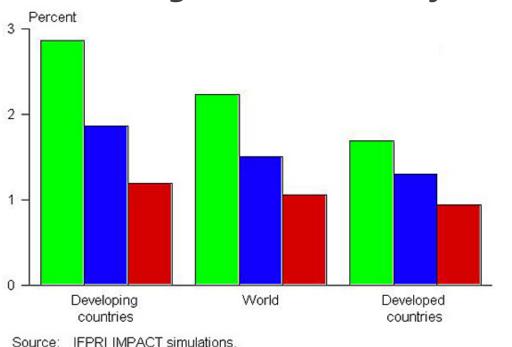






How do we reverse the current *decrease* in annual yield growth?

Annual growth in cereal yields



1967-1982

1982-1994

1993-2020

Need ~2.4% increase per year to feed the global population in 2050

How to feed the world

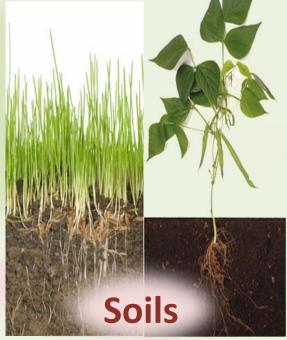
Plant-based Agriculture: A complex system





Viruses
Archaea
Bacteria
Amoeba
Oomycetes
Fungi
Algae
Nematodes

Their environment



Arthropods, Other Animals and Plants



Insects
Arachnids
Myriapods
Worms
Birds
Rodents
Ruminants
Weeds

Associated organisms

Climate

Roots recruit a myriad of microbes

Enhance uptake of nutrients and water

N₂ fixation P solubilization

piremongolia.files.wordpress.com

Enhance tolerance to environmental stresses

Drought, Salinity Heat/Cold, Heavy metals,...

Activate defenses against pests & pathogens

Increase plant health & productivity

Promote plant growth

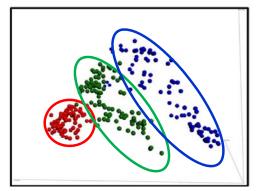
Root recruitment promotes clear community shifts

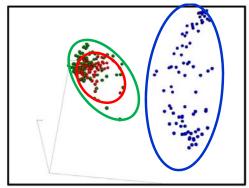
Bacterial Communities Fungal Communities

Soil •

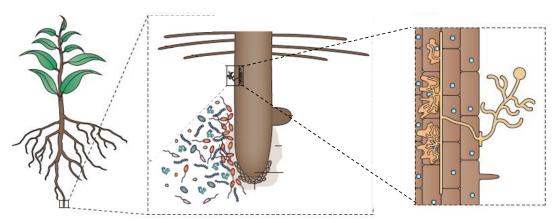
Rhizosphere •

Endosphere •





Soil = Microbial seed bank



Phillipot. 2013. *Nature Reviews Microbiology*

What are the primary factors driving microbiome assembly?

Abiotic factors

Soil properties

pH texture organic content

Climate

Temperature Water availability

Biotic factors

Microbe source seed, soil, air

Plant properties species genotype

age Plant community diversity

Macroorganisms

Protists, nematodes, aphids

- → Identify microbial indicators/predictors of crop and soil health
 - → Maximize the influence of seed microbiome manipulation on plant fitness and productivity

Can we improve plant-microbiome benefits through breeding?

- > Has domestication altered crop mutualisms?
- > Has breeding under ideal plant growth conditions reduced the recruitment of beneficial microbes?



Example: Can we use knowledge of microbiomes to enhance drought tolerance?







(Photo: Joseph Murphy/Iowa Soybean Association)

Soil water content influences community composition

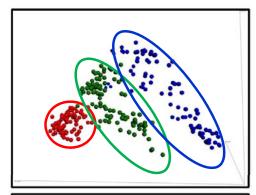
Bacterial Communities Fungal Communities

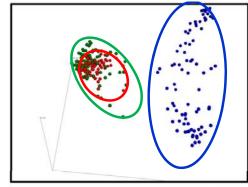
Proximity to the root

Soil •

Rhizosphere •

Endosphere •

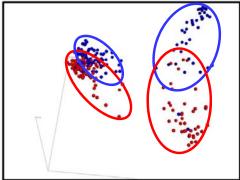




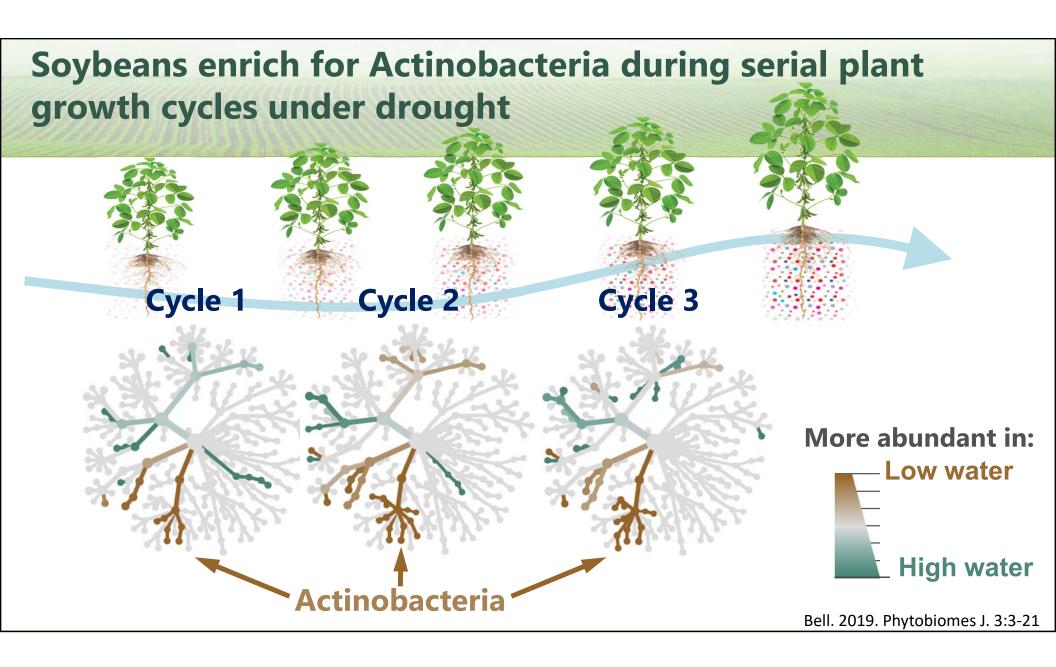
Soil water content

Low •

High •

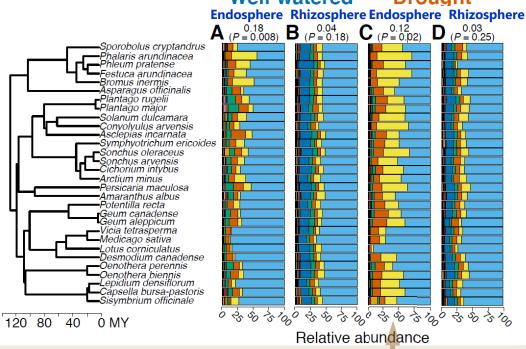


PCoA plots



Many plant species enrich for Actinobacteria under drought



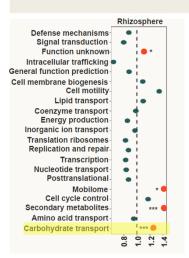


Fitzpatrick. 2018. PNAS 115:e1157

Actinobacteria

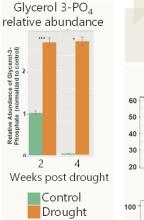
Sorghum

Metatranscriptomics



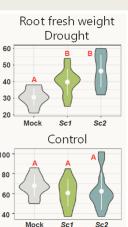
Control Drought





Xu. 2018. PNAS 115:e4284

Isolates



> Can we breed/engineer plants to better recruit beneficial microbes/microbiomes?

> Can we breed microbes/design biologicals to enhance their benefits to host plants?



What are the mechanisms by which specific management practices promote ecosystem health?

Plant diversity

Monoculture
Cover crops
Crop rotations
Intercropping

Inputs

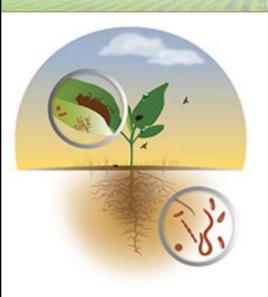
Organic/Inorganic N
Fertilizer rate, timing
Herbicides/Pesticides

Cultural practices

Till/No-till
Plant time
Irrigation/Drainage
Livestock mgt

→ Design novel or improved management practices

Fundamental understanding of plant microbiomes



- How do plant exudates/bacteriophage/ nematodes/protists affect the ecology/evolution/ function/development of plant microbiomes?
- ➤ How do microbes/microbiomes prime the plant immune system against pests and pathogens?
- How do microbe-microbe and microbe-plant-insect interactions influence
 - microbiome diversity, function and resilience? microbiome impact on nutrient cycling and C sequestration?

Common study systems

Model plant species

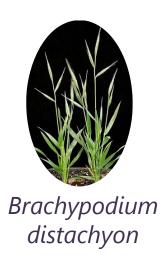




Medicago truncatula



Populus trichocarpa



Major crop species



Maize



Wheat/Barley



Soybean



Rice

Challenges in plant microbiome studies

Large number of plant species (~7,000 cultivated crop species)* and varieties (cultivars, landraces, elite varieties) of each

- ❖ Extensive permutations of Plant genotype x Environment x Management x Microbiome interactions → affect repeatability, generalizability and experimental sampling
- ❖ Wide diversity of spatial scales (µm → field → region
 → planet) and temporal scales
- High environmental heterogeneity

*Khoshbakht. 2008. Genet Resour Crop Evol 55:925

Technology needs

- Approaches to separate microbial metagenomes/ metatranscriptomes from host DNA/RNA
- Improved taxonomic identification and databases that allow for the integration of distinct information
- Whole genome-based identification and associated impacts on plants, animals and the environment for risk assessment and permits

Opportunities presented by plant microbiome studies

- * Experimental control, sample accessibility, high sample numbers
- Availability of long-term sites with known historical management
- Opportunities for comparisons such as evolutionary (e.g., pre-vs post-domestication) and ecosystem (agriculture vs. natural)
- Short path to translate fundamental findings into application
- Strong infrastructure for dissemination of applications & opportunities to engage farmers

Broad relevance of plant microbiome studies

- Sustainable crop production for food, feed and fiber (bioenergy)
- Environmental health (phytoremediation, forest health)
- Human health (human pathogens on plants, digestibility)
- Climate change mitigation (N cycling impacts, C sequestration)
- Rehabilitating degraded and depleted lands

*1.5 billion people depend on degraded lands for survival!



Funding & Organizations

Primary US funding

\$\$\$ \$\$\$ \$\$\$

USDA NSF DOE

Commodity groups
Private industry
Nonprofit foundations
(FFAR, Noble)





American Phytopathological Society

American Society for Agronomy Crop Science Society of America Soil Science Society of America American Society of Plant Biologists





Private sector companies (agricultural biologics, seeds, crop protection chemicals)

International Alliance for Phytobiomes Research Sponsors













































