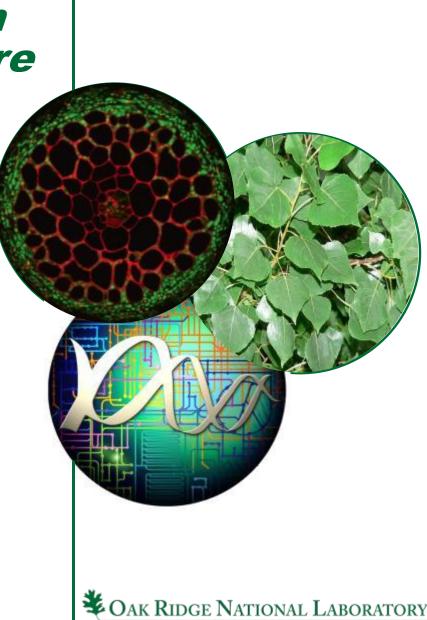
Fungal diversity within the Populus rhizosphere and endosphere

Gerald A.Tuskan Oak Ridge National Laboratory

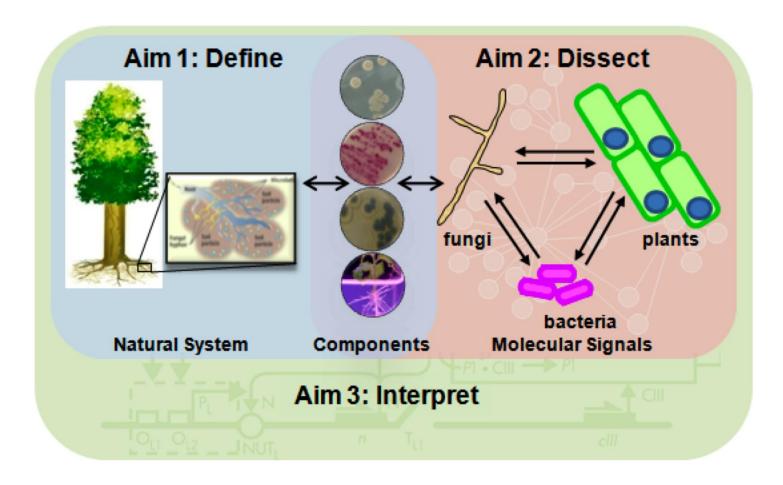
Jessy Labbe Nora Dunkirk Pascale Frey-Klett Aurelie Deveau Dale Pelletier Gregory Bonito Rytas Vilgalys





MANAGED BY UT-BATTELLE FOR THE DEPARTMENT OF ENERGY

Defining, dissecting and interpreting the plant-microbe interface



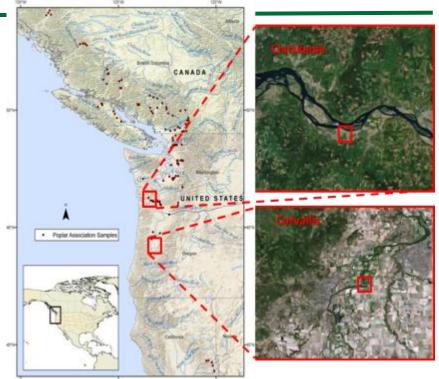


Next steps in defining the *Populus* microbiome

- Common garden studies
 - Partitioning of genotype by environment interactions through host resequencing and community metagenomics (JGI Collaboration)
- Understanding how members of the community function in concert
 - Metagenomic and metaproteomic measurements
 - Isolation of missing phyla
 - Model community studies

Beginning studies on well characterized common-garden populations that will allow examination of more detailed genomic-level information in a environmental setting





Ectomycorrhiza Bioassay Experiment

- Allows experimental assessment of what portion of the available arbuscular (AM) and ectomycorrhizal (ECM) population, is capable of forming relationships with *Populus* roots of different genotypes (e.g., soil inoculum potential x genotype effects)
- Serves as source for ECM isolation & cultivation efforts in Task 1.3.1
- 5 genotypes of *P. deltoides*, 1 *P. dxt h*ybrid, oak/pine & *Tuber* positive controls, are being used to access 5 different soils in the first experiment

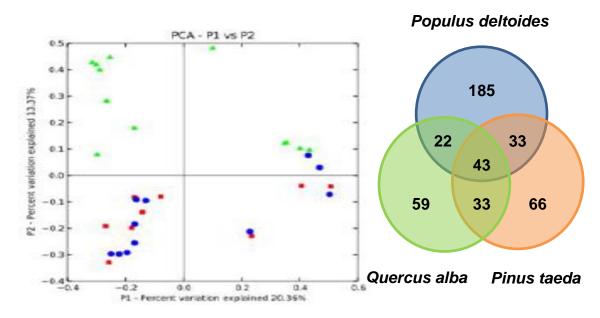


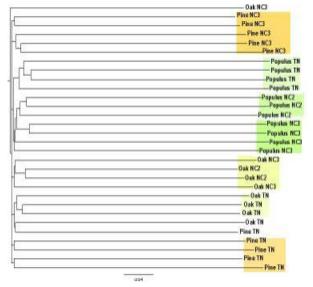


J. Labbe, G. Bonito & R. Vilgalys

Trap-Plant Experiment 1

Main finding: *Populus deltoides* hosts distinct endosphere fungal and bacterial communities from *Quercus alba* and *Pinus taeda*





Fungal communities on three hosts

Fungal species (OTUs) shared between 3 hosts

Unifrac distances of bacterial communities based on 10248 16S sequences

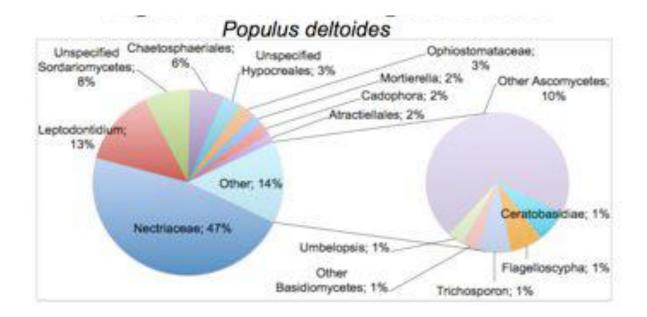


Bonito et al. In prep

PMI Fungal Culture Collection

>1800 isolates ITS DNA barcoded Selected Taxa for:

- Koch's postulates
- Characterization
- Genomics
- Transcriptomic
 Experiments



| Ab | P. Deltoides | Shared | P. trichocarpa |
|-------|--------------------|----------------|------------------------|
| | Chaetosphaeriaceae | Cylindrocarpon | Ilyonectria radicicola |
| | Ophiostomataceae | Leptodontium | Phomopsis |
| | Atractiellales | Ilyonectria | Corticiaceae |
| ŀ | Denodrosporium | Neonectria | Hypocrea |
| | Thozetella | Fusarium | Paecilomyces |
| Dance | Trichosporon | Exophiala | Chaetomiaceae |
| | Aspergillus | Mortierella | Cryptosporiopsis |
| | Phoma | Lecythophora | Bionectria |
| | Chaetosphaeria | Hymenoscyphus | Phialophora |
| | Diaporthales | Mortierella | Aureobasiidum |



Koch's Postulates

- 1) Causal organism must be associated with disease
- 2) Causal organism must be isolated from an infected plant & grown in culture
- 3) When a healthy susceptible host is inoculated with the pathogen from pure culture, symptoms of the original disease must develop
- 4) The same organism must be re-isolated from plant



Grow fungus on millet



Inoculate plants with colonized millet

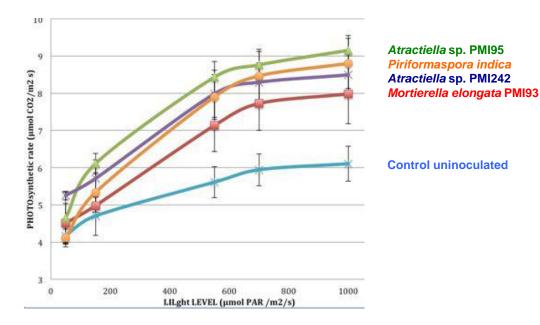


Re-isolated fungus, characterize fungal and plant growth and health



Effects of fungal endophytes on plant growth & photosynthesis

Mean photosynthesis of oaks inoculated with root endophytes



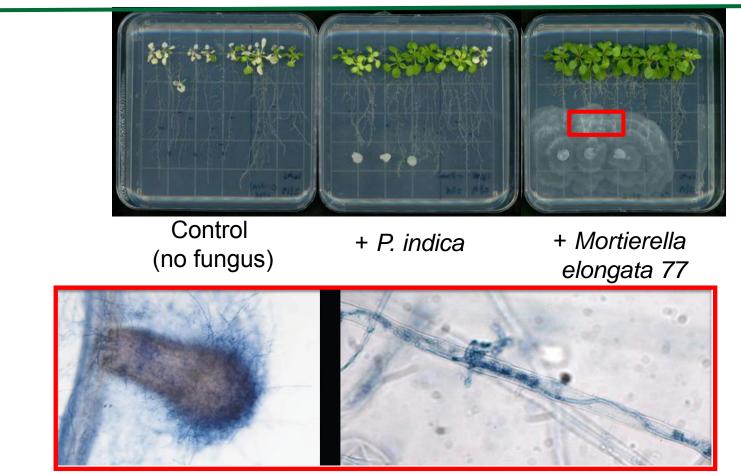
Atractiella affects leaf area of red oak



Control uninoculated PMI95 PMI242 PMI152



Effect of fungal colonization on heat shock tolerance



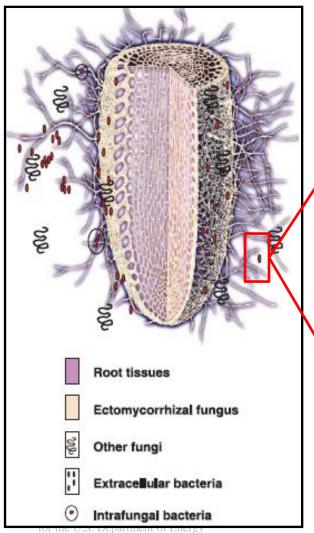
Mortierella elongata colonizing Arabidopsis roots

Jeremy Lipkowitz (Duke Univ.)



Experimental characterization of Mycorrhizal Helper Bacteria (MHB)

Goal: identification of the molecular determinants of the helper effect



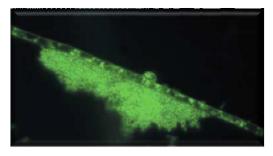


Electron microscopic image

Mycorrhizal fungi are surrounded by complex microbial communities, which modulate the mycorrhizal symbiosis.

'Mycorrhiza helper bacteria' (MHB) assist mycorrhiza formation,

and interact positively with the functioning of the symbiosis.



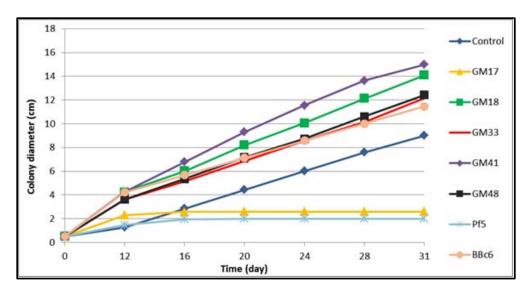
Epifluorescence microscopy of biofilm on fungal hyphae

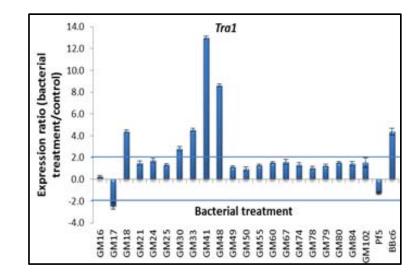


Frey-Klett et al 2005

Summary of MHB Characterization

MHB strains of the Laccaria-Populus interaction among the native Populus deltoides microbial communities





Microscopic observation of L. bicolor mycelium (10X) 12 DPI



BBc6 effect +

11 Managed by UT-Battelle

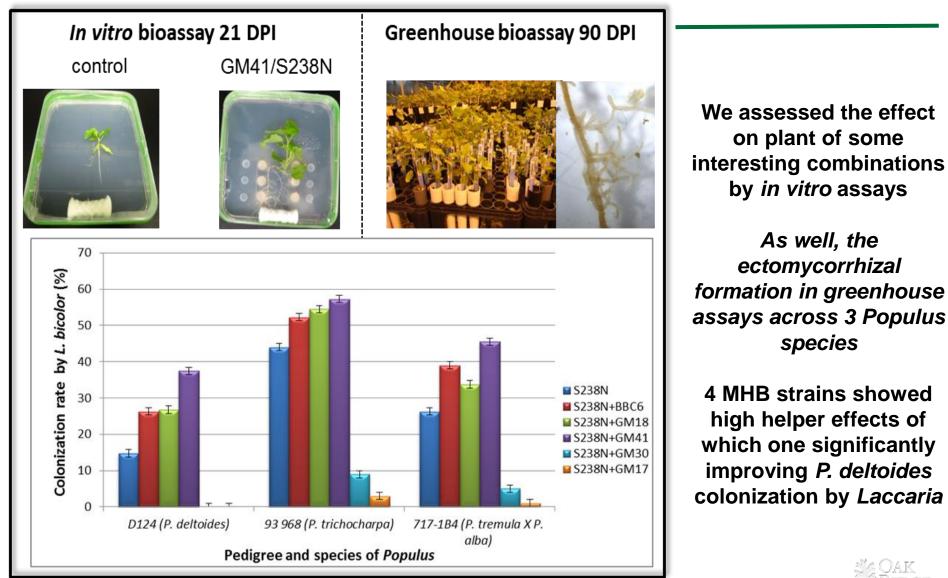
GM41 effect ++

We looked at the bacterial effect on Laccaria growth and mycelium morphology (branching and density)

and

the expression level of seven target genes in L. bicolor shown as regulated by the MHB strain BBc6R8

Summary of MHB Characterization - con't



for the U.S. Department of Energ

Evaluating the characterized MHB strains effect on other Fungi isolates

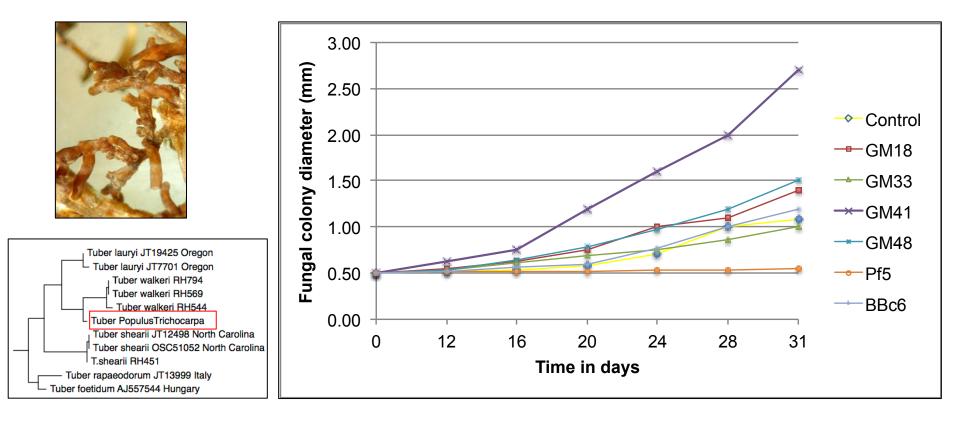
(experimental evaluation done on 20 fungal strains)



Look at the fungal growth and the branching density over a month of culture



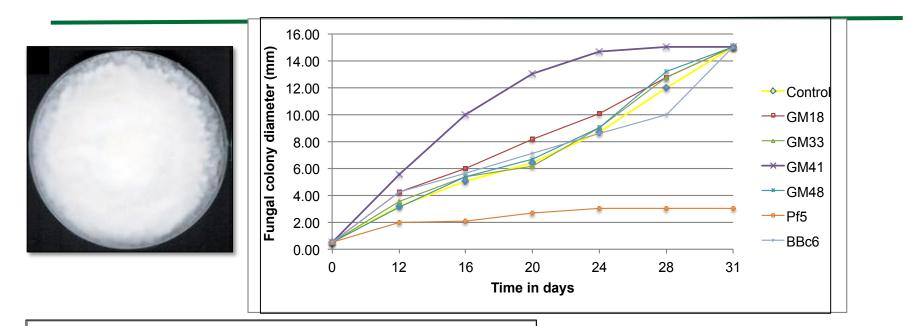
Focus on the new isolated species of Tuber (not named yet)

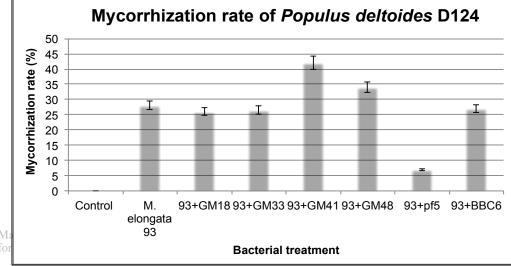


The *Pseudomonas fluorescens* strain GM41 has a significant effect on the growth of the mycelium (ANOVA, p<u>></u>0.01)



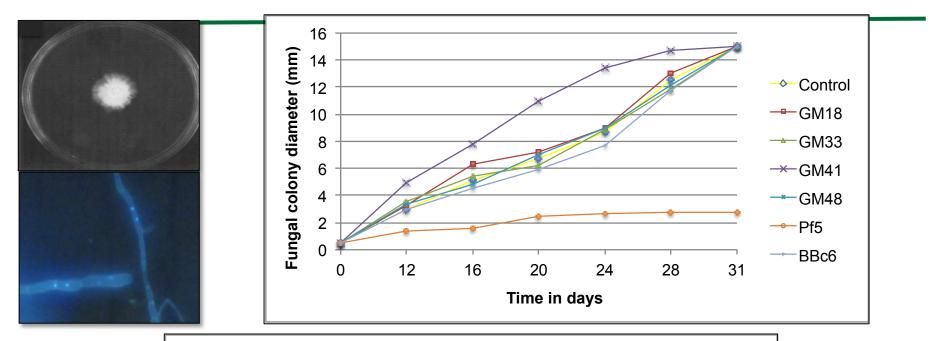
Focus on the sequenced *Mortierella elongata* strain #93

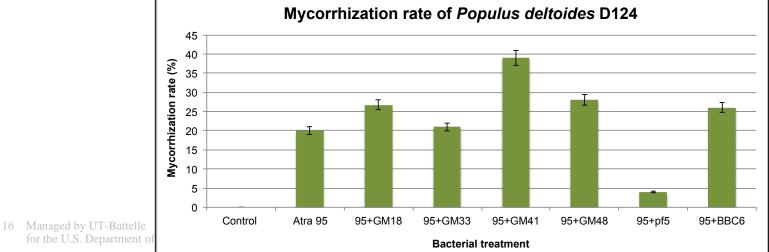




The *Pseudomonas fluorescens* strain GM41 has a significant effect on the growth of the mycelium and root colonization as well (ANOVA, p≥0.01)

Focus on the sequenced Atractiella sp. strain #95



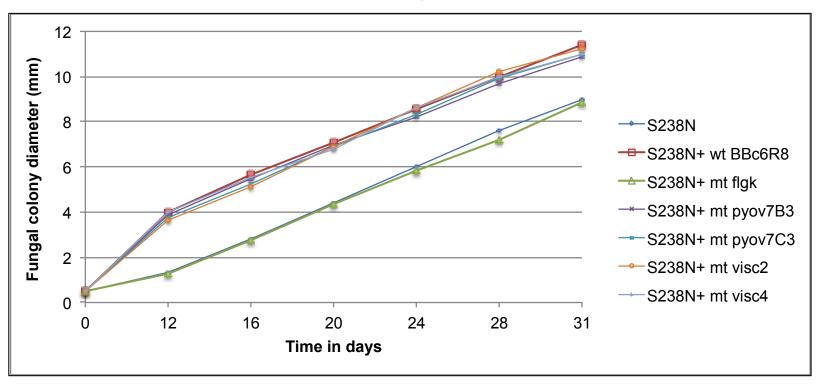


OAK RIDGE

Transition to the molecular determinants of MHB

Effect of the BBc6R8 mutants on the L. bicolor S238N

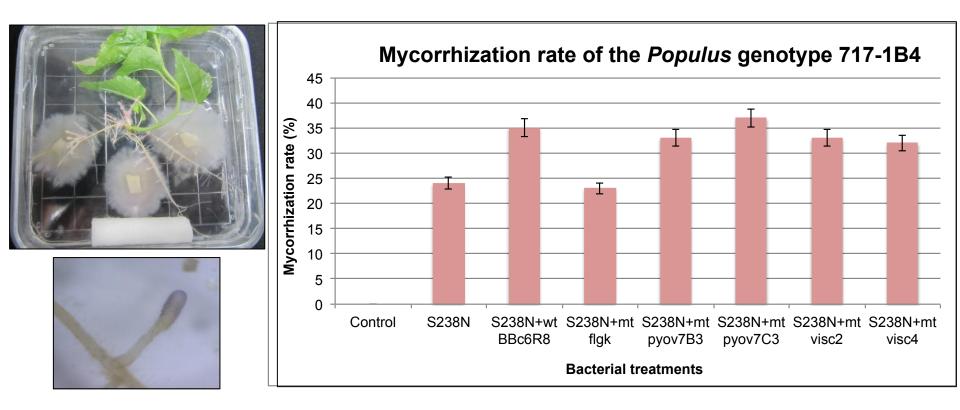
radial growth



The *P. fluorescens* BBc6R8 mutant $\Delta flgk$ has a significant impact on the Laccaria growth (ANOVA, p \geq 0.01)



Effect of the BBc6R8 mutants on the colonization of *Populus* by *Laccaria* (*in vitro*)



The *P. fluorescens* BBc6R8 mutant *∆flgk* has a significant effect on the helper ability (ANOVA, p≥0.01)



Acknowledgments

About Us



opulus root sampling in Clat

Plant-Microbe Interfaces Scientific Focus Area

INRA

The Plant-Microbe Interfaces (PMI) project is a Scientific Focus Area directed towards understanding the dynamic interface that exists between plants, microbes and their environment. A specific focus is on defining the genetic bases of molecular communication between Populus and its microbial consortia. Understanding the inherent chemical and physical processes involved will facilitate natural routes to the cycling and sequestration of carbon in terrestrial environments, ecosystem response to climate change, and the development and management of renewable energy sources.

The project integrates expertise in the areas of plant genomics, longal and bacterial research, longal acclogy, analytical tool development and computational biology and is based at the Oak Ridge National Laboratory, with collubrators at the University of Washington, Dulke University, and IRRA- Hanny (Trance). The project is a Foundational Genomics Scientific Focus Area supported by the Genomic Science Program of the OBCe of Biological and Environmental Research of the U.S. Department of Energy.



pmi.ornl.gov



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