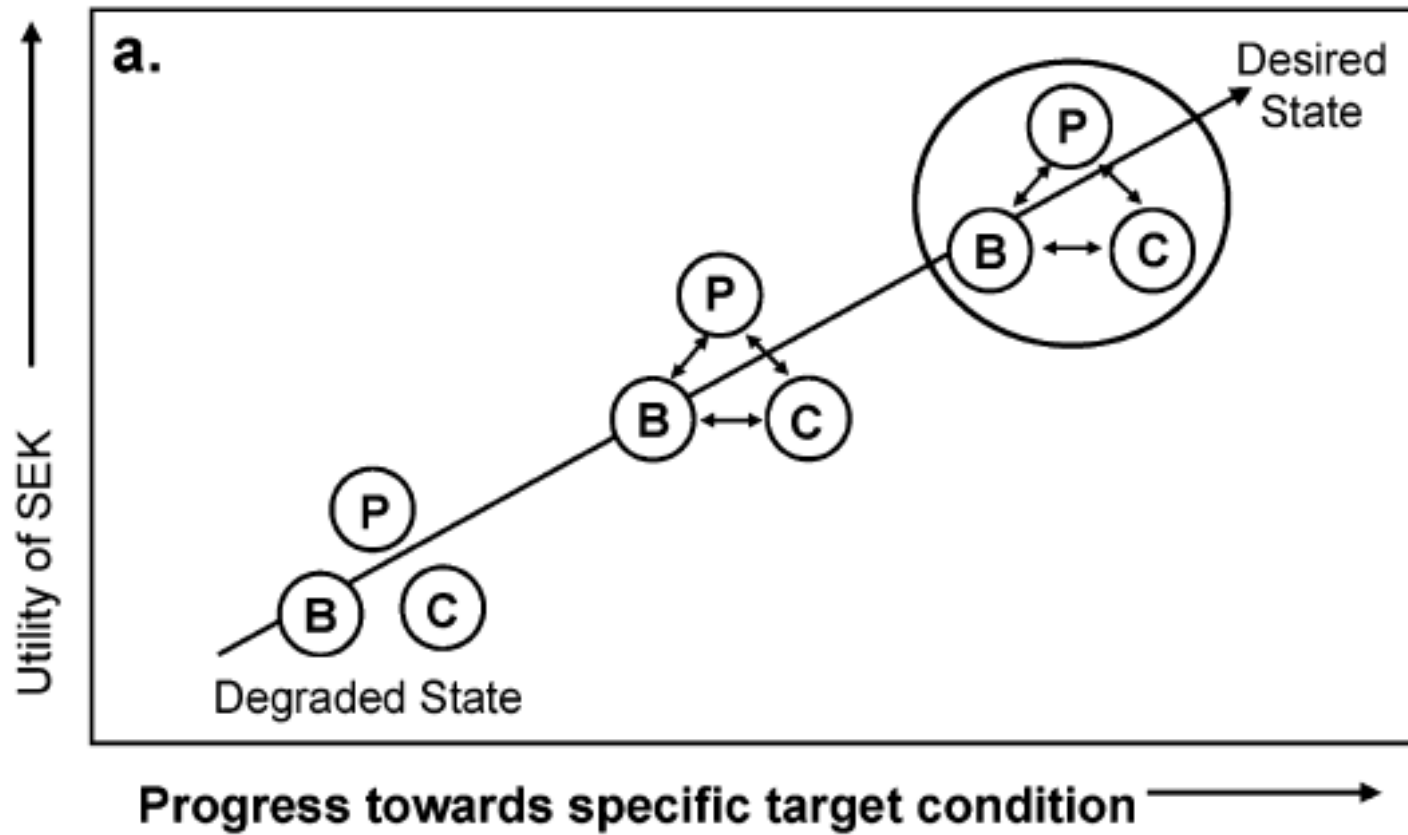
University of Texas at Austin



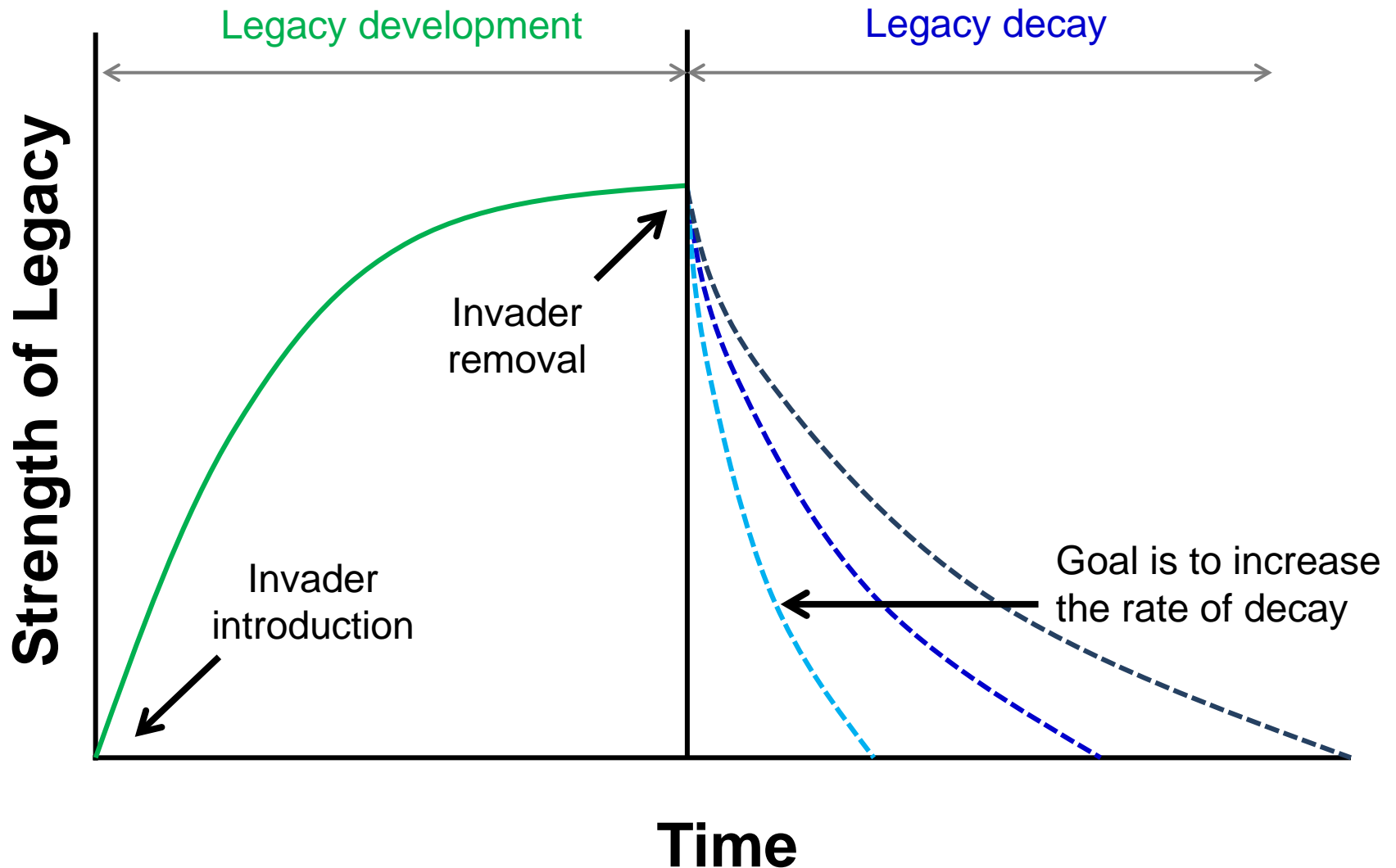
# Targeting legacies in restoration

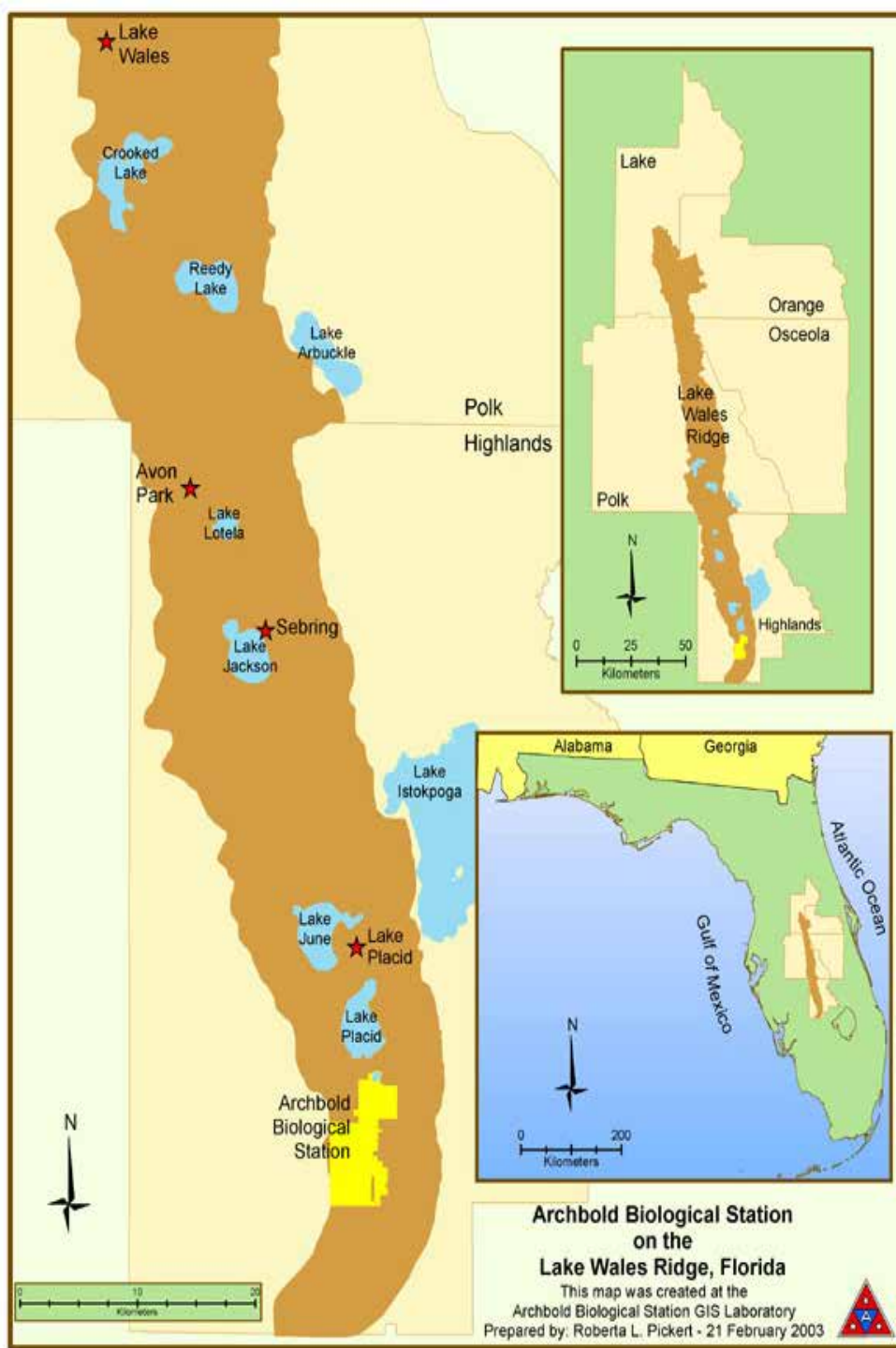


# Need to target legacy effects explicitly



# Can we change rate of legacy decay?





Native Scrub



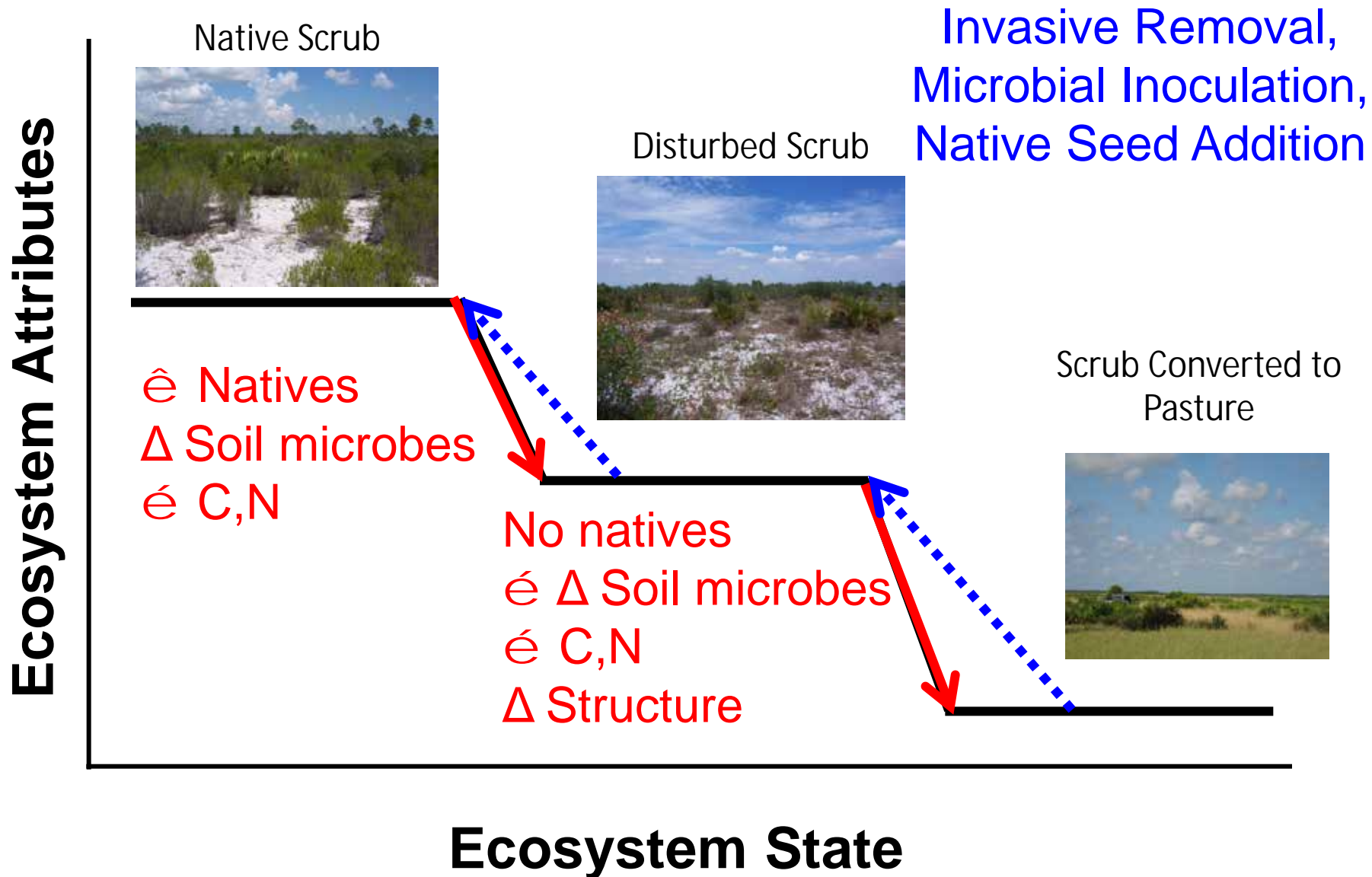
Disturbed Scrub



Scrub Converted to Pasture



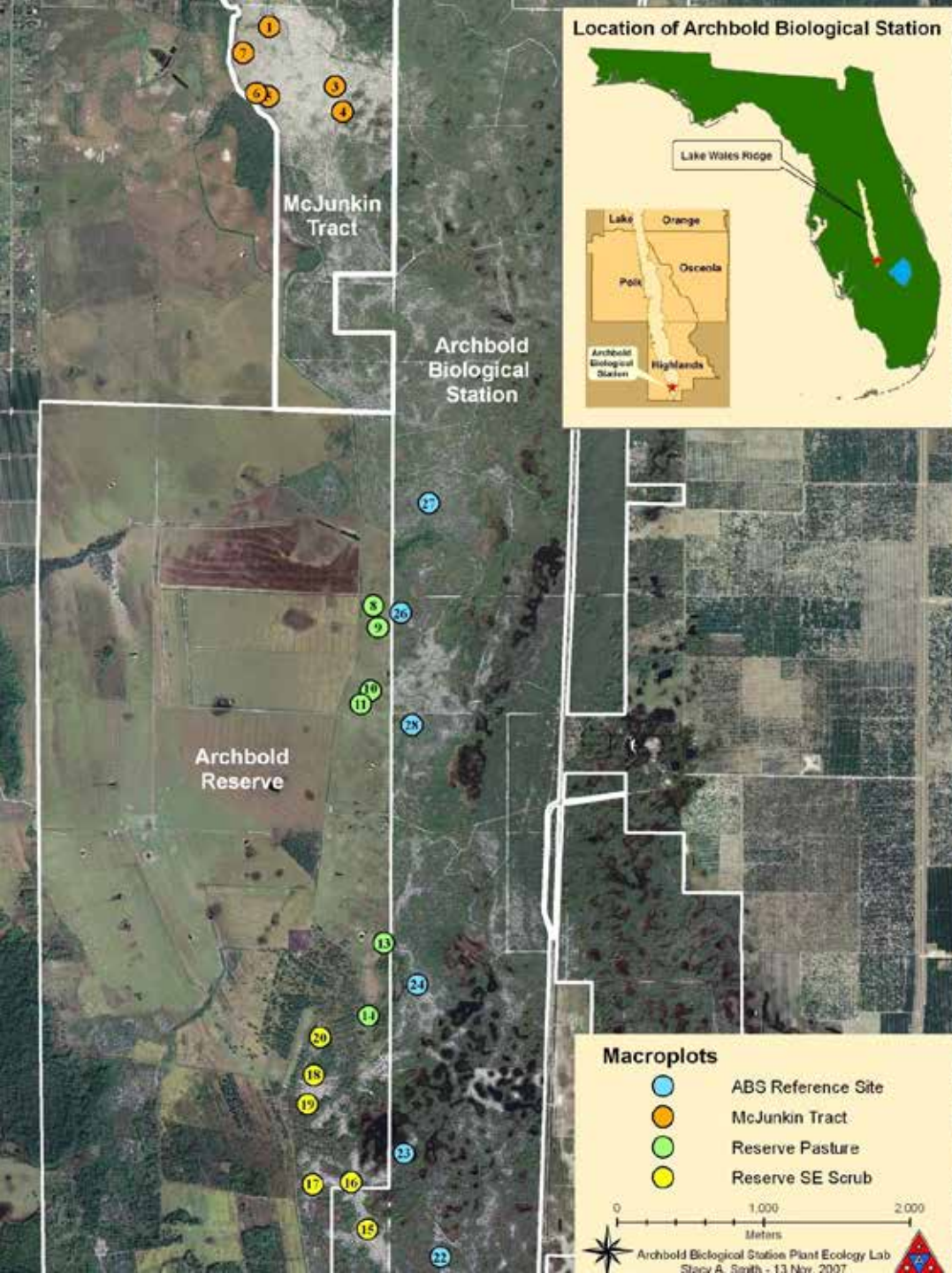
# Targeting legacies in restoration



# Restoration community & ecosystem goals

- Mitigate structural, biogeochemical, and microbial legacies
- Initiate self-sustaining native populations





Restoration sites  
-disturbed  
-pasture

Native sites  
-undisturbed  
scrub patches

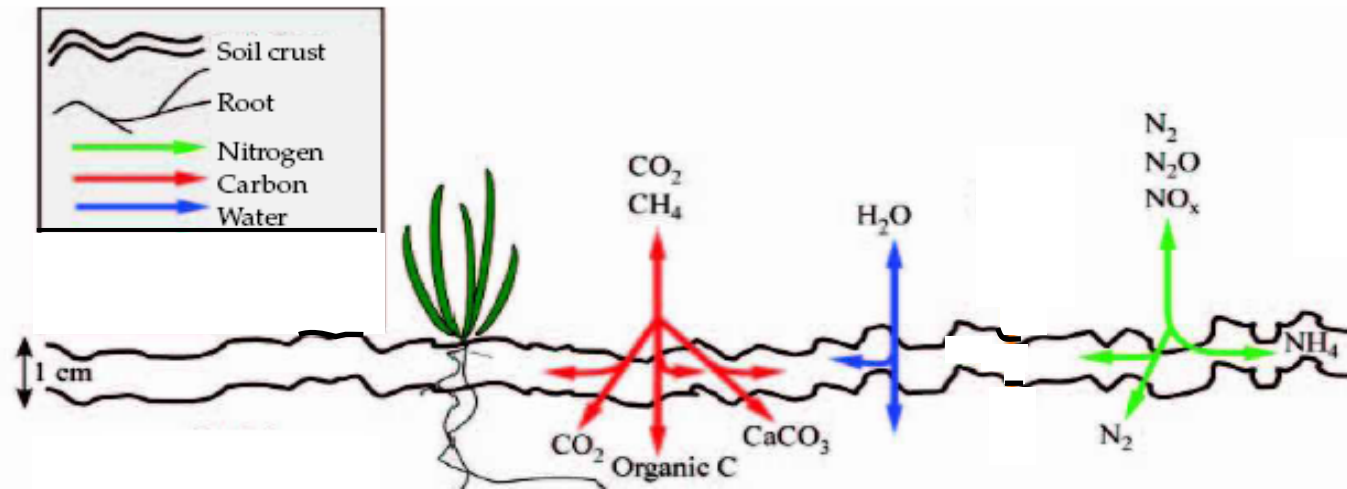


# Restoration treatments: herbicide



Credit: Archbold Biological Station

# Restoration treatments: soil microbial addition





# Restoration experiment

Native Scrub



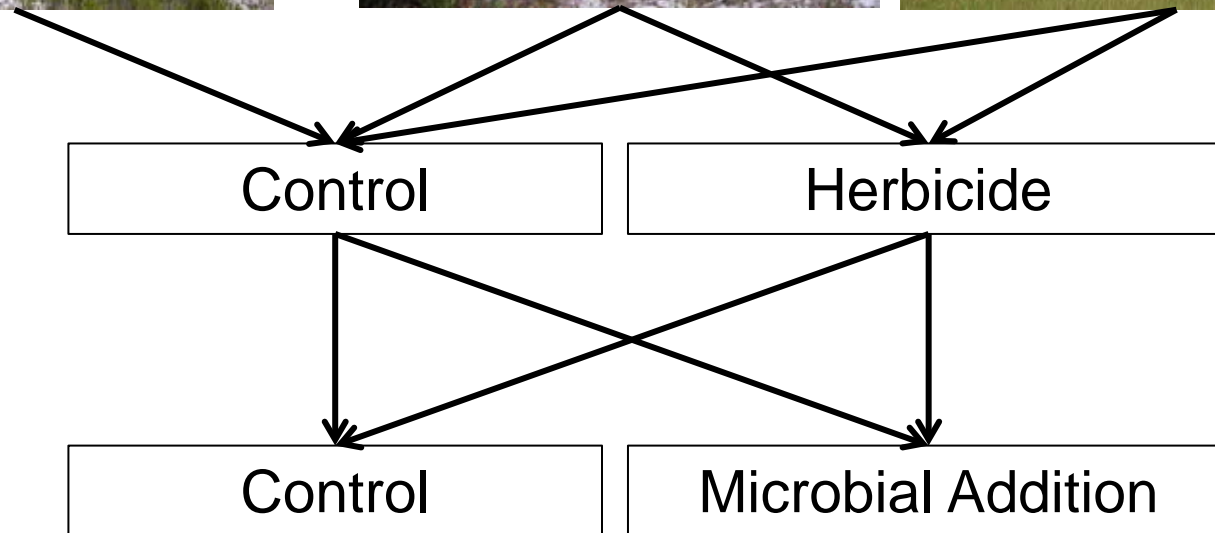
Disturbed Scrub



Pasture



Plots were set  
up in fall 2006  
and tracked  
through  
spring 2010.



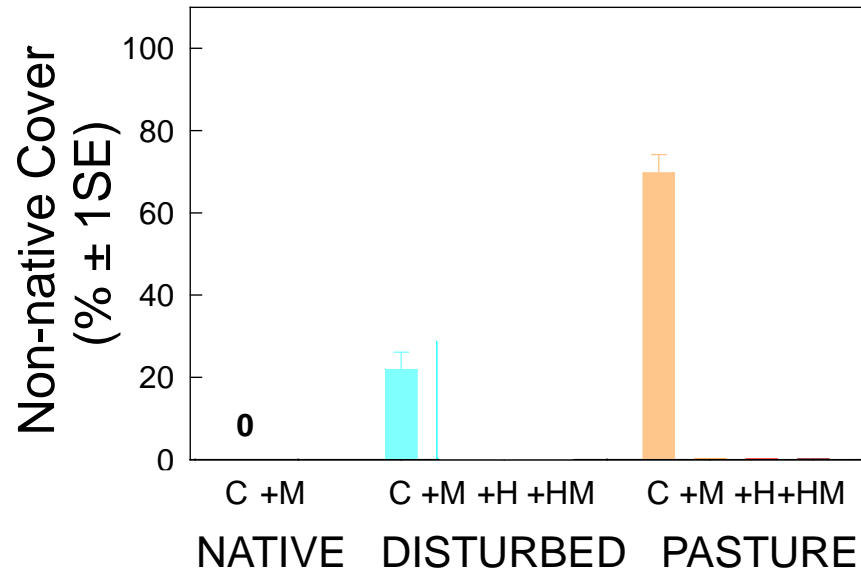
# Measuring restoration success

- Return site characteristics?
- Reduce nutrient legacies?
- Improve plant recovery?

# Measuring restoration success

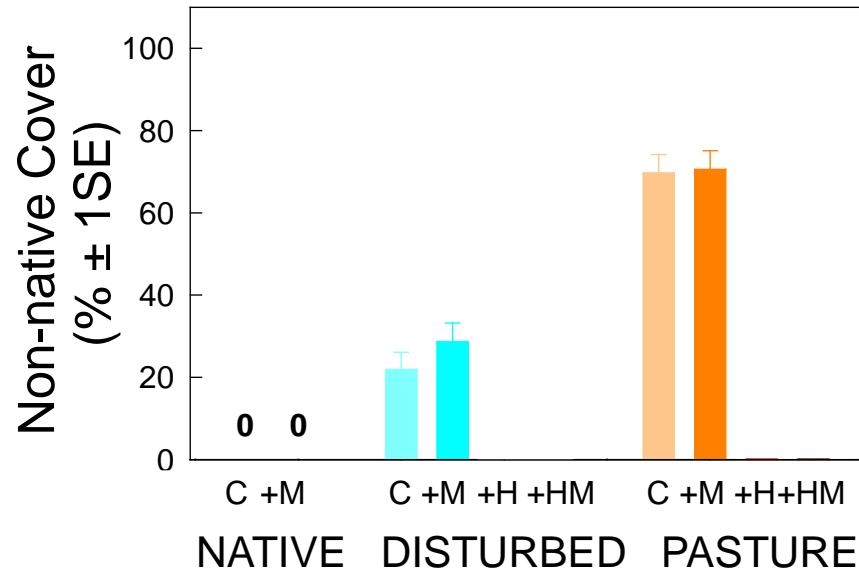
- Return site characteristics?
- Reduce nutrient legacies?
- Improve plant recovery?

# Success on some (visible) fronts: plant removals largely restore structure

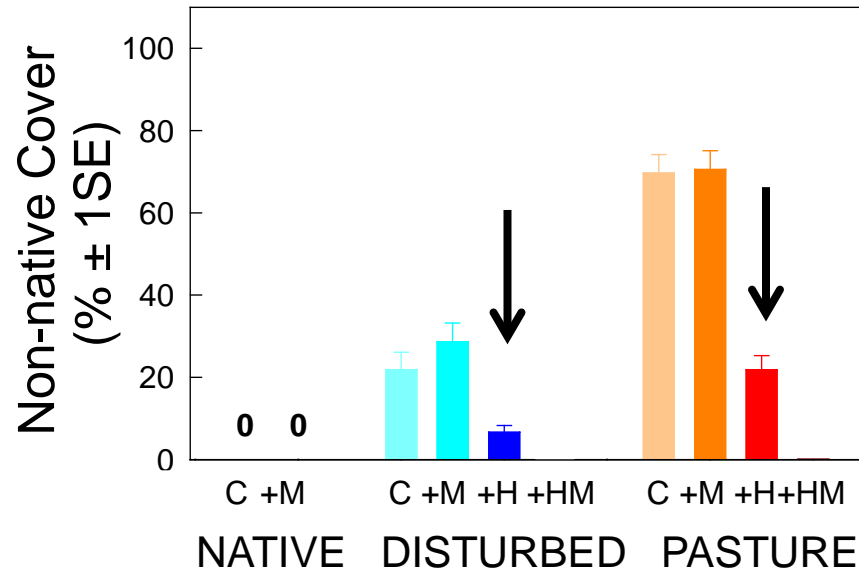




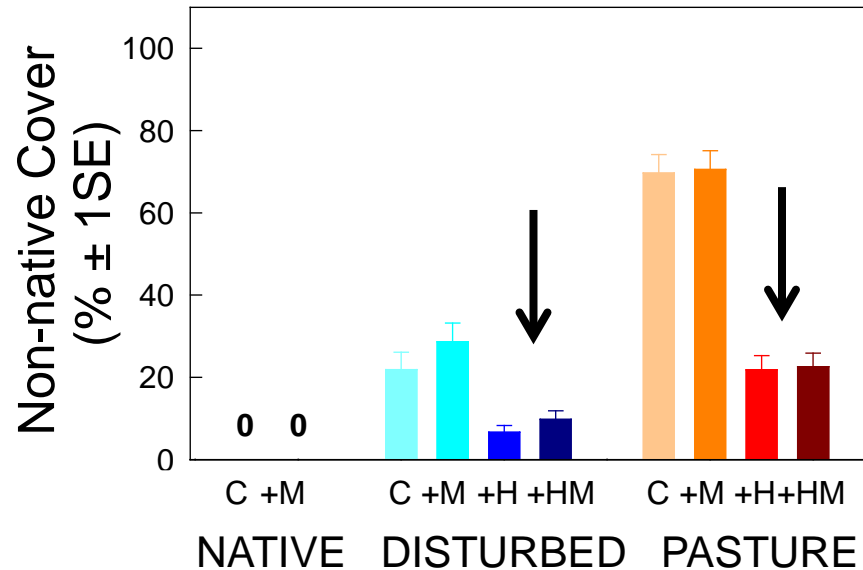
# Success on some (visible) fronts: plant removals largely restore structure



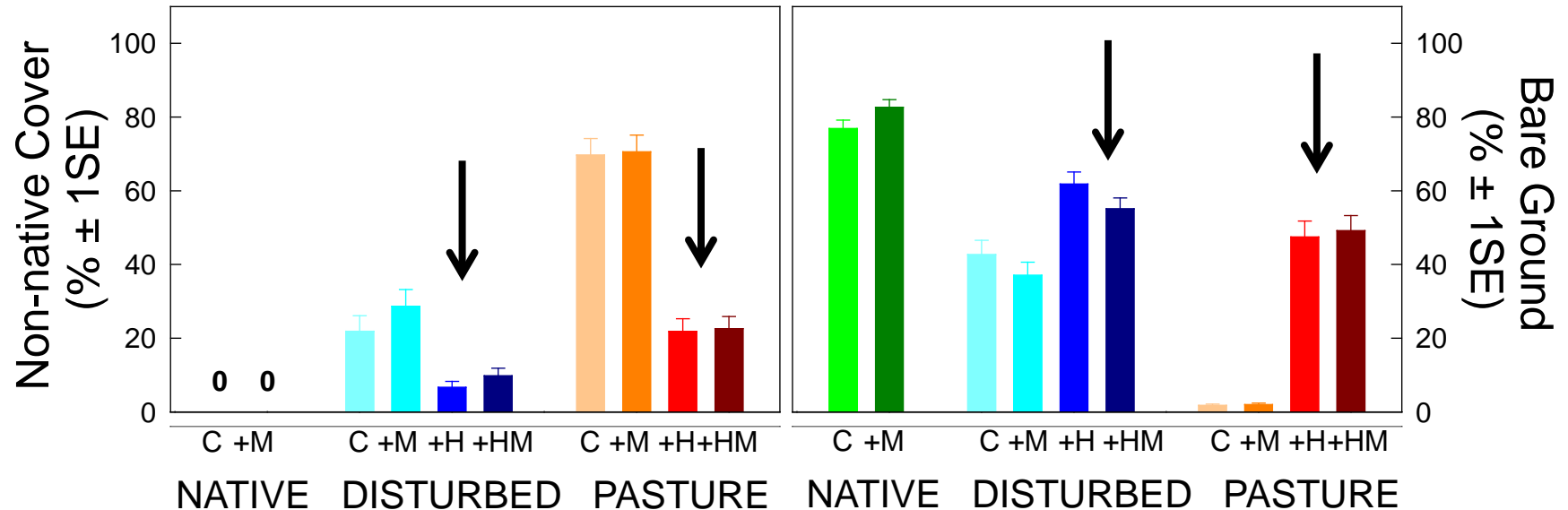
# Success on some (visible) fronts: plant removals largely restore structure



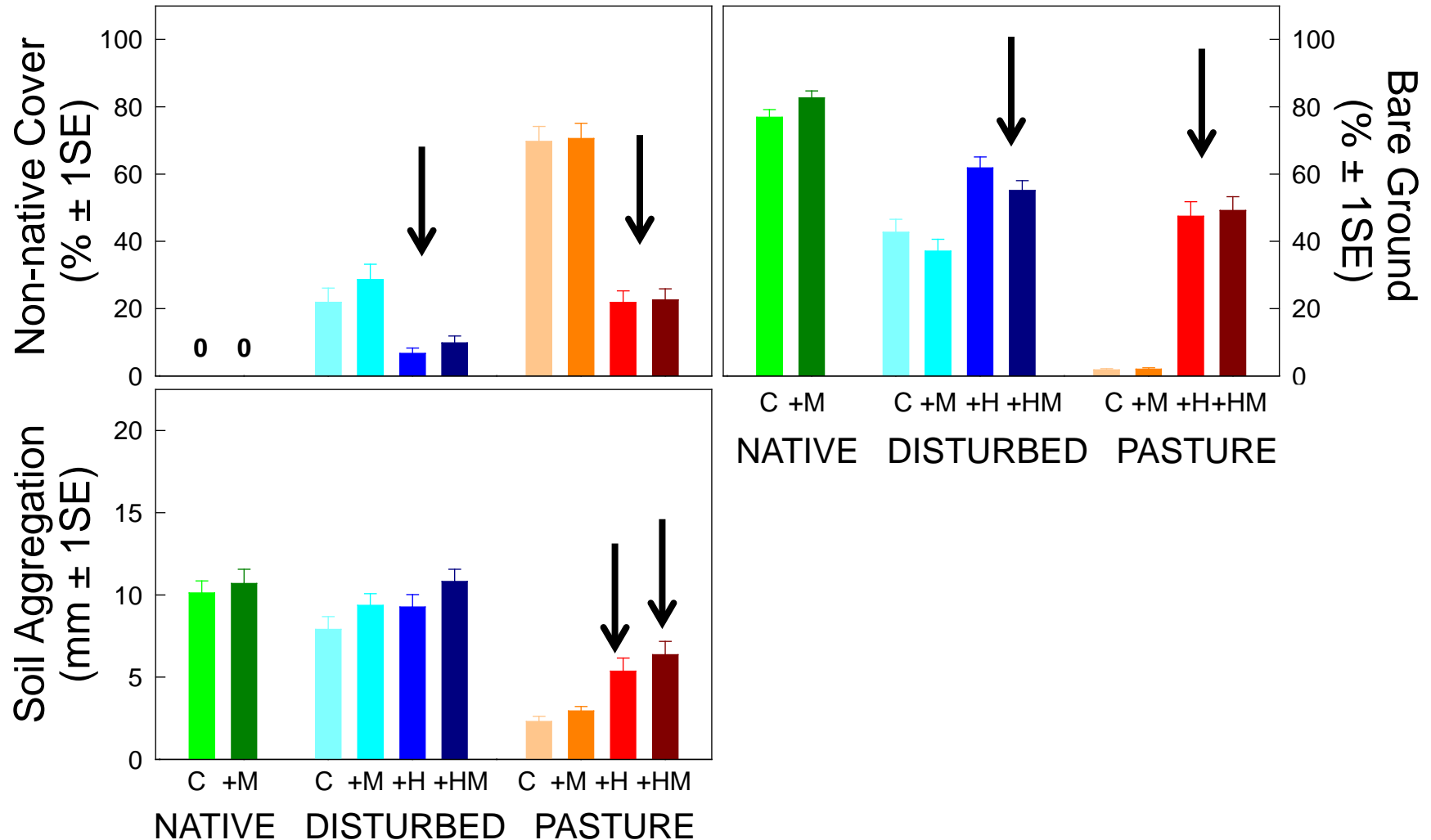
# Success on some (visible) fronts: plant removals largely restore structure



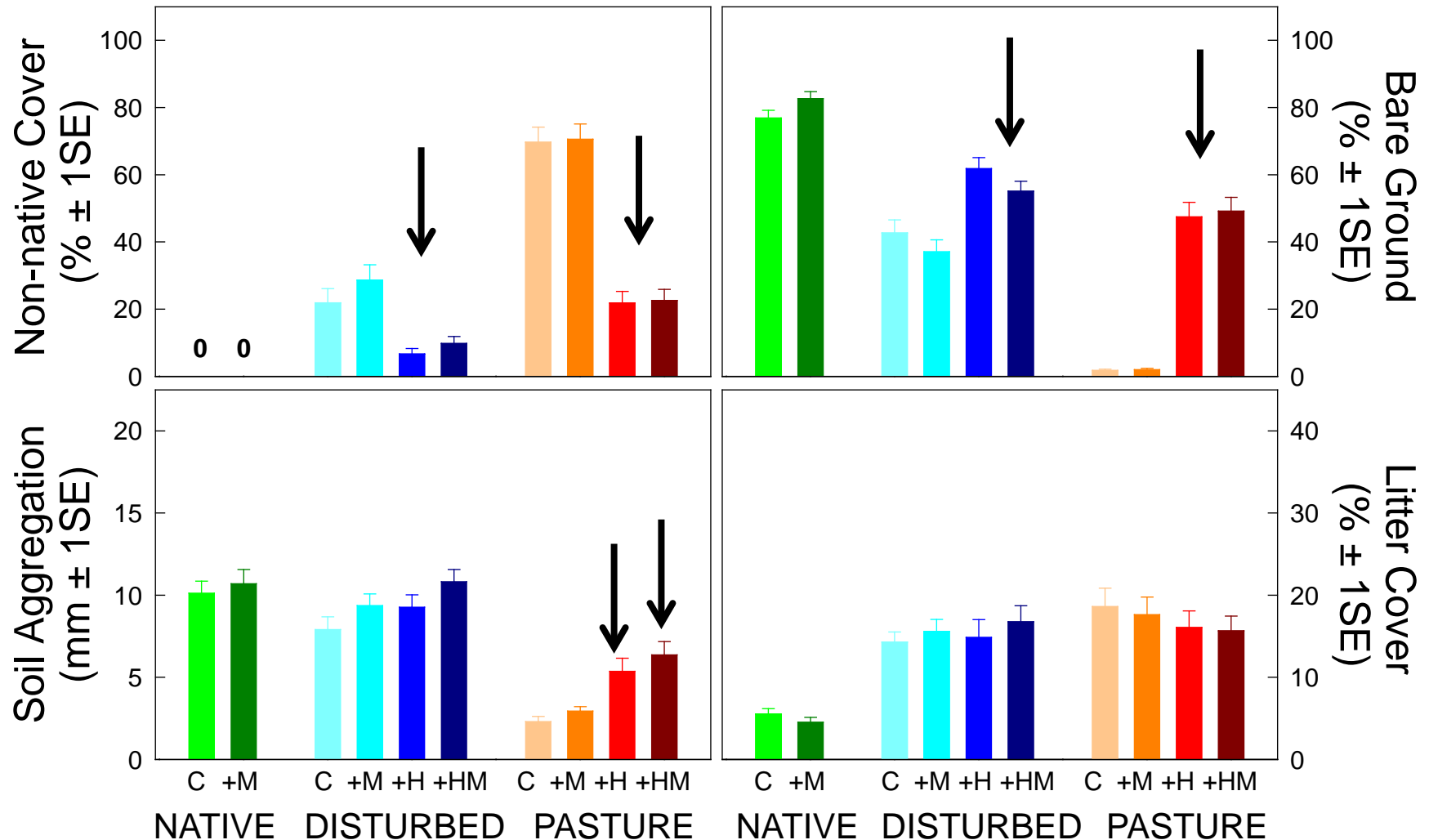
# Success on some (visible) fronts: plant removals largely restore structure



# Success on some (visible) fronts: plant removals largely restore structure



# Success on some (visible) fronts: plant removals largely restore structure





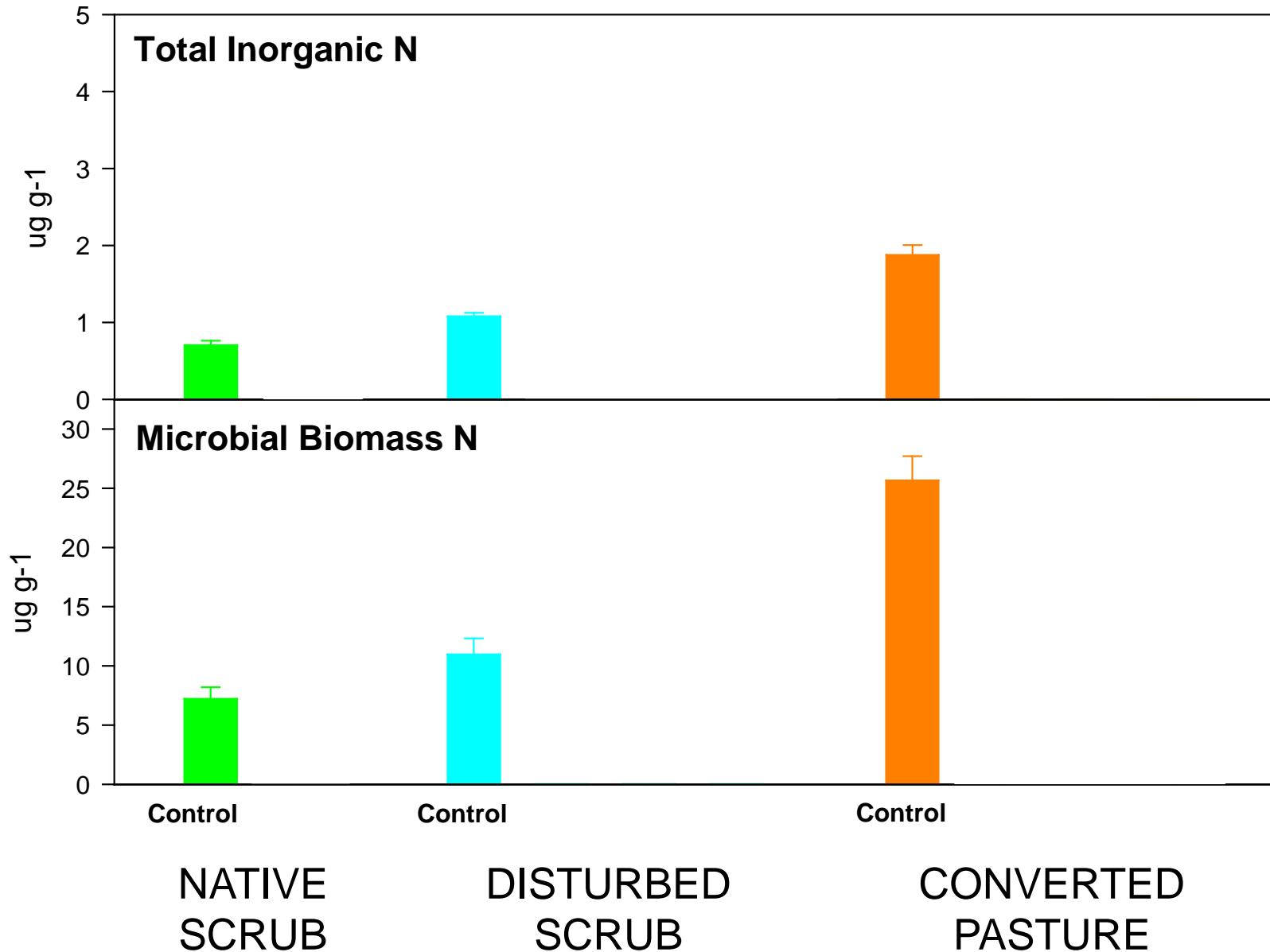
# Measuring restoration success

- Return site characteristics?
  - Yes, site cover, openness, & soil crust aggregation were improved in degraded sites
- Reduce nutrient legacies?
- Improve plant recovery?

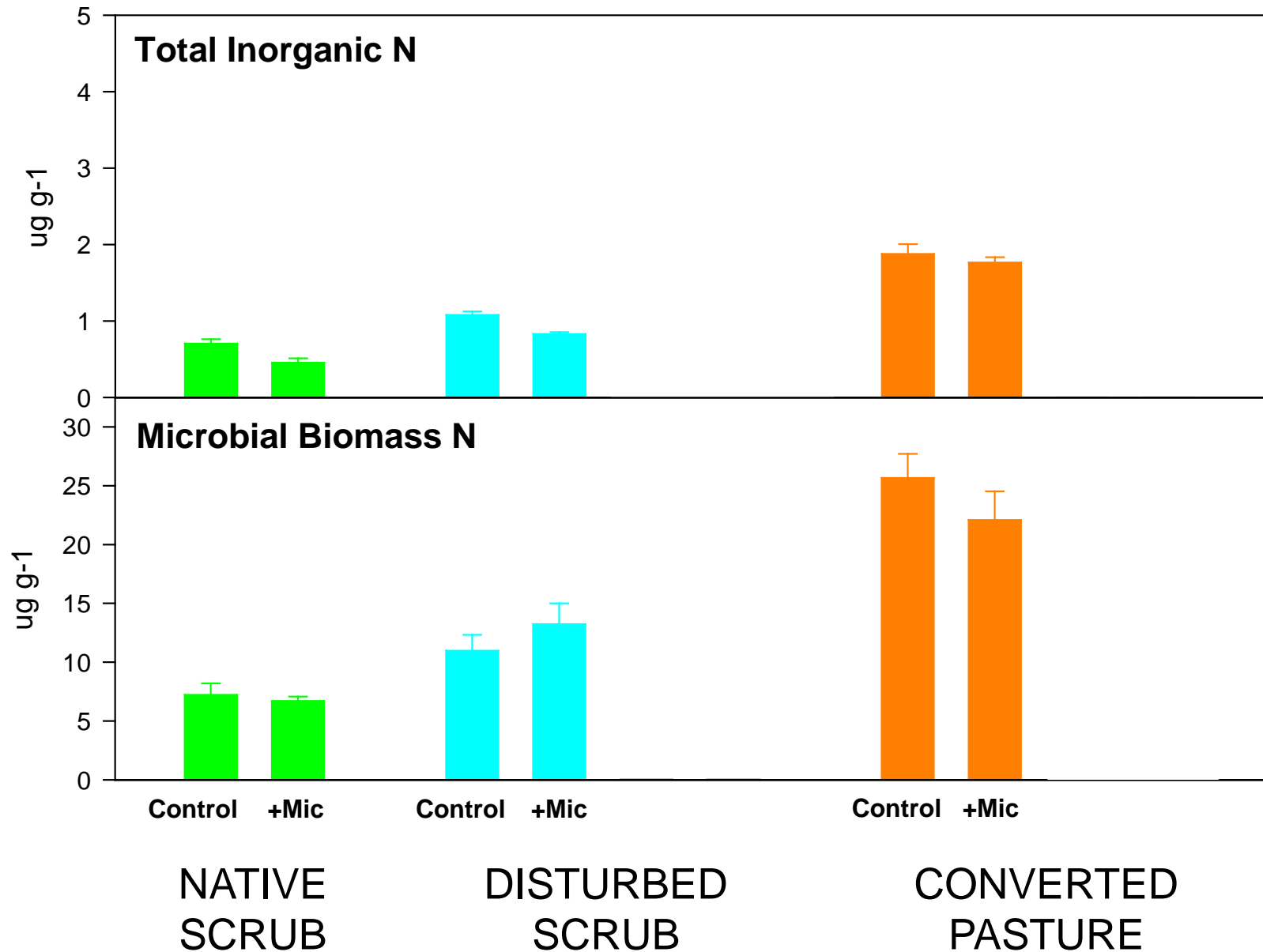
# Measuring restoration success

- Return site characteristics?
  - Yes, degraded site cover, openness, & soil crust aggregation were improved
- Reduce nutrient legacies?
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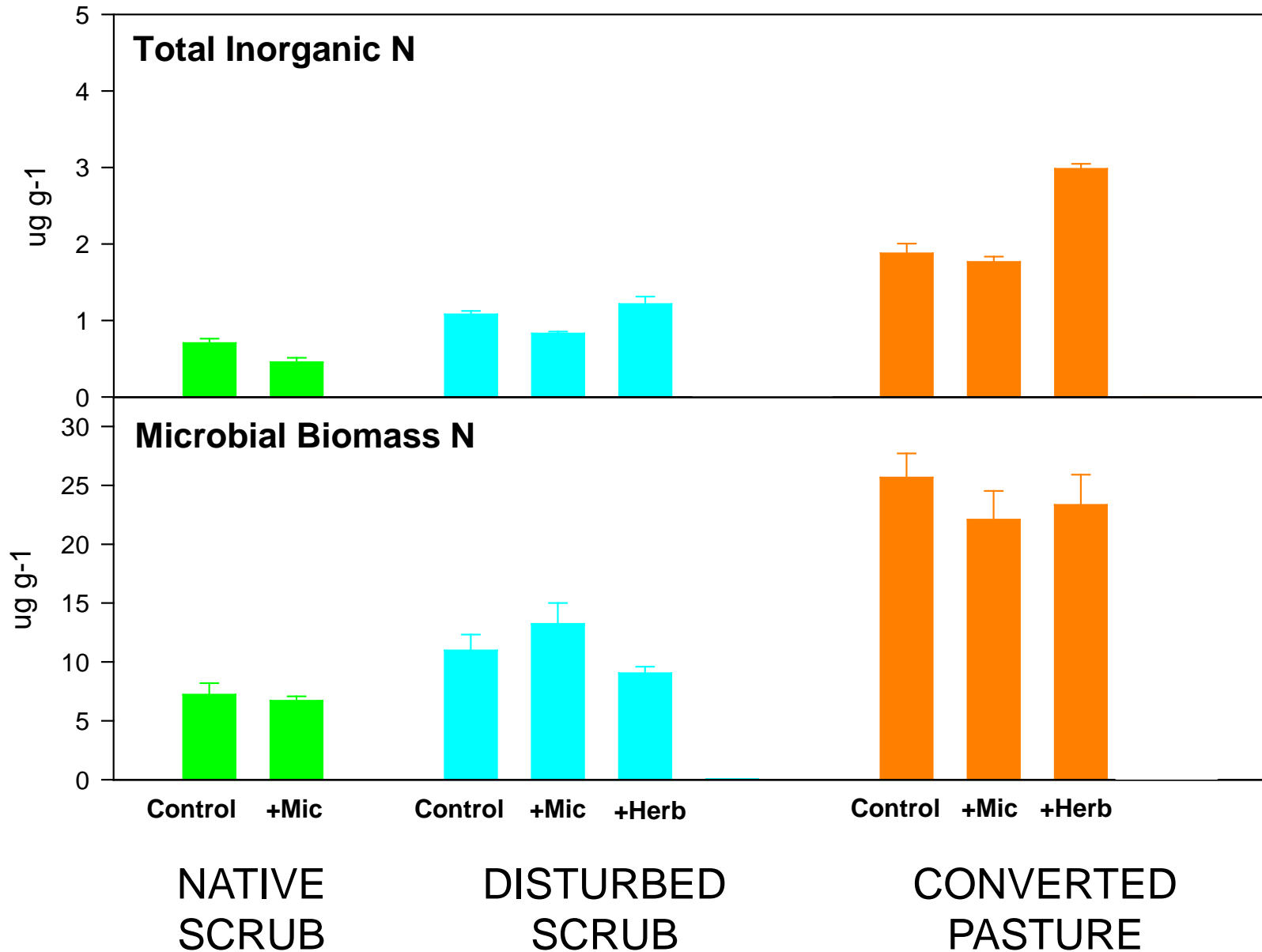
# Soil N differs among veg types



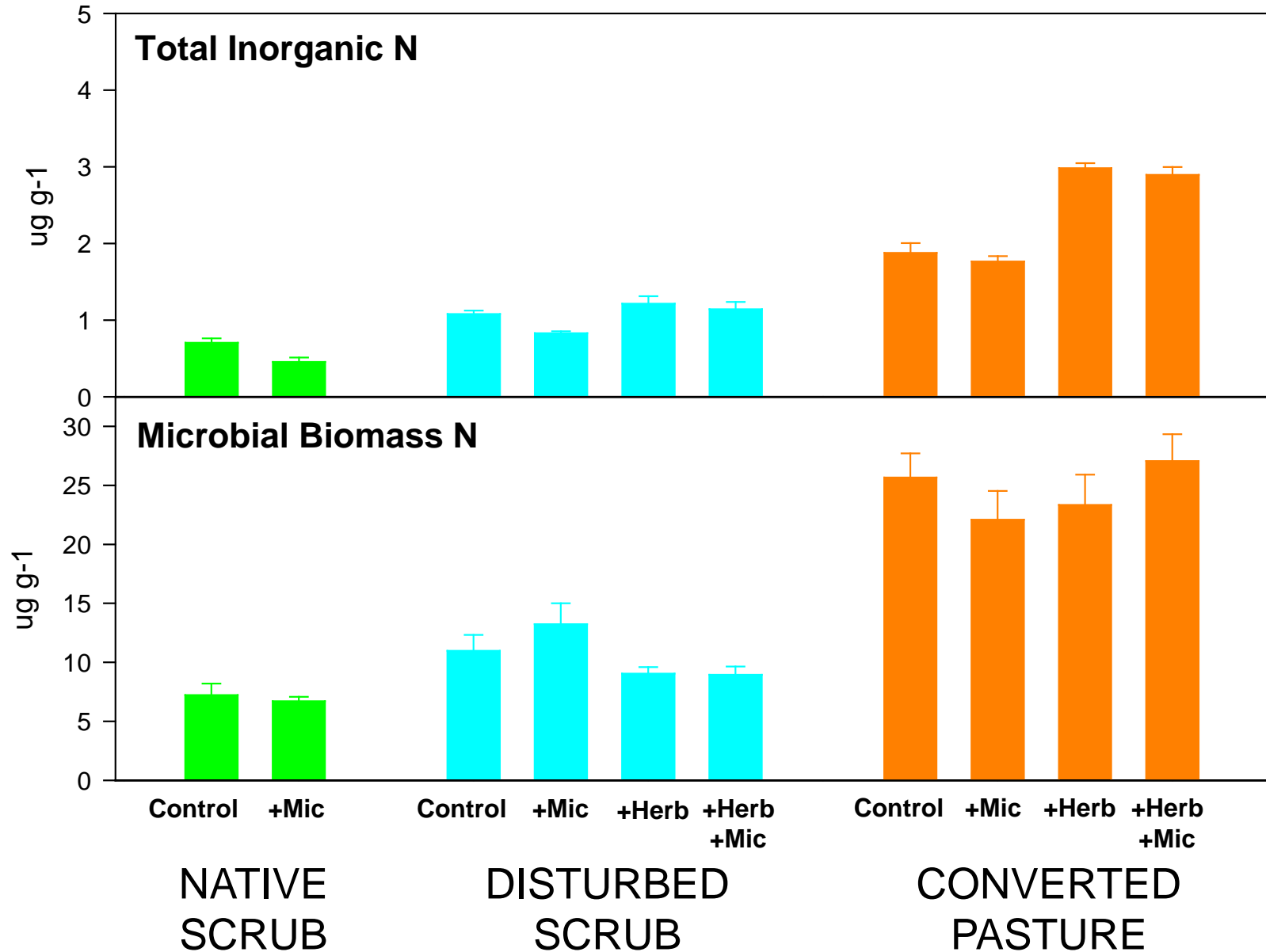
# Microbial additions have little effect on N



# High N persists after veg removal



# Treatments do not change N legacies





# Measuring restoration success

- Return site characteristics?
  - Yes, degraded site cover, openness, & soil crust aggregation were improved.
- Reduce nutrient legacies?
  - No. May need more time. Possibly remove topsoil if N reduction is necessary.
- Improve plant recovery?

# Measuring restoration success

- Return site characteristics?
  - Yes, degraded site cover, openness, & soil crust aggregation were improved.
- Reduce nutrient legacies?
  - No. May need more time. Possibly remove topsoil if N reduction is necessary.
- Improve plant recovery?

# The cast of characters

*Hypericum cumulicola*\*\*\*

*Eryngium cuneifolium*\*\*\*

*Lechea cernua*\*

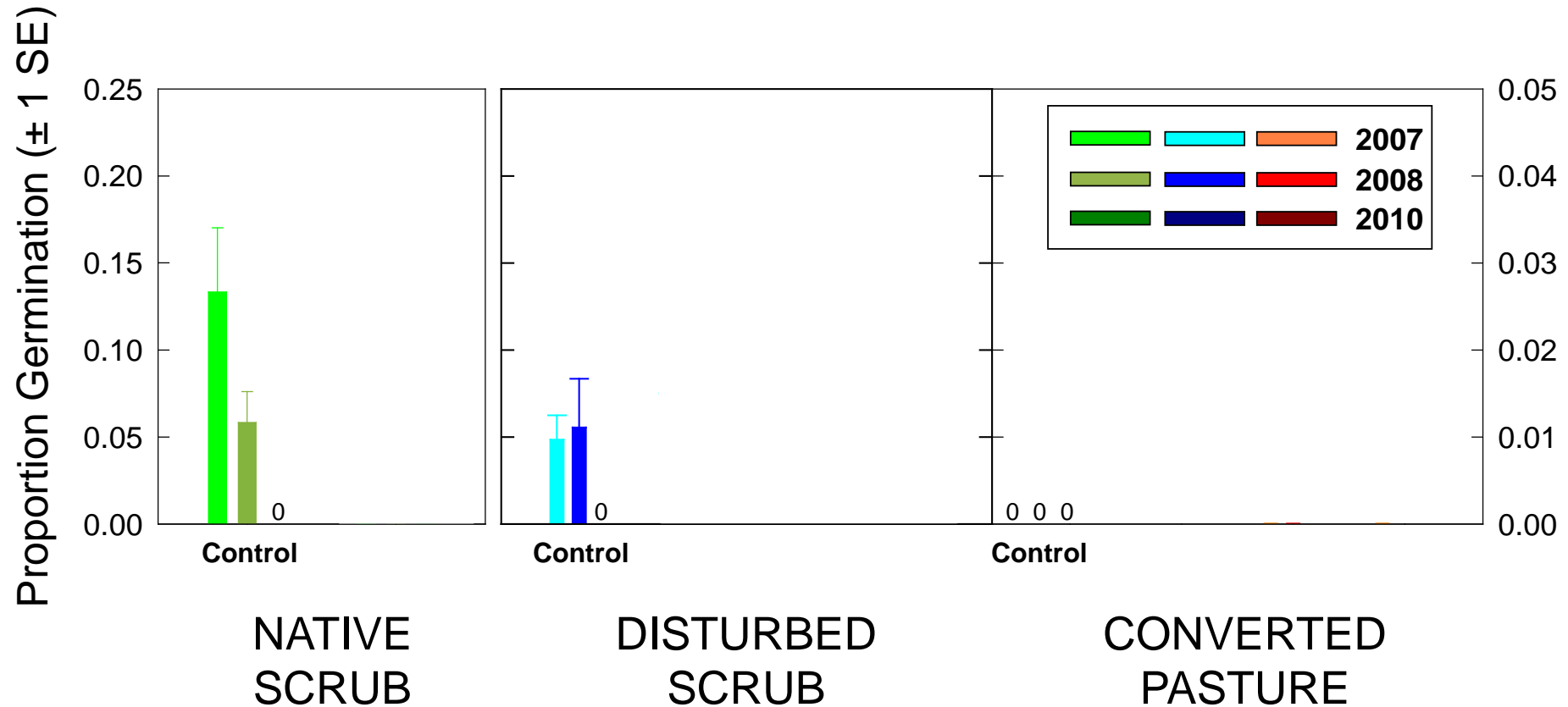
*Lechea deckertii*

*Polygonella basiramia*\*\*\*

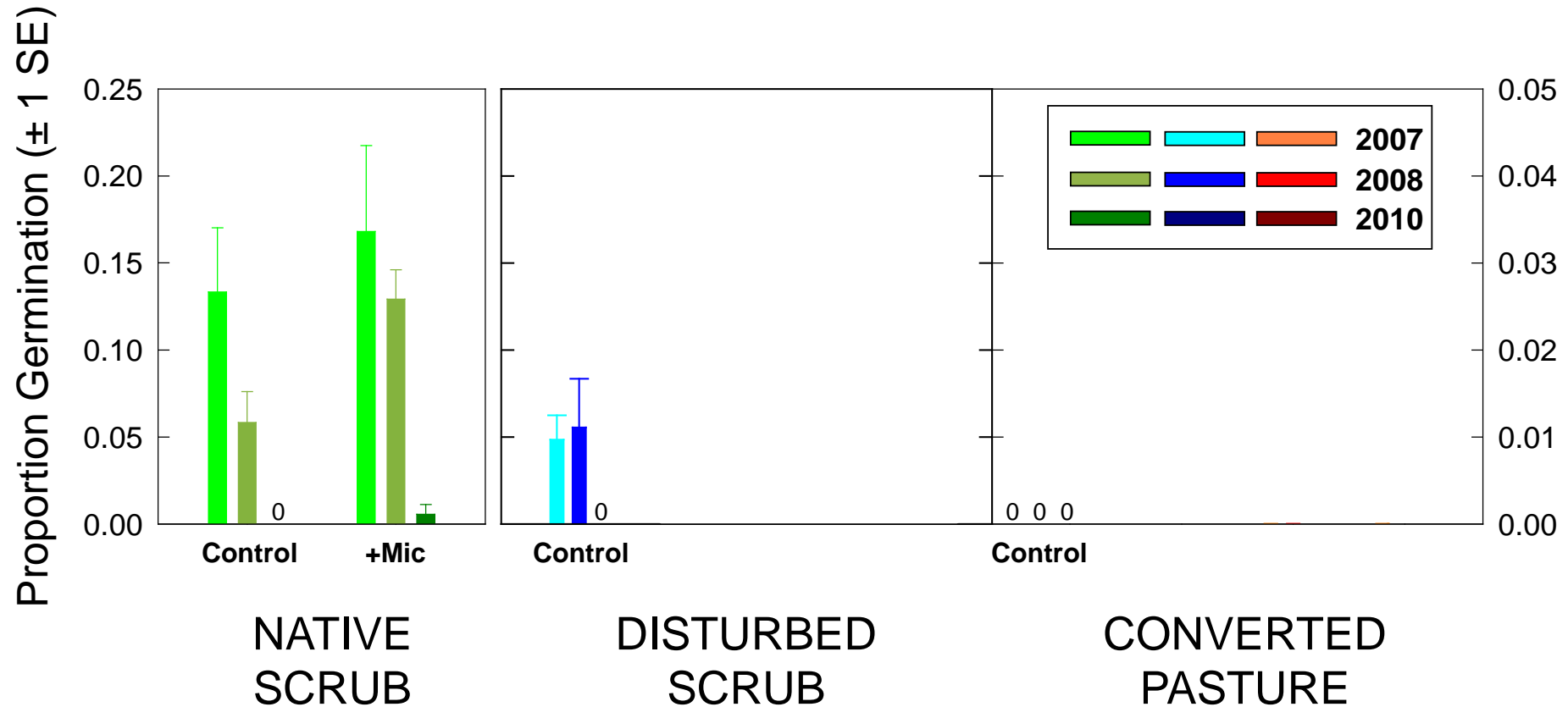
*Paronychia chartacea*\*\*



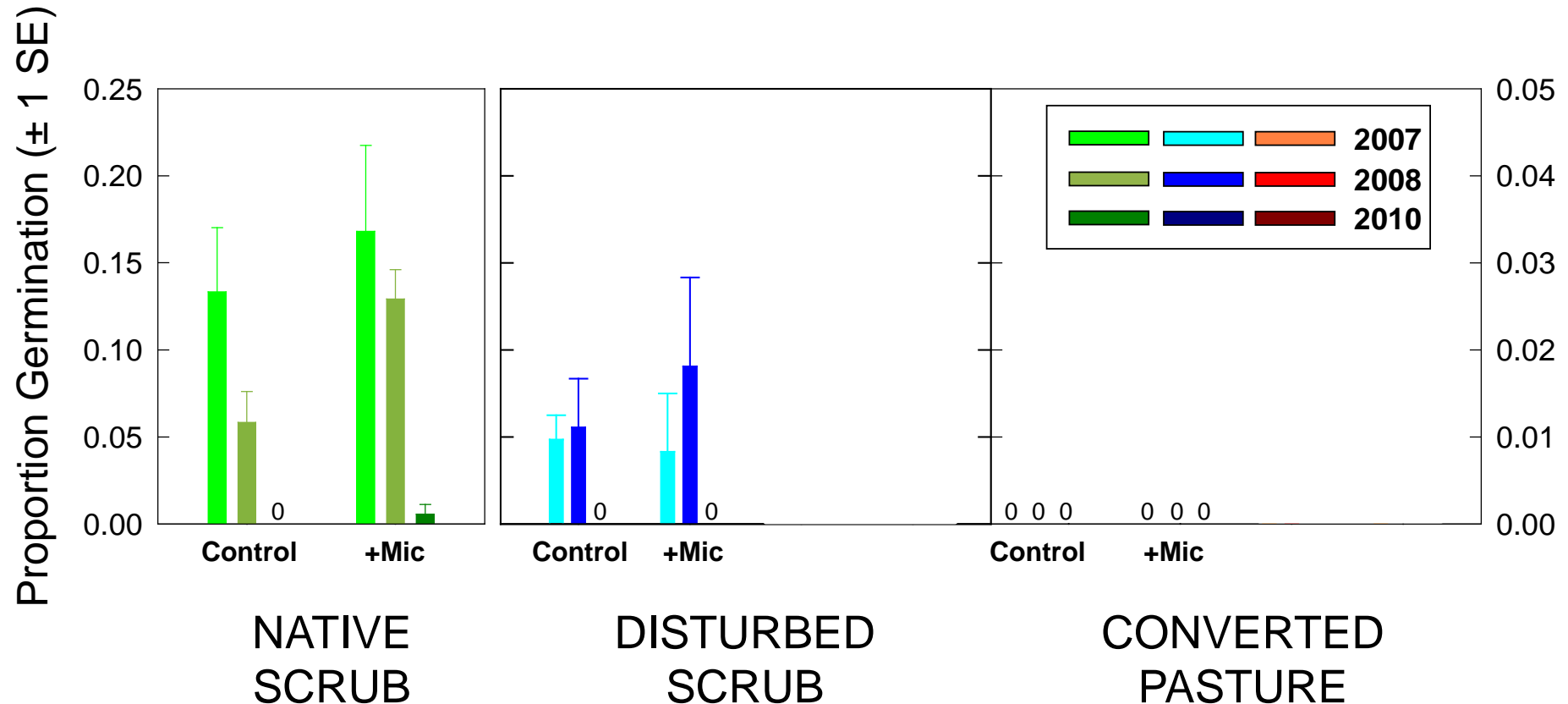
# More germination in native sites



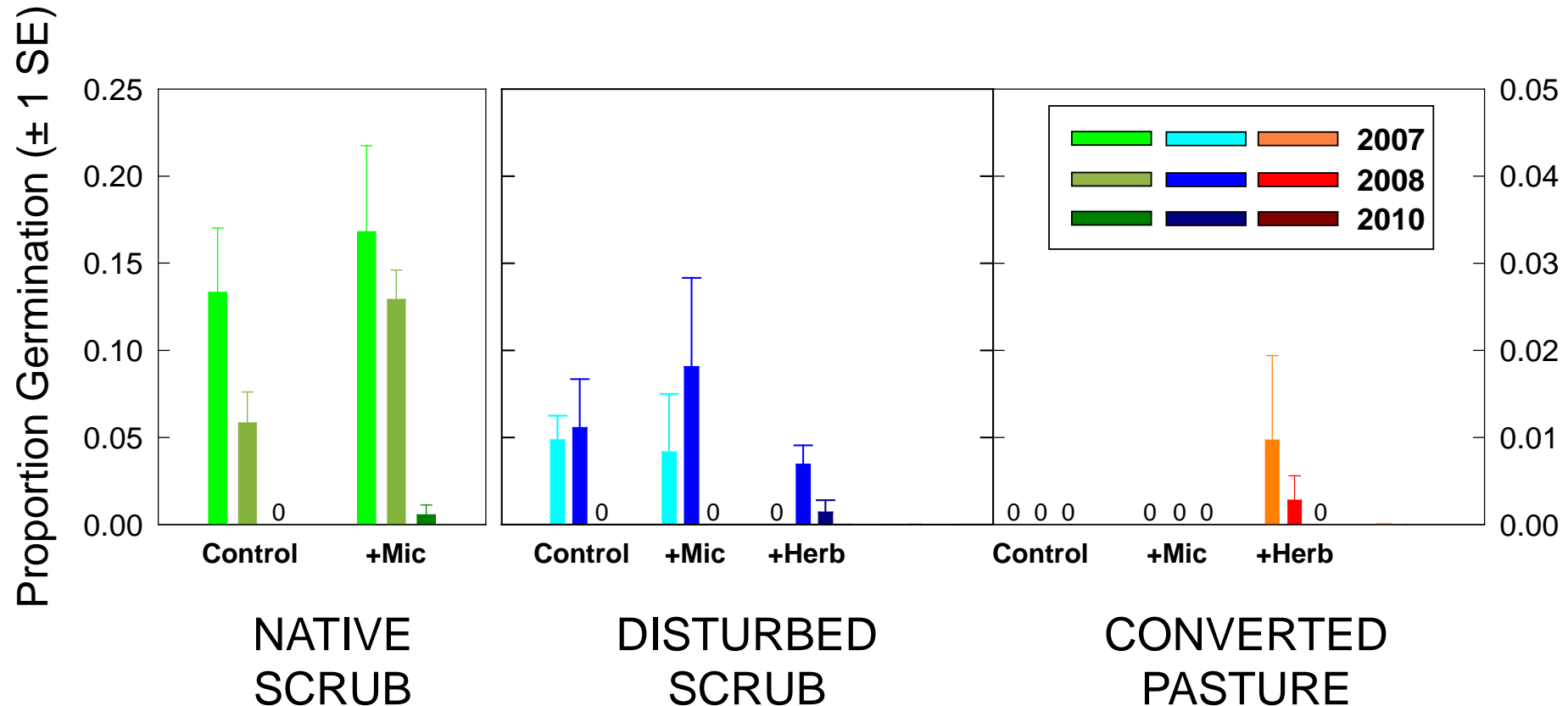
# More germination in native sites



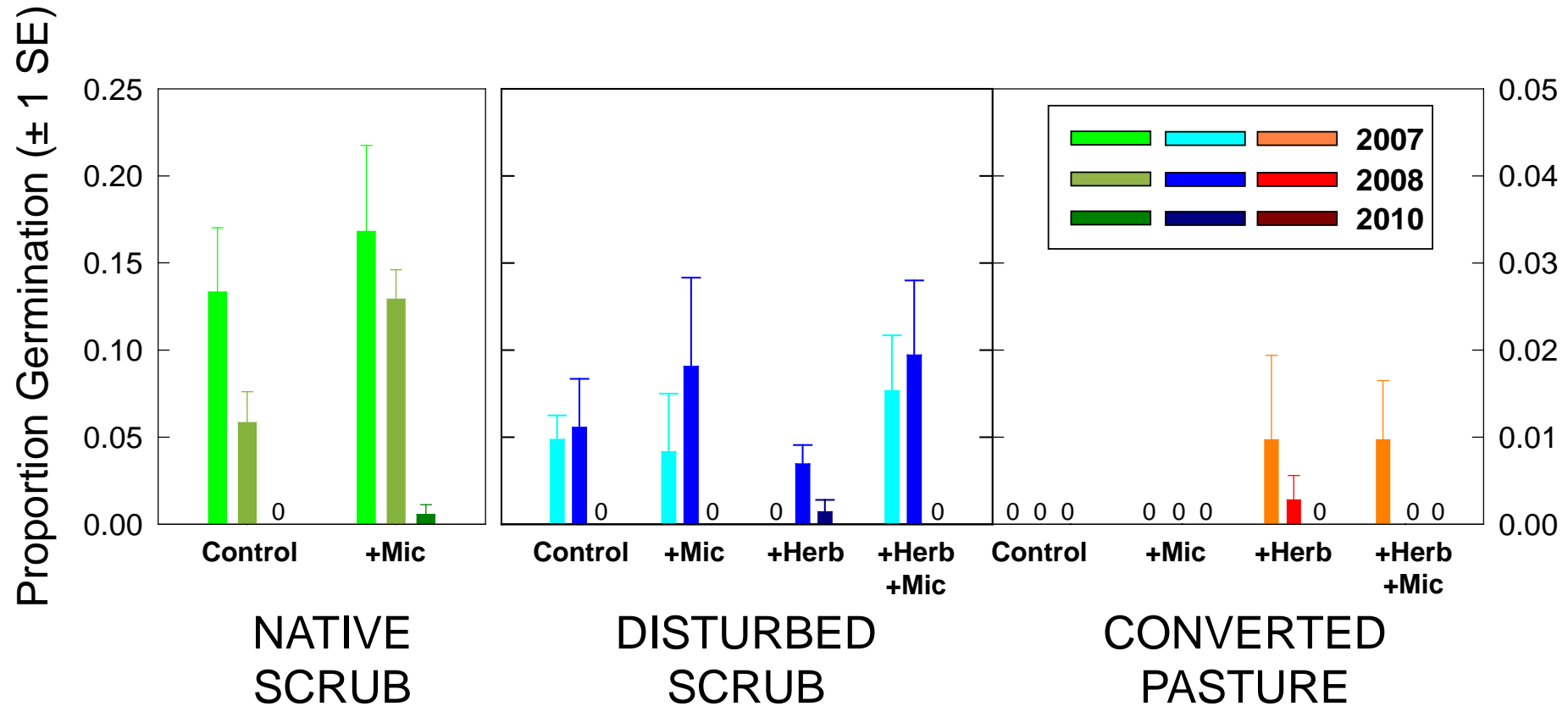
# Microbial additions alone do not improve germination



# Plant removals improve germination only in pastures

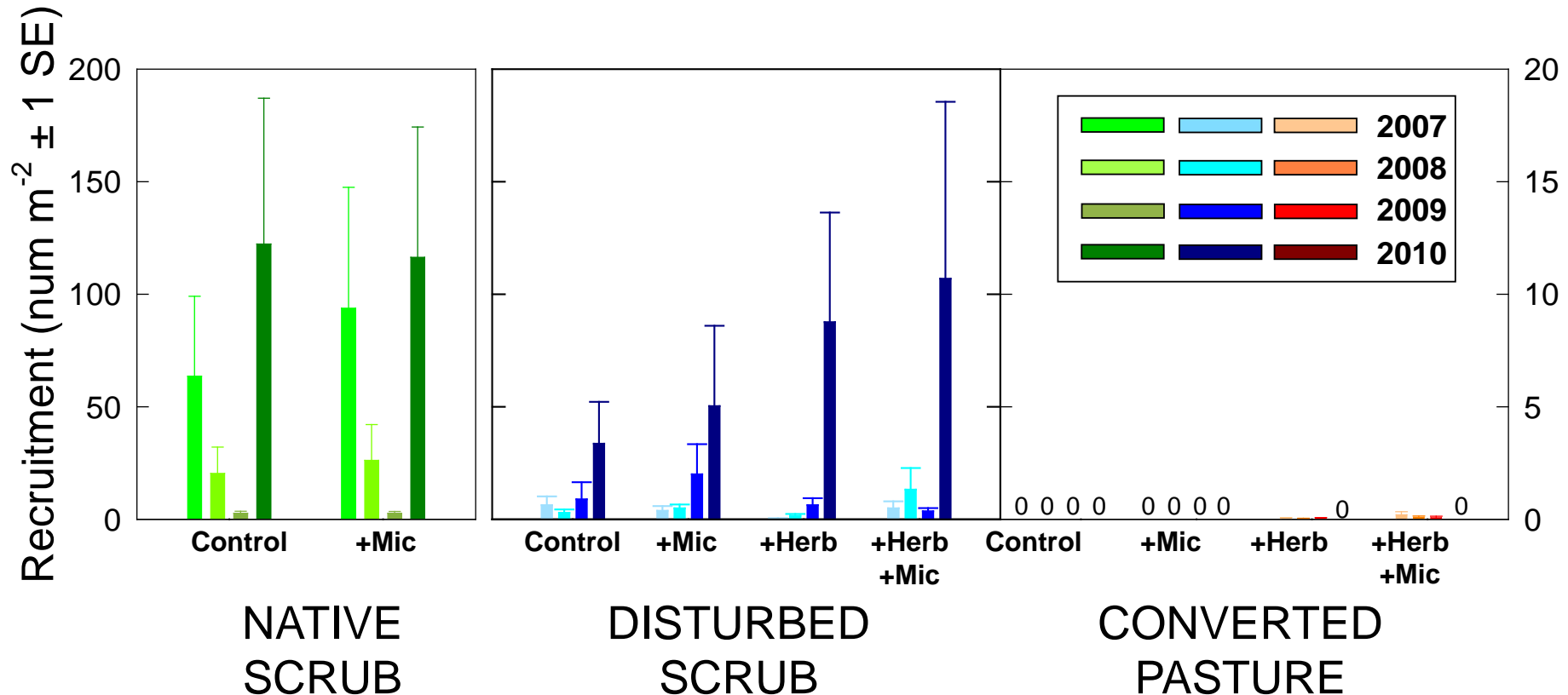


# Plant removals with microbial additions recover germination in disturbed sites



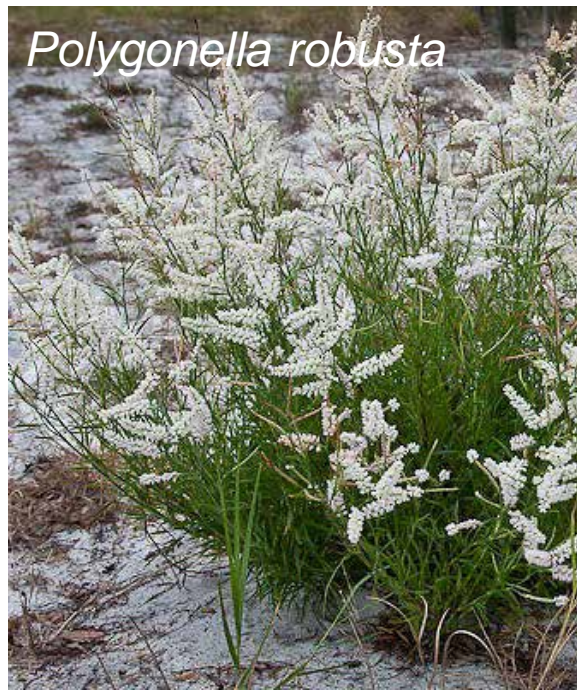


# Substantial native background recruitment, but not in pastures

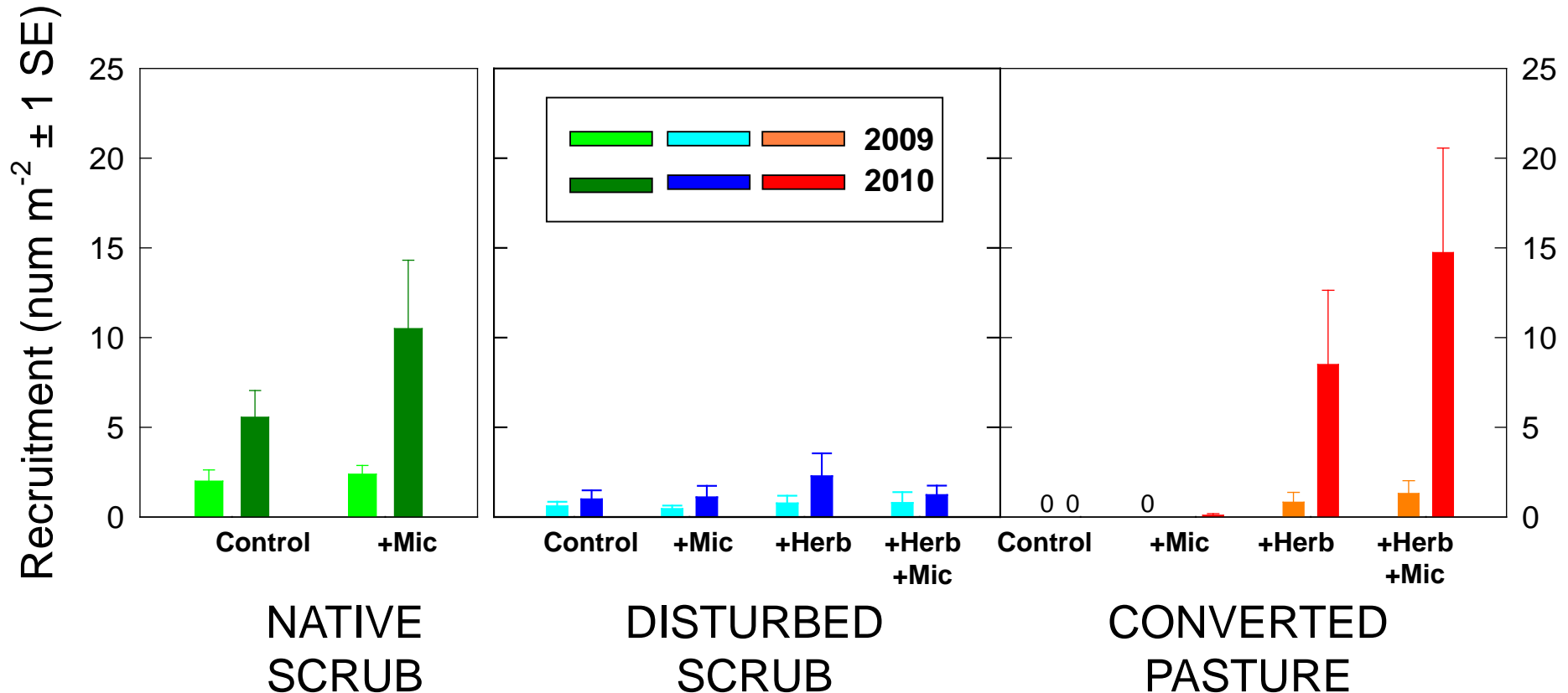


Open space, moisture, and non-native veg cover explain 38% of variation in recruitment

# Background recruitment of non-targeted species



# Opportunities created for recruitment of non-targeted species in pastures



Open space and soil aggregation explain 43% of variation in non-target recruitment

# What does this mean for population viability?

- Demographic modeling of taxa to estimate population growth rates
- Started with one of the most abundant plant species, *Polygonella basiramia*



# *Polygonella* population growth rates increased with restoration treatments

| VEGETATION | TREATMENT | $\Delta$ LAMBDA RELATIVE TO CONTROLS |
|------------|-----------|--------------------------------------|
|------------|-----------|--------------------------------------|

|                 |                    |       |
|-----------------|--------------------|-------|
| Disturbed Scrub | Microbes           | 0     |
|                 | Herbicide          | 0     |
|                 | Herbicide+Microbes | +2.8% |

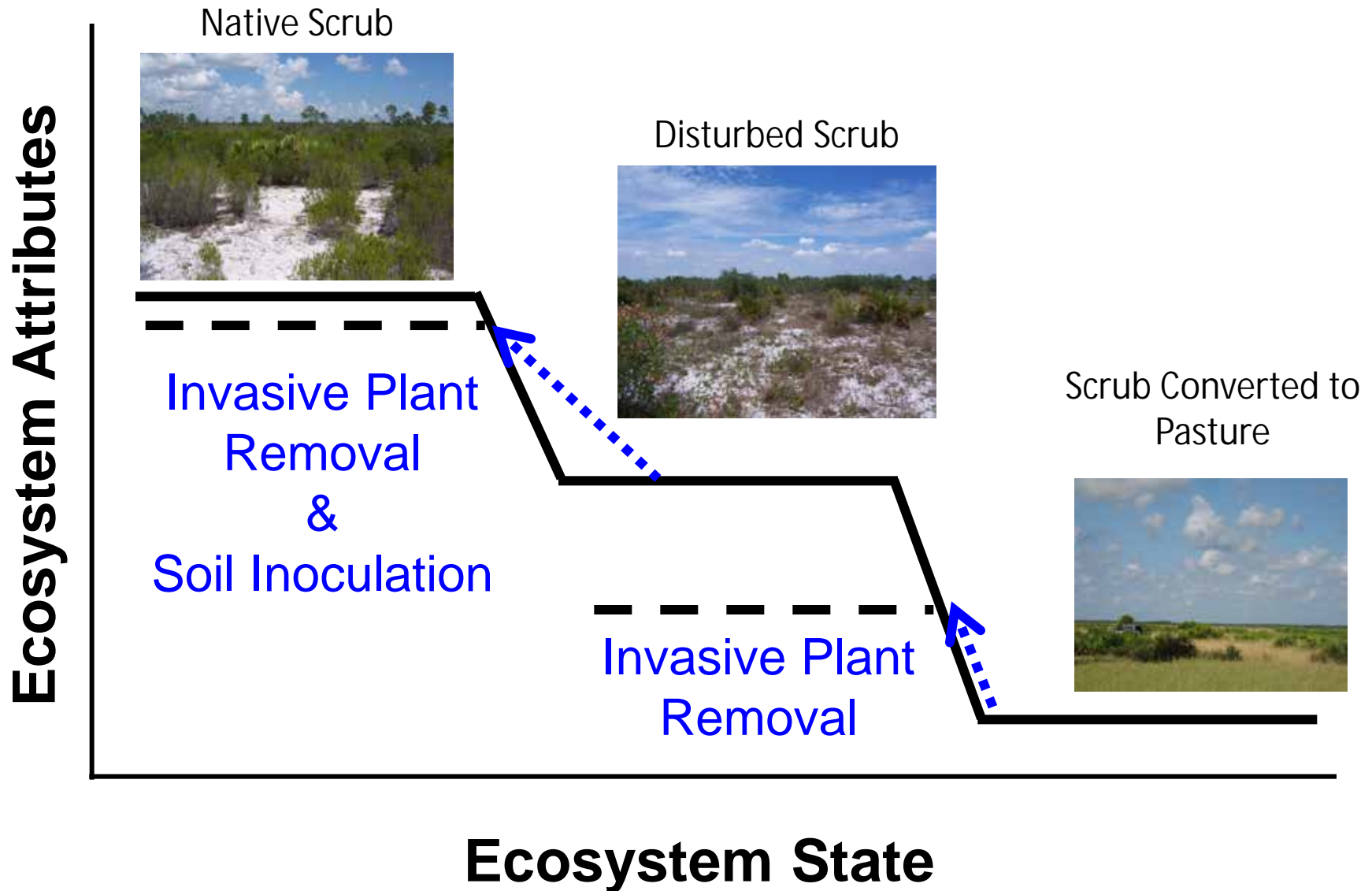
|                   |                    |       |
|-------------------|--------------------|-------|
| Converted Pasture | Microbes           | 0     |
|                   | Herbicide          | +2.5% |
|                   | Herbicide+Microbes | +1.1% |

# Measuring restoration success

- Return site characteristics?
  - Yes, degraded site cover, openness, & soil crust aggregation were improved.
- Reduce nutrient legacies?
  - No. May need more time. Possibly remove topsoil if N reduction is necessary.
- Improve plant recovery?
  - Yes, but veg-specific effects on germination, establishment, and population trajectories.



# The probability of successful restoration differs among sites



# Can we open up the soil black box?

- How do microbial communities differ among native, disturbed, and pasture sites?
- Do those differences persist?
- Can we use what we learn to further enhance restoration success?



# Focus on fungi

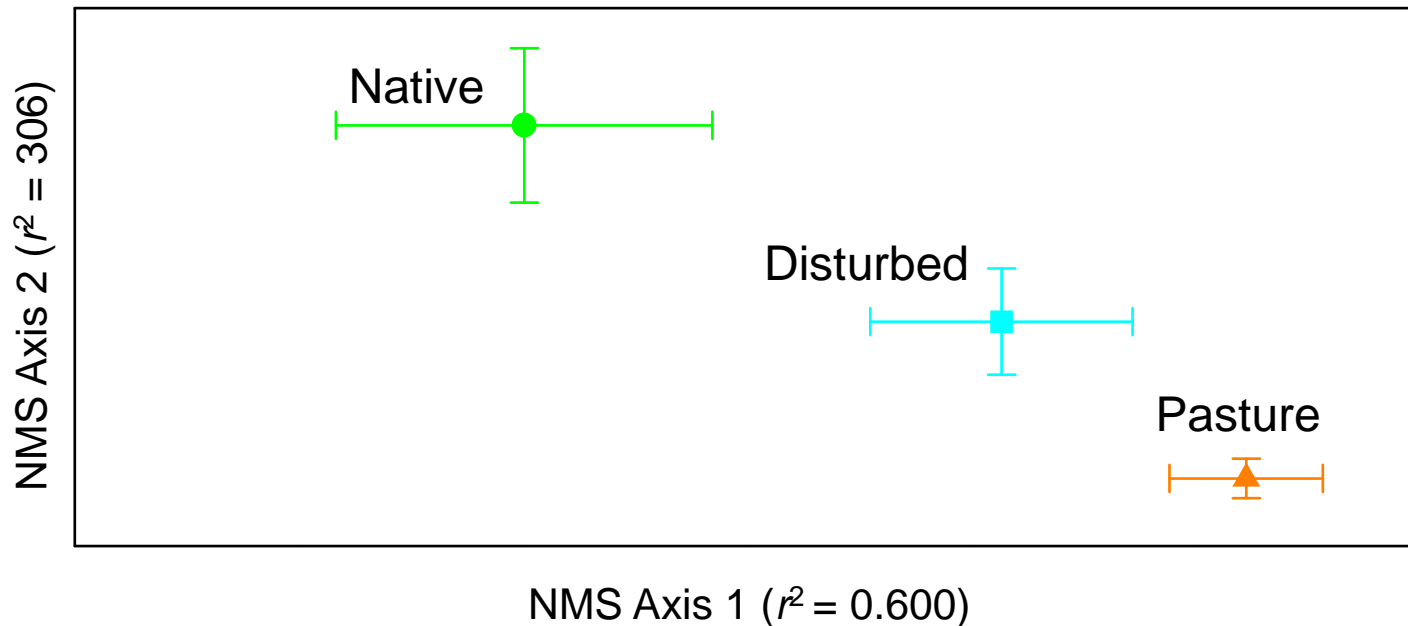
- Root fungi
  - Involved in nutrient and water uptake
- Soil fungi
  - Responsible for decomposition and nutrient recycling in soil
- Both abundant in this ecosystem and should play important functional roles given the low nutrient, xeric soils



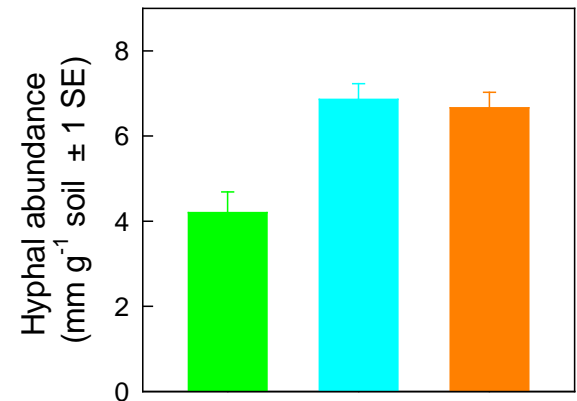
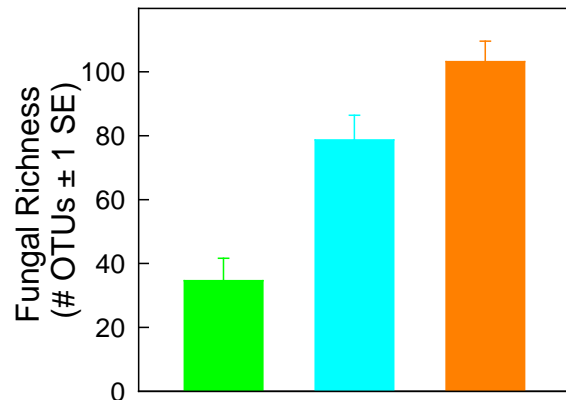
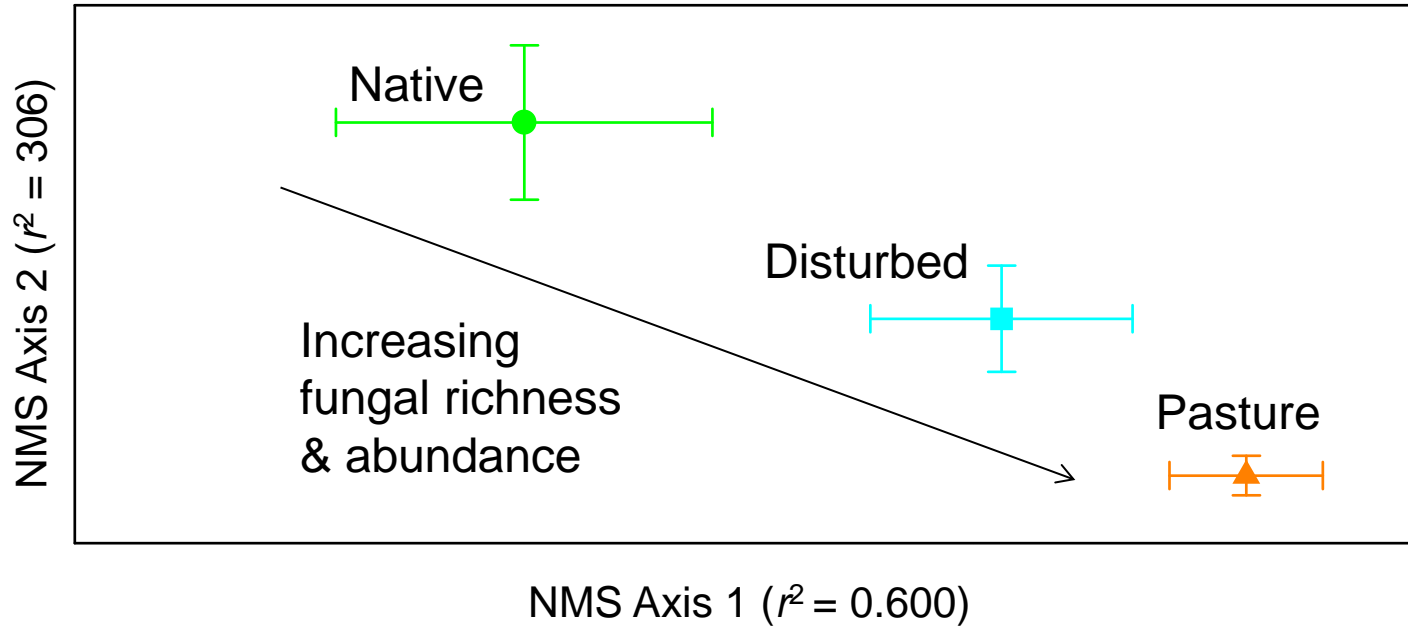
# Can we open up the soil black box?

- How do **fungi** communities differ among native, disturbed, and pasture sites?
- Do those differences persist?
- Can we use what we learn to further enhance restoration success?

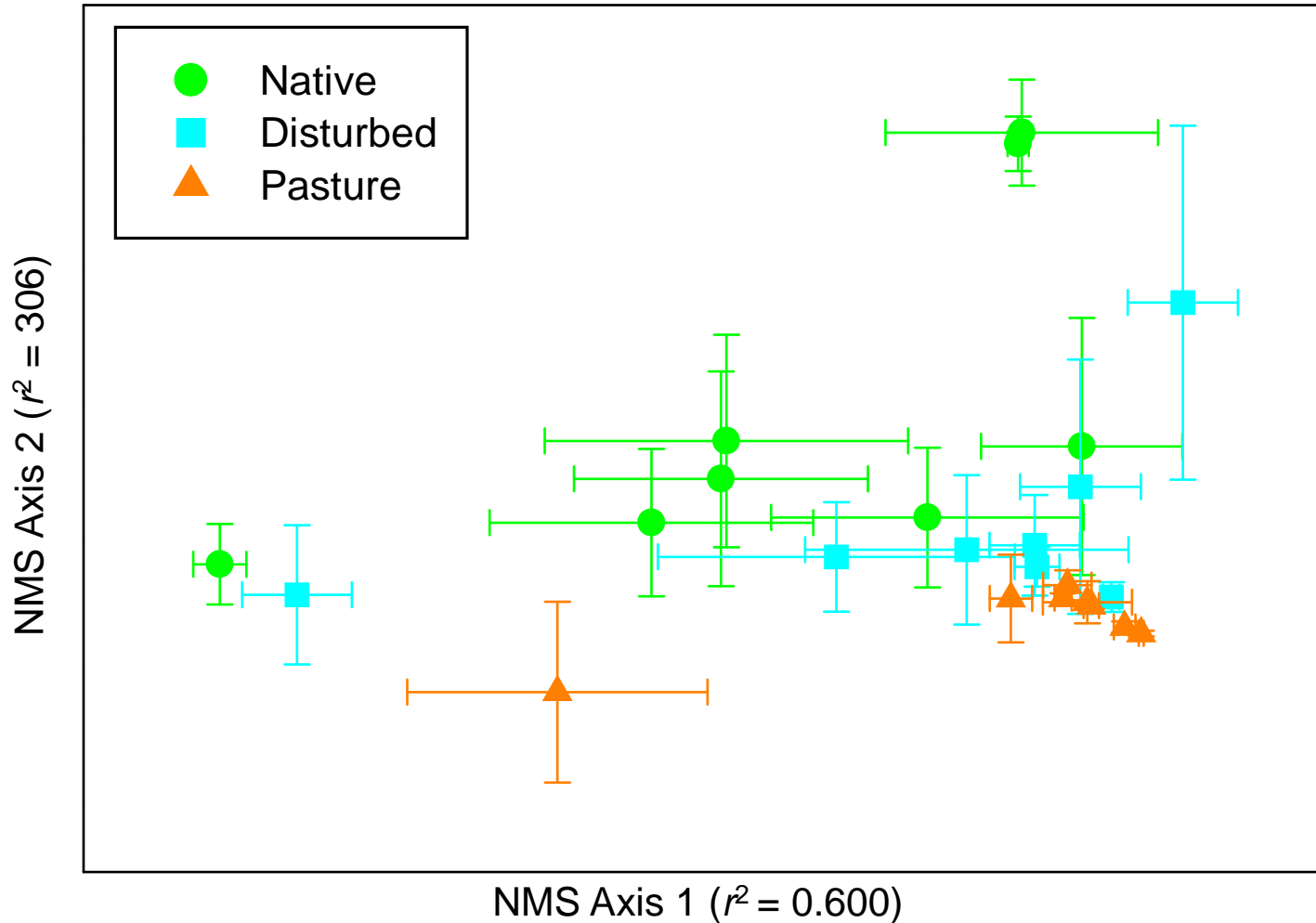
# Soil fungi had little overlap among veg types regardless of treatment



# Native sites have lower diversity and fewer fungi



# Strongest fungal legacy in pastures



# Can we open up the soil black box?

- How do **fungi** communities differ among native, disturbed, and pasture sites? Do differences persist?
  - Yes! There are strong differences over three years, likely related to changes in soil organic matter.
- Can we use what we learn to further enhance restoration success?

# Can we open up the soil black box?

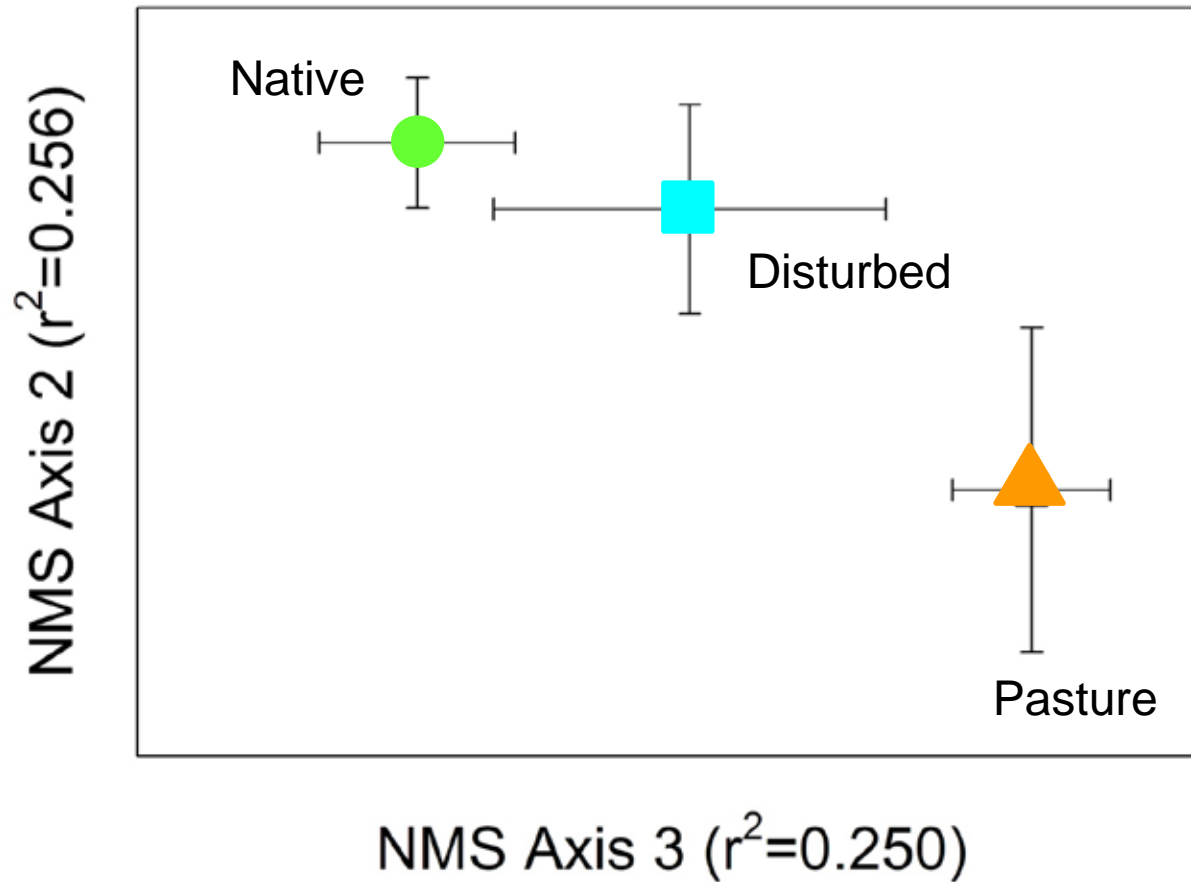
- How do **fungi** communities differ among native, disturbed, and pasture sites? Do those differences persist?
  - Yes! There are strong differences over three years, likely related to changes in soil organic matter.
- Can we use what we learn to further enhance restoration success?

# Manipulating specific fungi





# Cultured fungi reflect whole community patterns



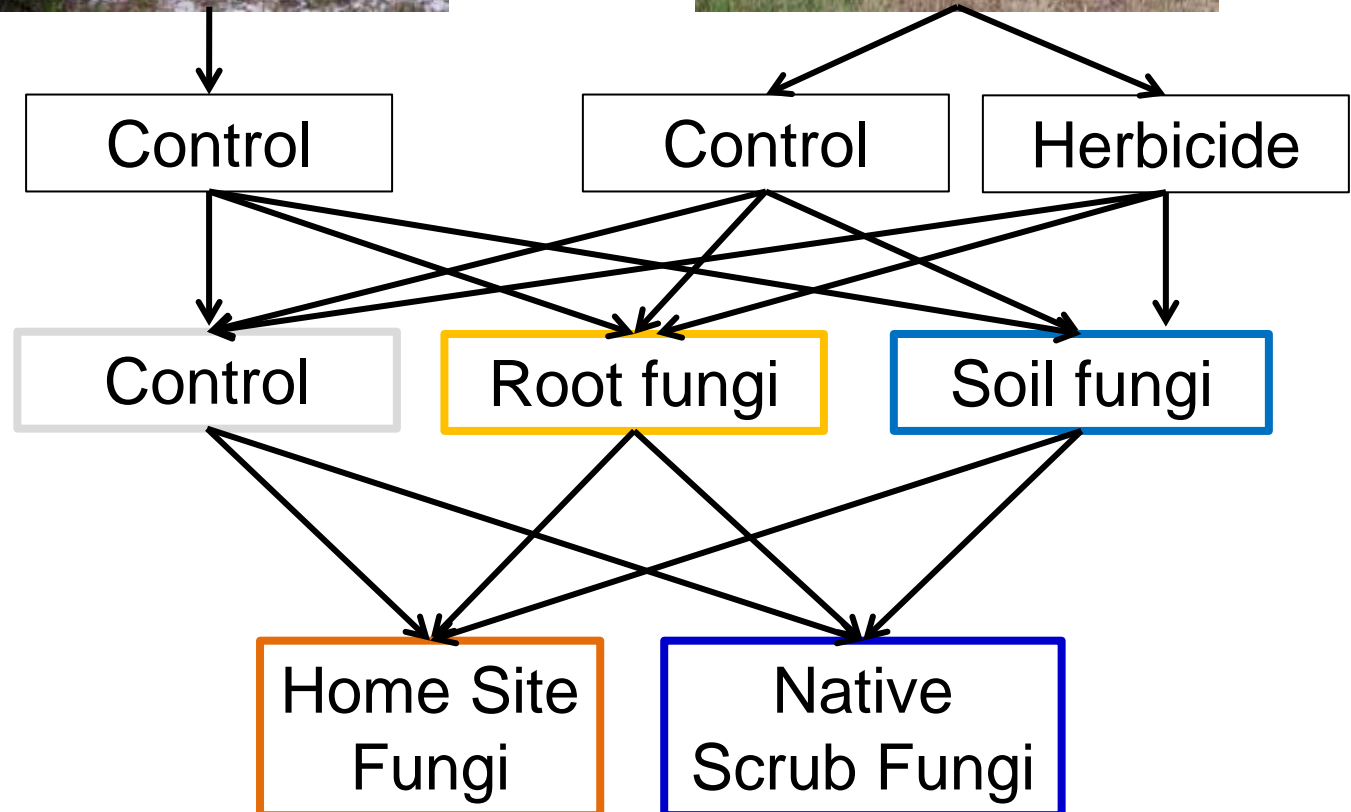


# Select fungi for amendments

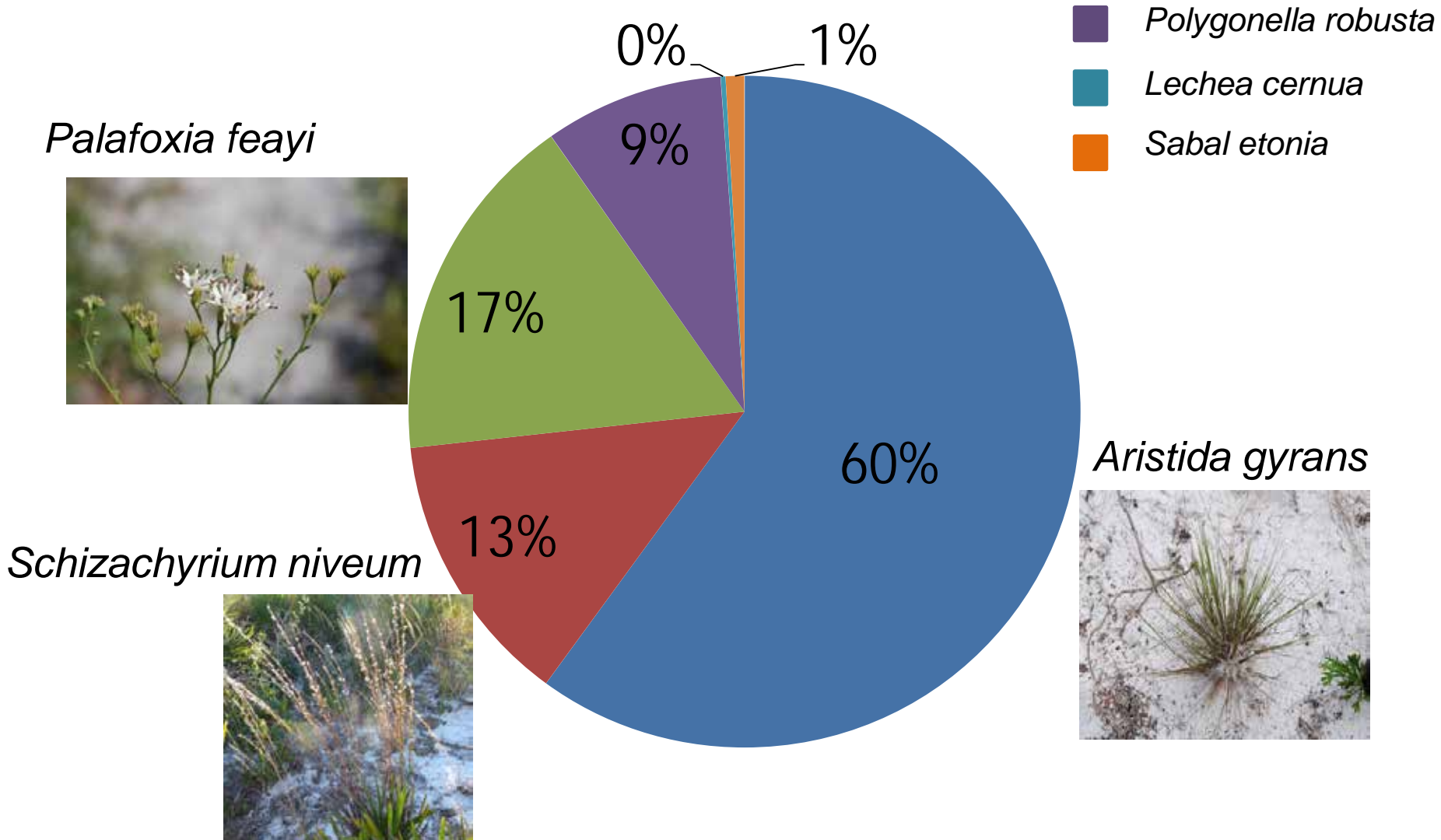
- Fungi selected for variable function based on prior tests of competition and decomposition
- 8 root and 5 soil fungal isolates per site type



# Testing fungal effects on restoration efforts

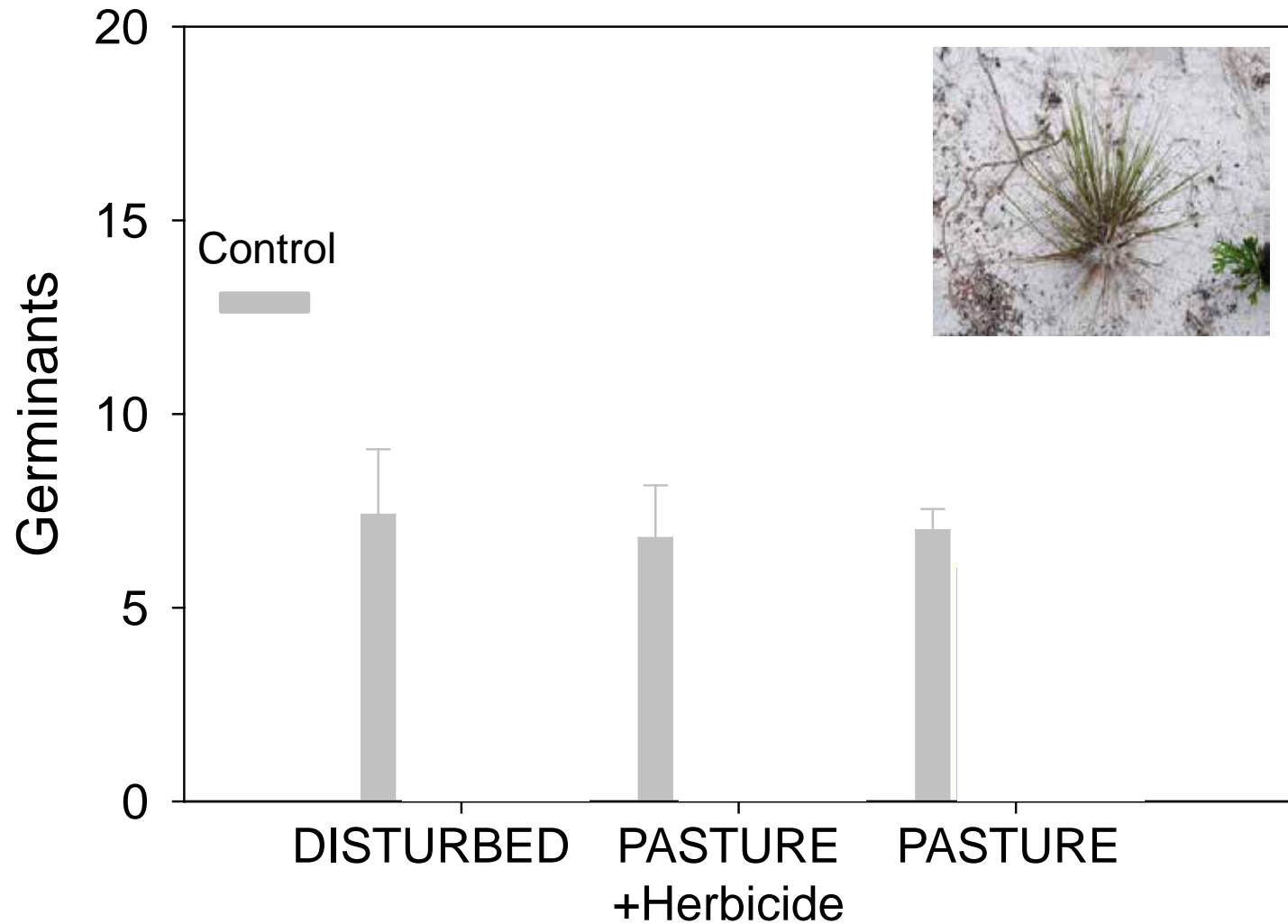


# Overall native plant germination was dominated by three species

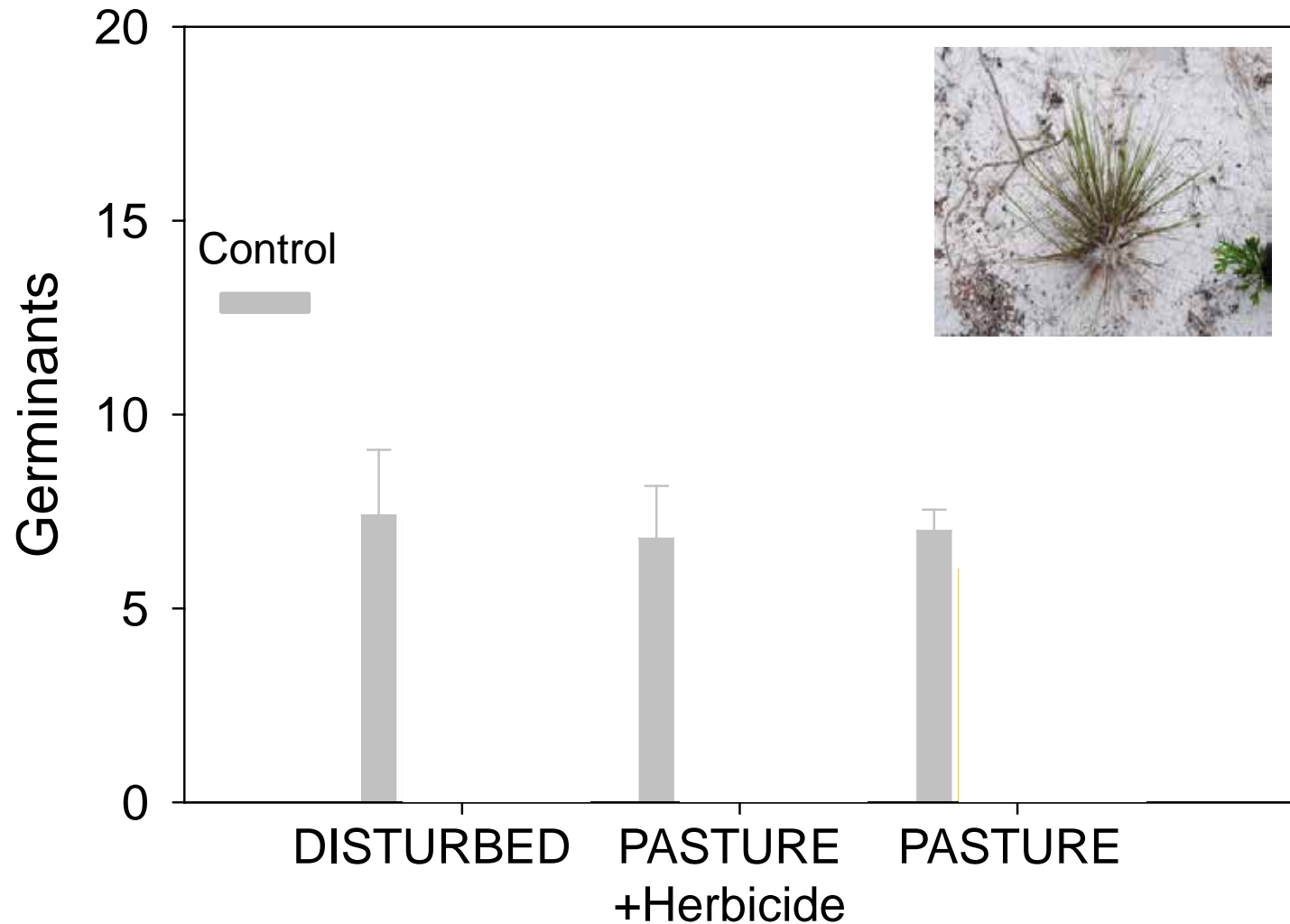




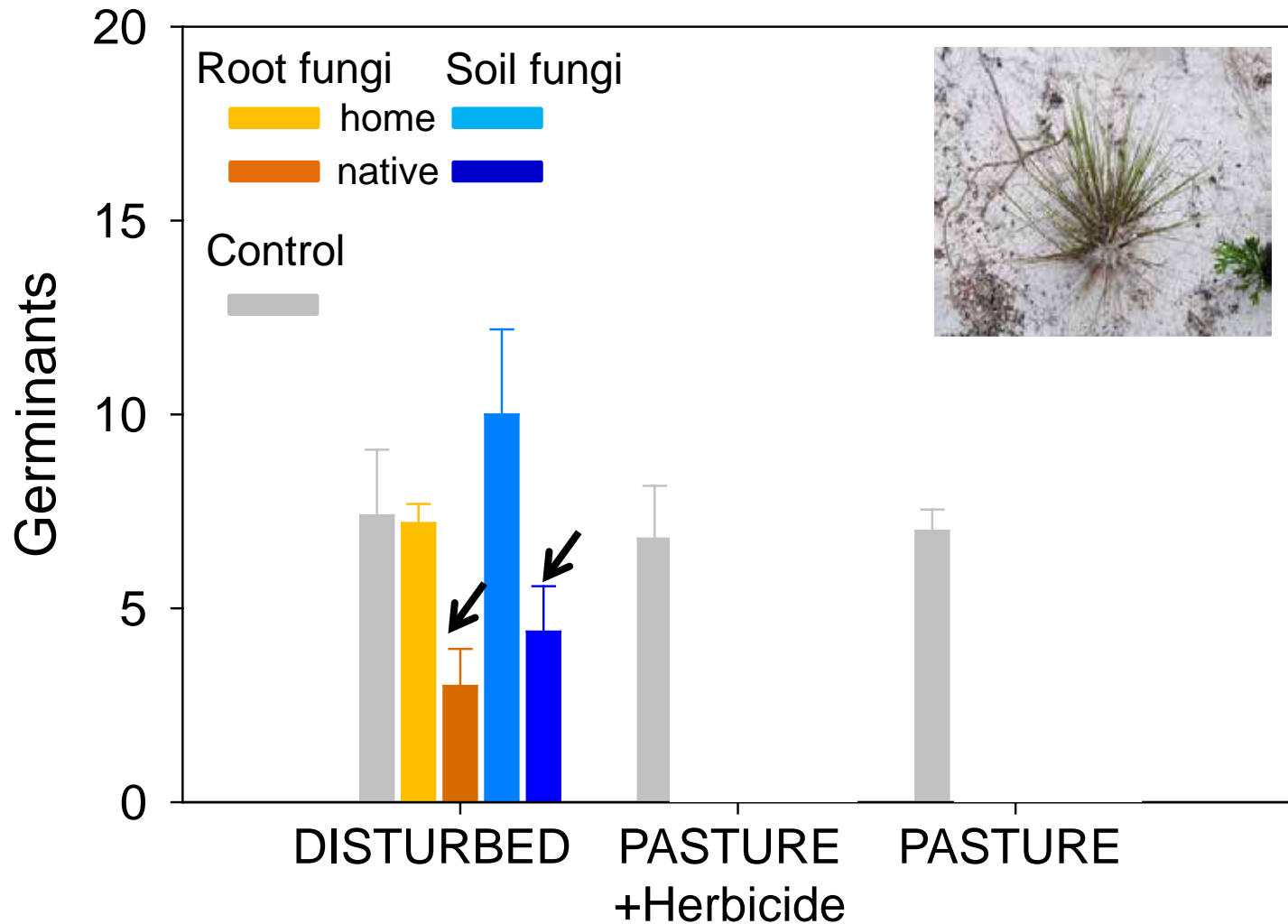
# No variation in *Aristida* in the absence of fungi



# Fungal effects on *Aristida* were highly context dependent

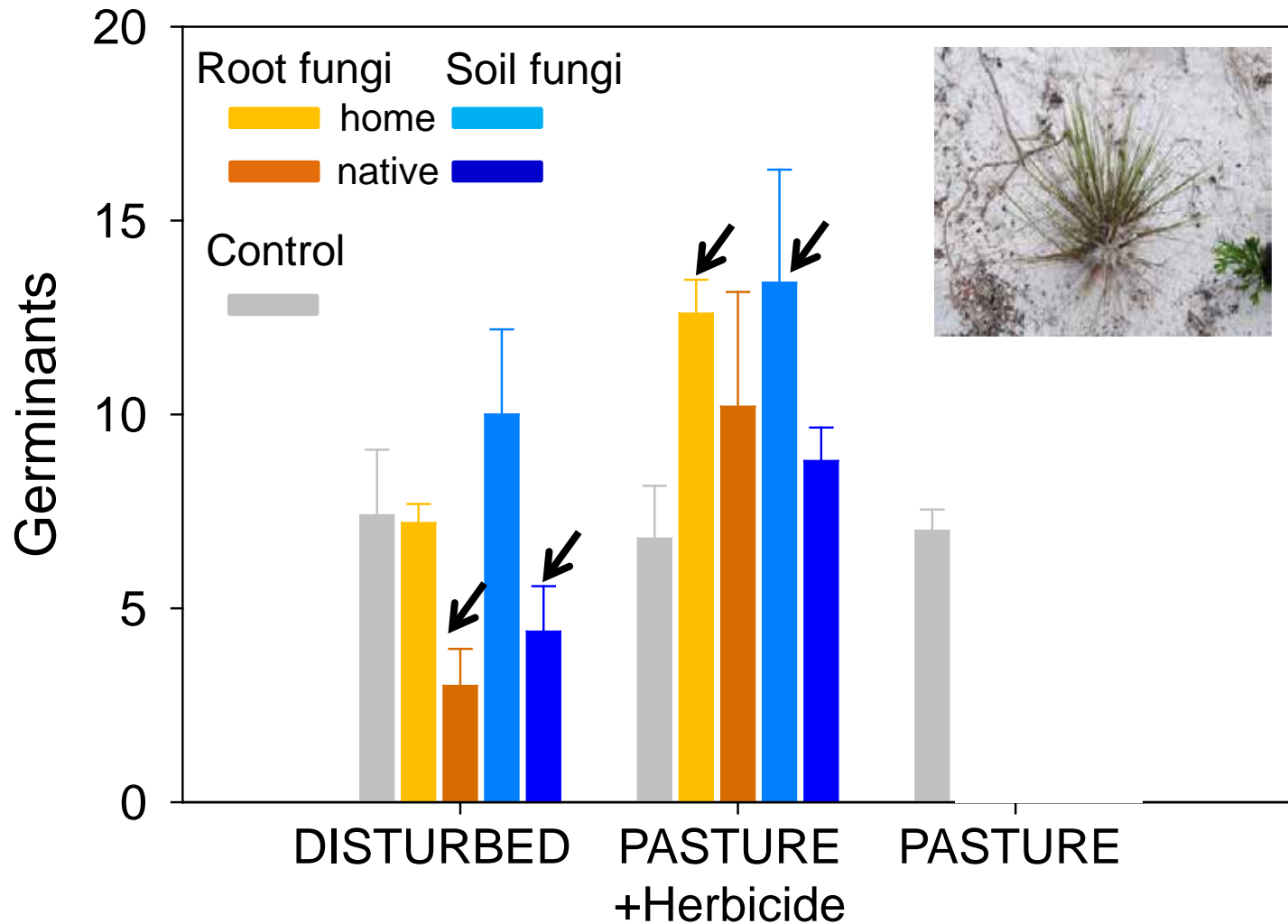


# *Aristida* germination inhibited in disturbed sites by fungi from native scrub

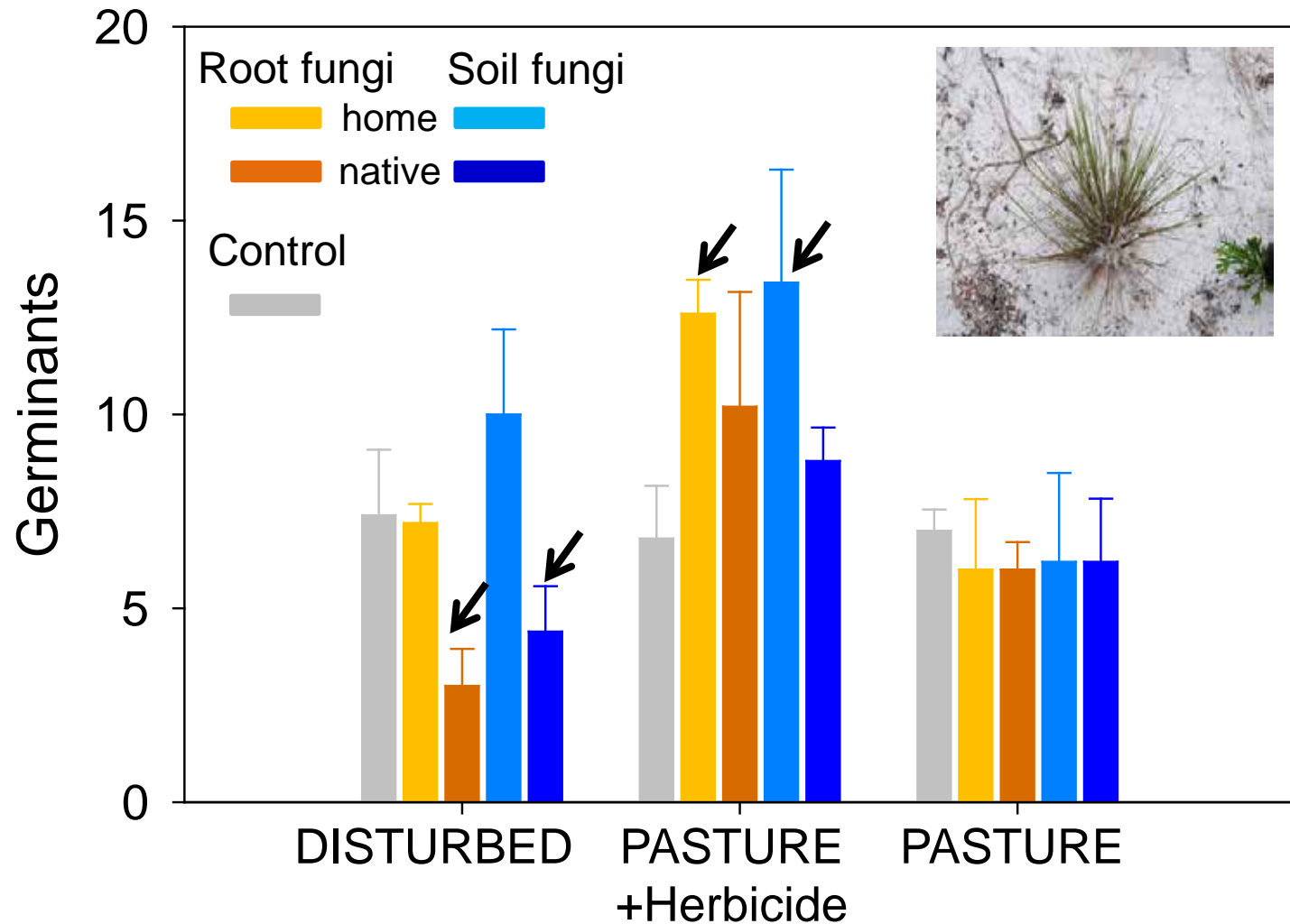




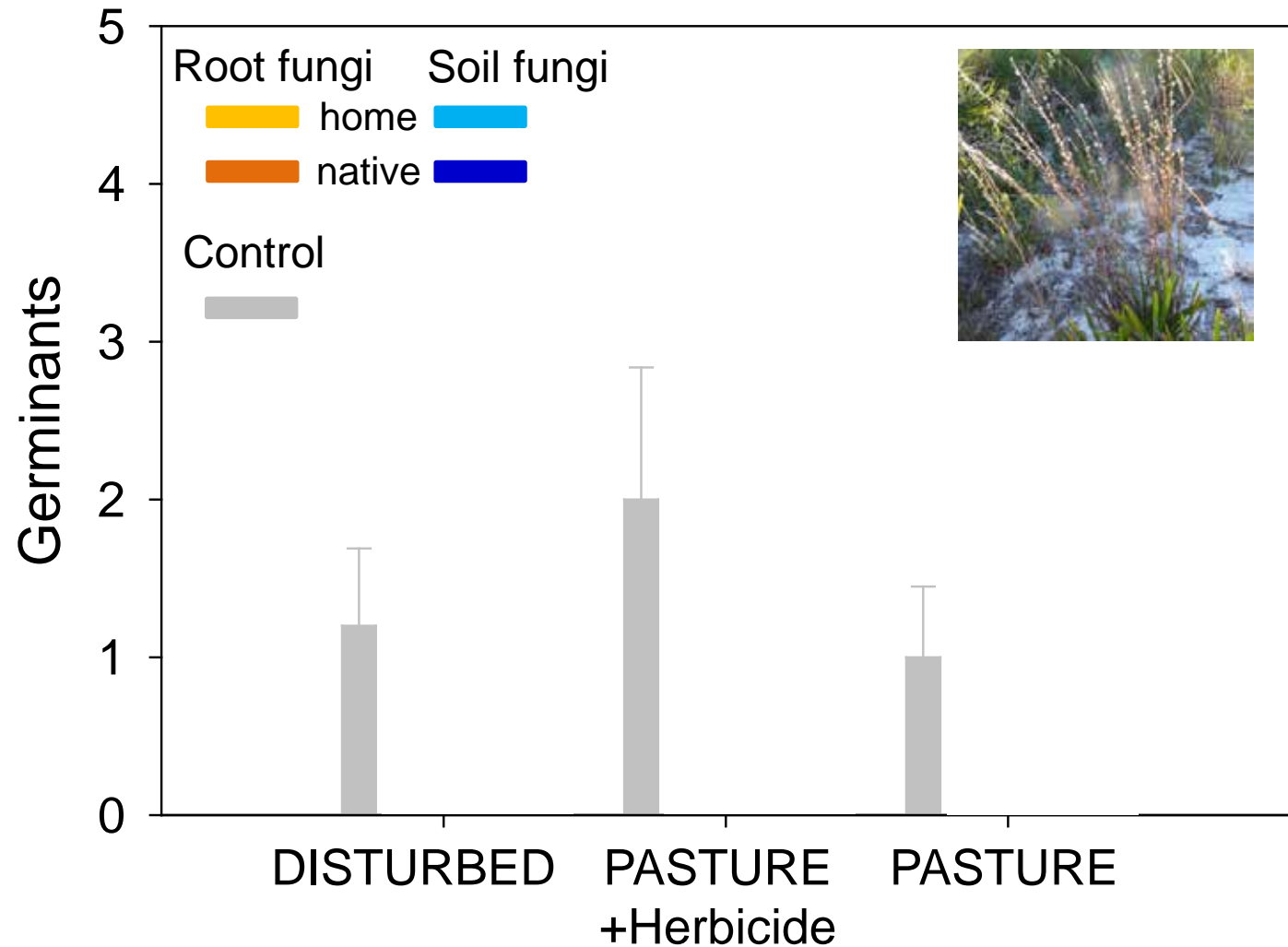
# Fungi from pastures improved *Aristida* germination in pastures with veg removed



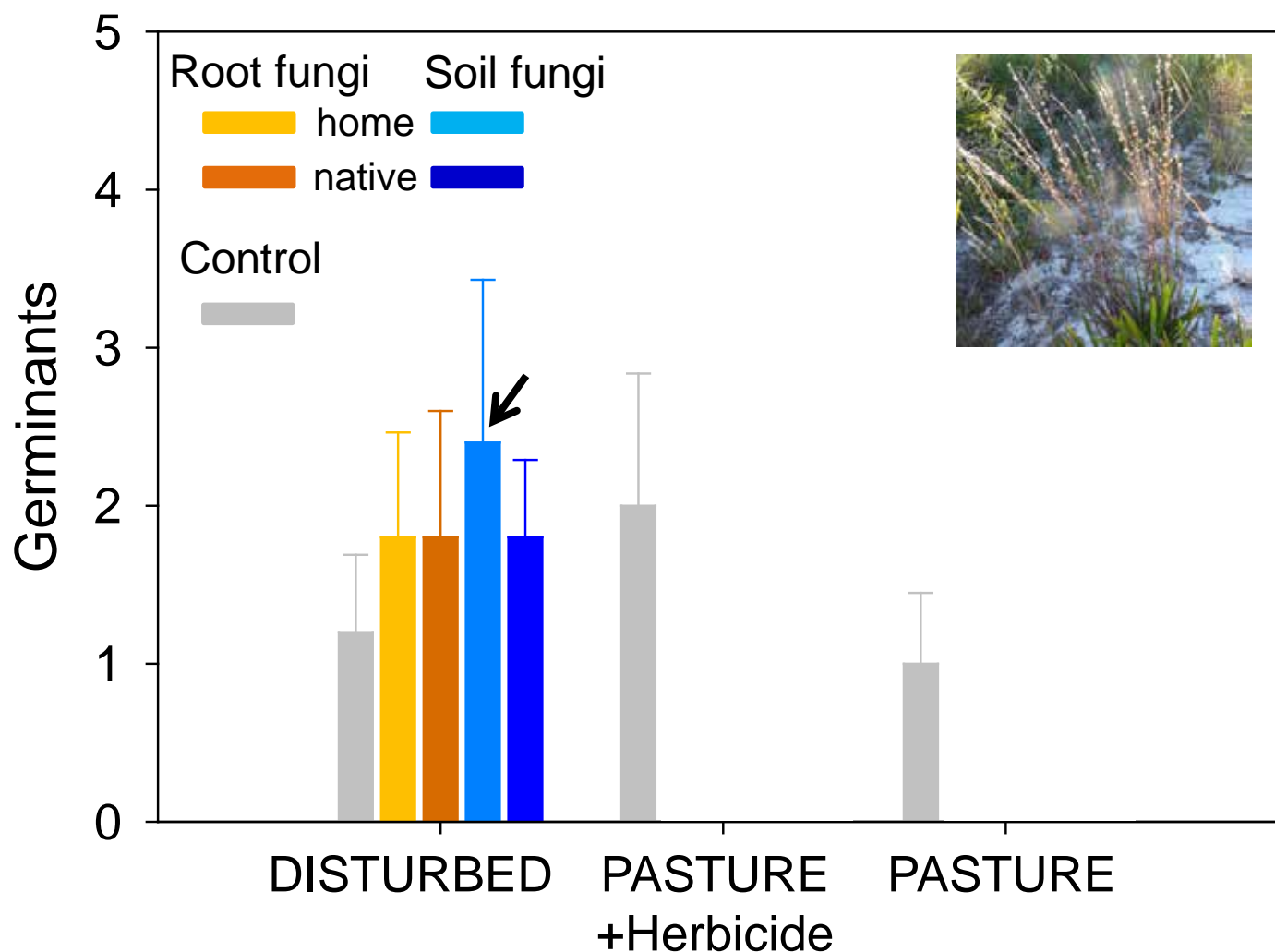
# Fungi had no effect on *Aristida* germination in pastures when grasses were not removed



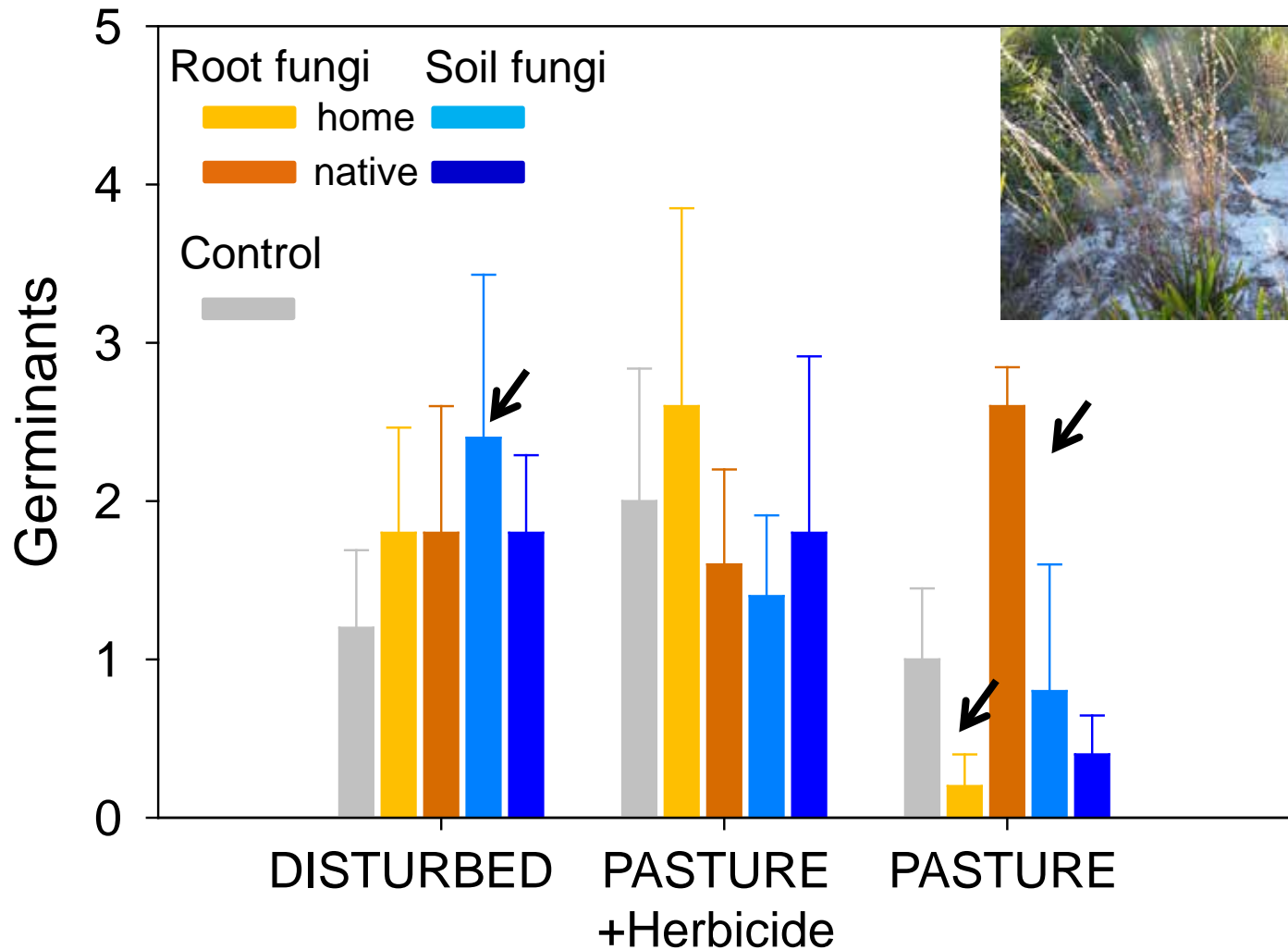
# Without fungi, *Schizachyrium* germination varied little across sites



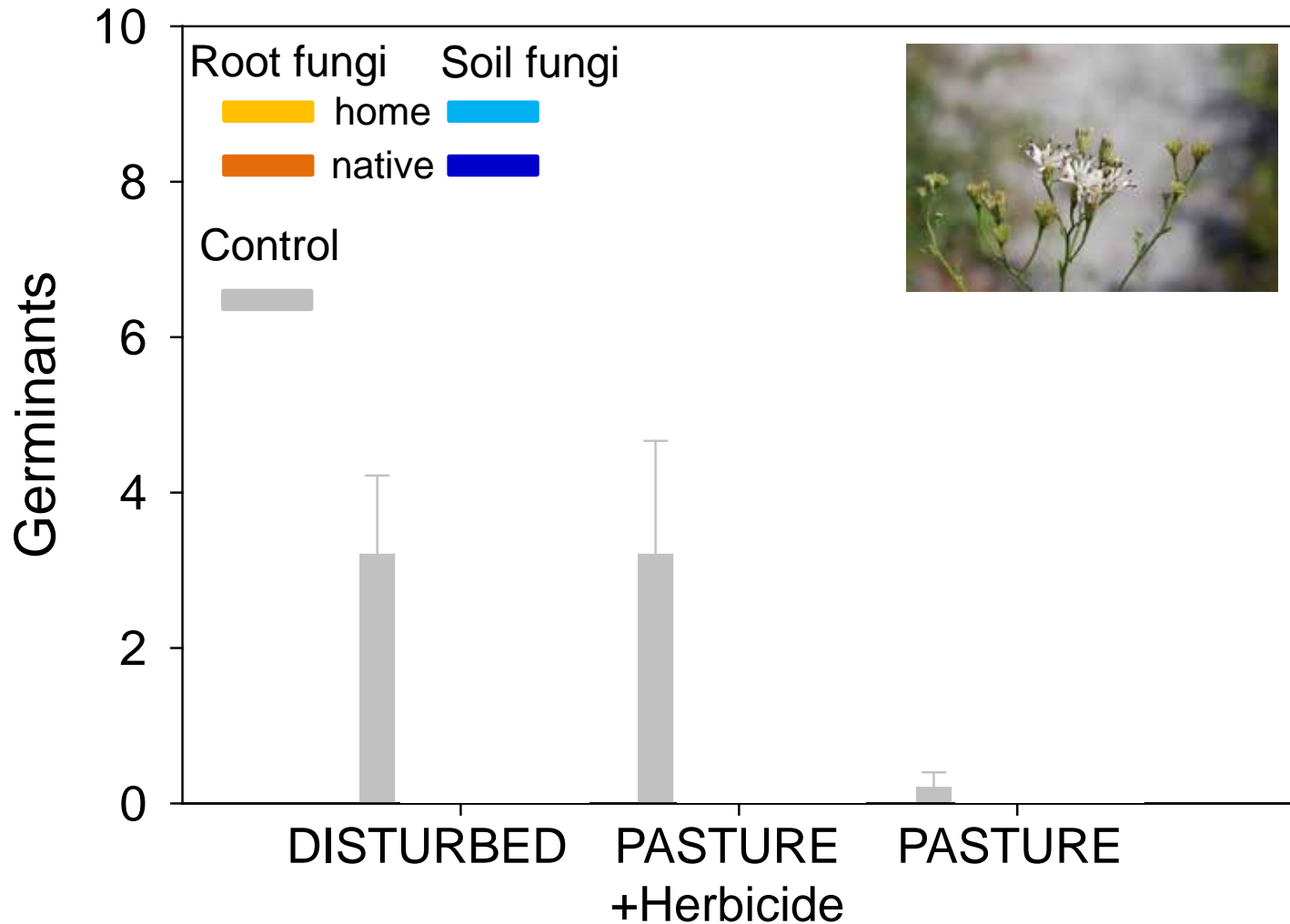
# Fungi from disturbed sites improved *Schizachyrium* germination in disturbed sites



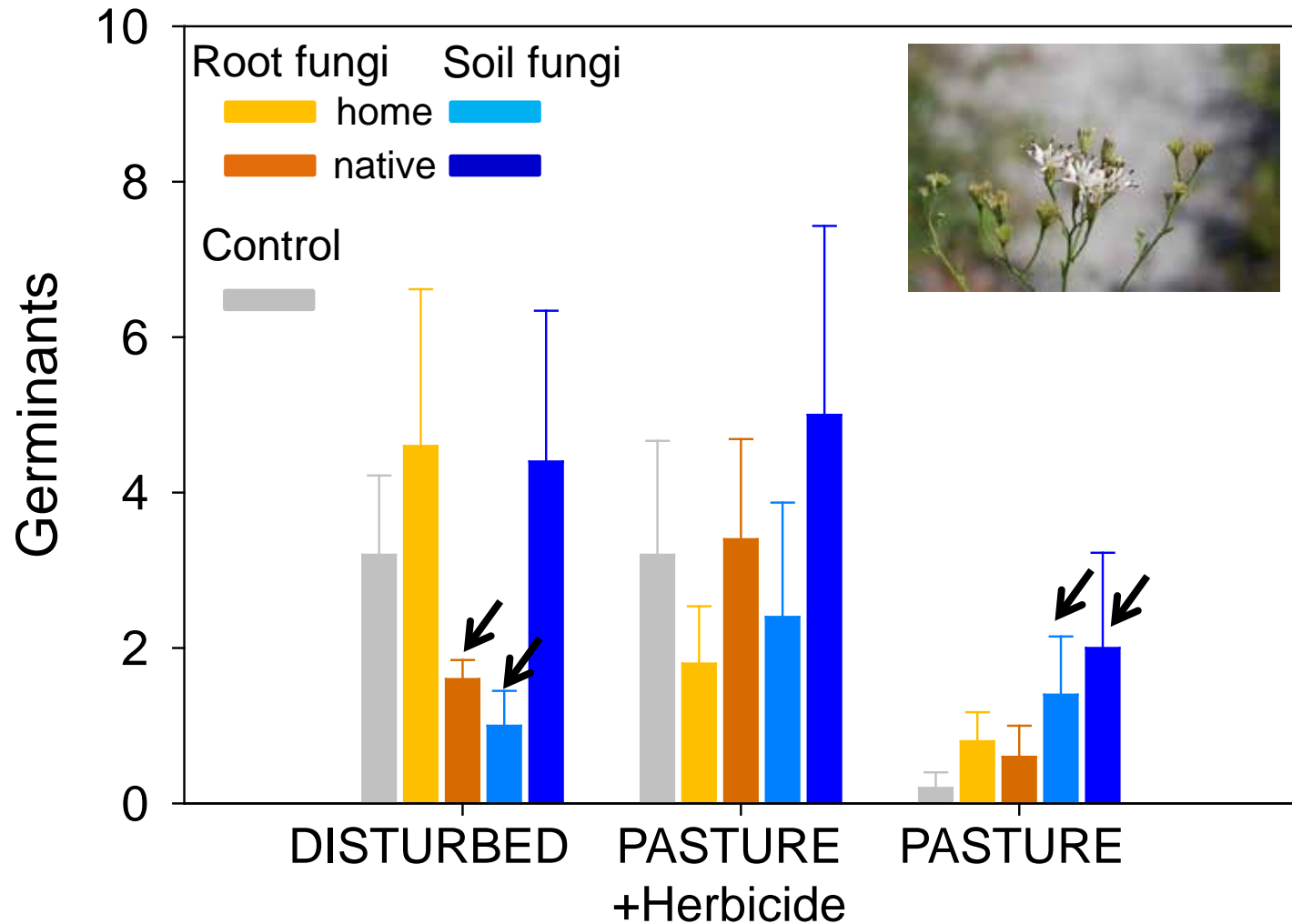
# In pastures, root fungi from native scrub improved *Schizachyrium* germination



# *Palafoxia* germination lower in pastures without fungi



# Fungal effects on *Palafoxia* are also highly context dependent

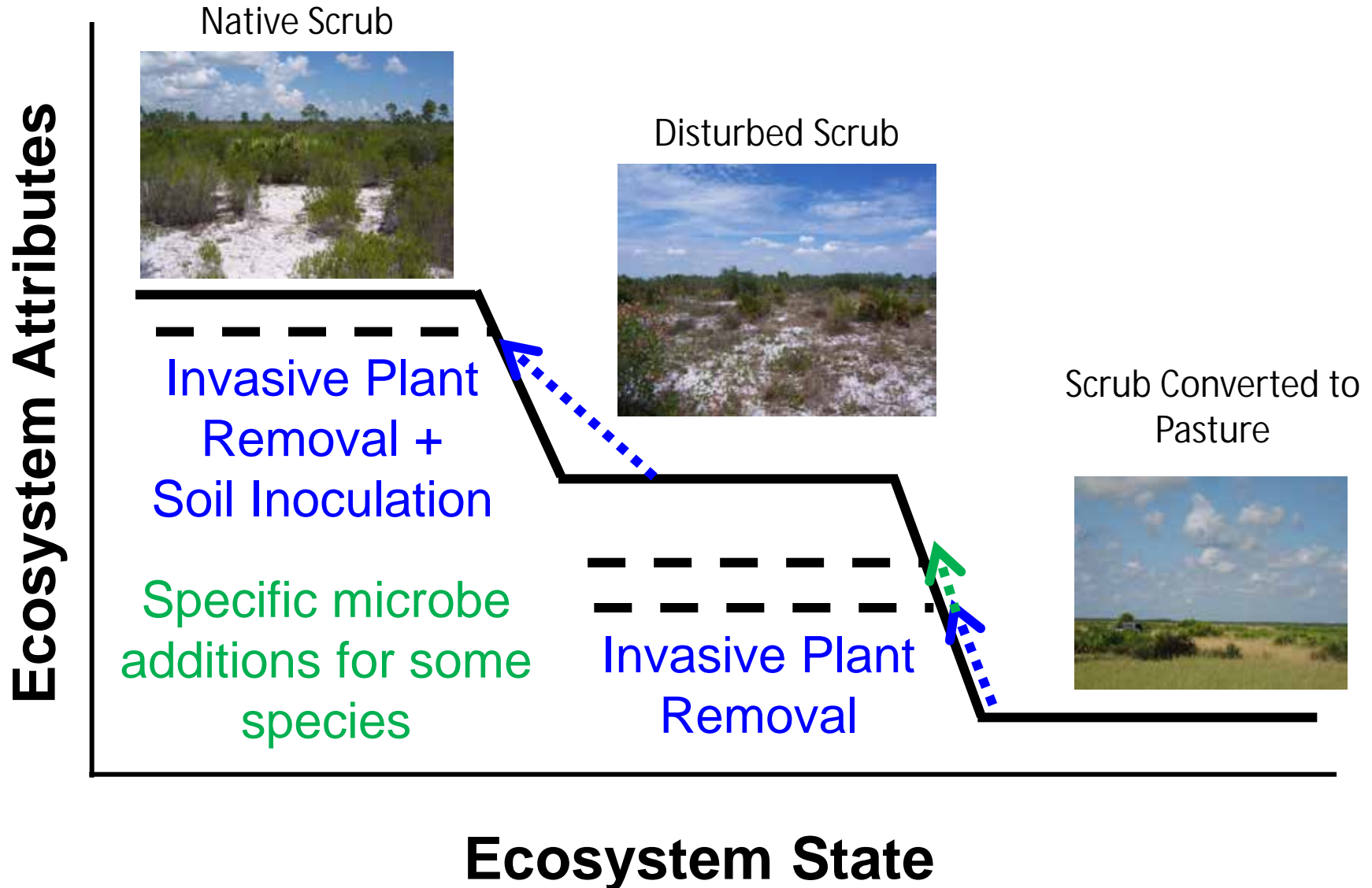


# Can we open up the soil black box?

- How do **fungi** communities differ among native, disturbed, and pasture sites? Do those differences persist?
  - Yes! There are strong differences over three years, likely related to changes in soil organic matter.
- Can we use what we learn to further enhance restoration success?
  - Potentially, but it is not straightforward and may be both species- and site-specific

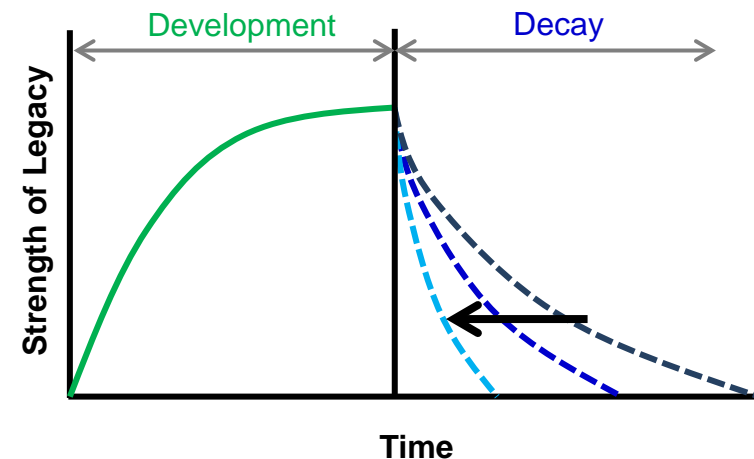


# Different approaches overcome legacies to different degrees



# When should we address legacy effects?

- Amount of effort should be based on the balance of resources required, potential efficacy, and degree of need
- Many new studies coming out manipulating microbes and finding improved restoration, but the field is still in its infancy
- General rules?



# Thank you!



- Hawkes Lab
  - Erin Brault
  - Clare Glinka
  - Nick Johnson
  - Ben Sikes
- Archbold Biological Station
  - Eric Menges
  - Hilary Swain
  - Patrick Bohlen
  - Stacy Smith
- USDA NRI Managed Ecosystems Program
- Smith Fellows Program



Smith Fellows

# Four of these species were also differentially sensitive to microbes

