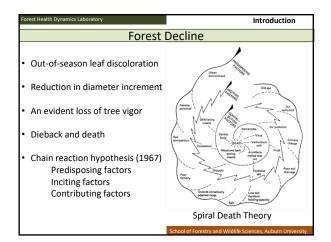
Variation in Tolerance of Several Loblolly Pine Families to *Leptographium terebrantis* and *Grosmannia huntii*

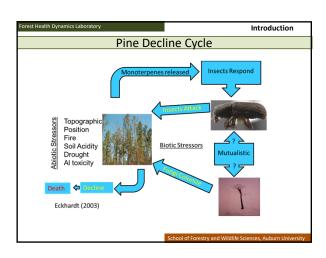
¹Pratima Devkota, ²Scott A. Enebak, ²Joseph W. Kloepper, ²Ryan L. Nadel, ²Zhaofei Fan, ²Lori G. Eckhardt

¹Plant Biology Laboratory, Michigan State University, Michigan, ²School of Forestry and Wildlife Sciences, Auburn University, Auburn, Alabama

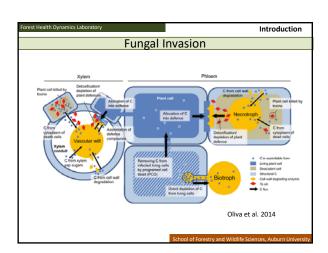
Forest Health Dynamics Laboratory













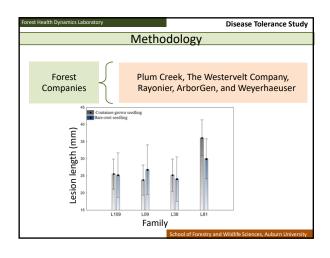
Motivation Published reports Pinus taeda is relatively more susceptible to fungi among the southern pines (Matusick et al. 2010) Intra-specific variation in tolerance of P. taeda to ophiostomatoid fungi (Singh et al. 2014) Resistance to other disease like fusiform rust has been incorporated in breeding (20 - 25% infection reduction) (Liu et al. 1999)

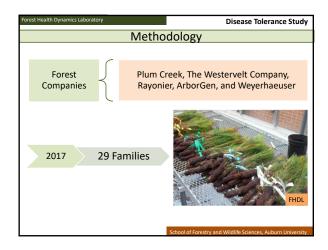
Research gap

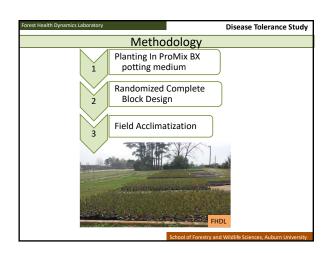
 Need for further screening of more commonly out-planted P. taeda families

Cabanda Signatura and Marildinia Cabanda Andreas Marildinia

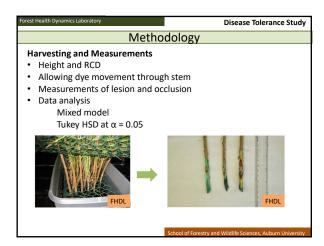
Objectives To determine intra-species variation in tolerance of loblolly pine to Leptographium terebrantis and Grosmannia huntii To determine the yearly performance of connector families



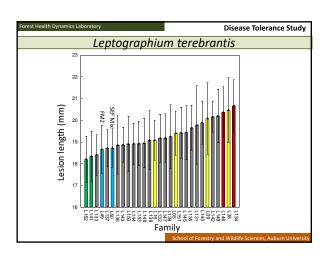


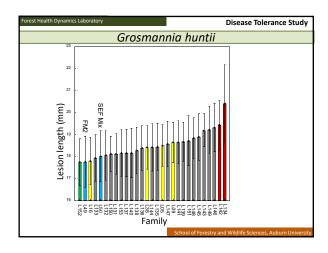


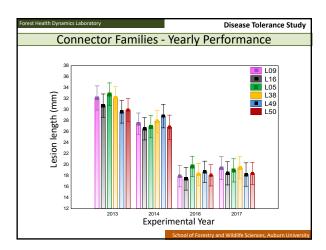




est Health Dynamics Laborator	У	Disease Tolerance Study
	Results	
Effect	P-value (lesion length)	P-value (occlusion length)
Family	0.001	0.045
Treatment	<0.0001	<0.0001
Family*Treatment	0.991	0.259
	Sahaa	l of Forestry and Wildlife Sciences, Auburn University

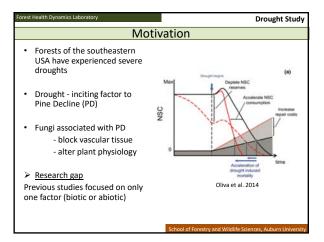






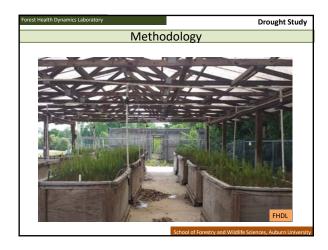
Conclusion Intra-specific variation in tolerance to L. terebrantis and G. huntii Great potential for selection of tolerant families to mitigate potential problem due to decline Genotype and environment (GXE) interaction more likely Relative susceptibility of the families remain same

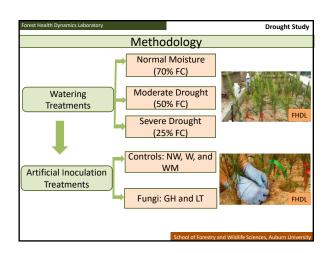
The Role of *Leptographium terebrantis* and *Grosmannia huntii* Invasion in Driving Drought-Related Decline in Loblolly Pine

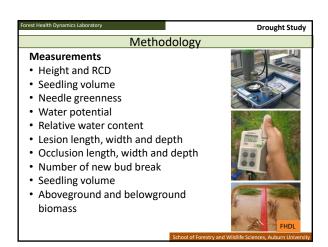


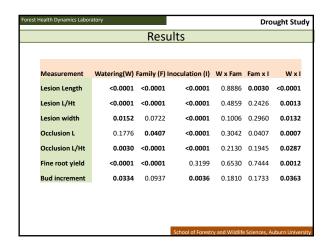
To determine the combined effect of the drought and ophiostomatoid fungi on loblolly pine To understand how loblolly pine families susceptible and tolerant to ophiostomatoid fungi will respond to *G. huntii* and *L. terebrantis* under drought conditions

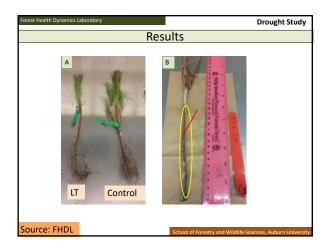
Drought Study

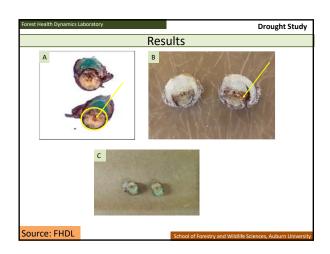


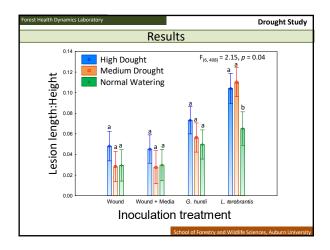


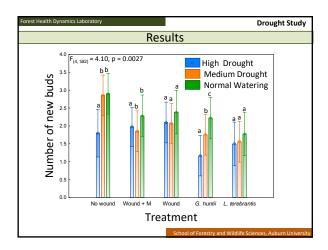


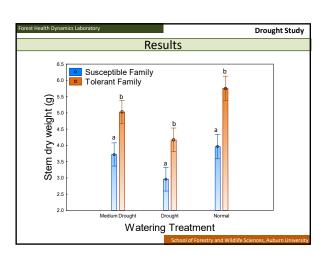








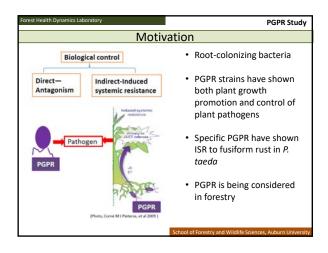




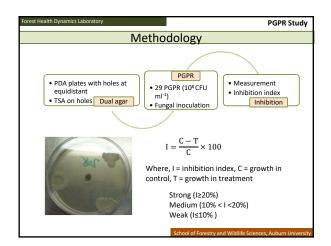
Conclusion Tolerant families tended to be more productive under severe droughts Virulence of two fungi increase under drought through different mechanisms Drought and fungal interactions impact lesion size, fine root formation, and new bud-break Trade off between adaptive traits and seedling growth Interactions put health of P. taeda at risk

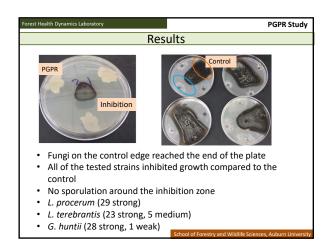
School of Forestry and Wildlife Sciences, Auburn Universit

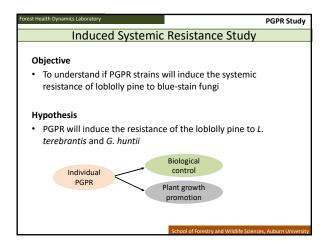
Driving Towards Biocontrol of Ophiostomatoid Fungi By Plant GrowthPromoting Rhizobacteria

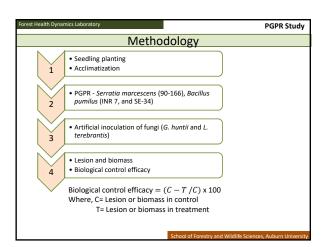


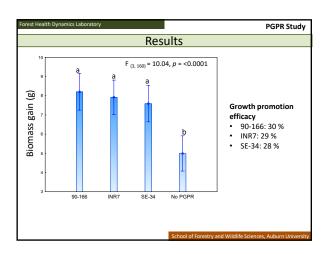
Forest Health Dynamics Laboratory	PGPR Study
Antibios	is Study
Objective To understand if specific strain	
	School of Forestry and Wildlife Sciences, Auburn University

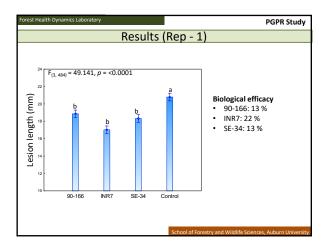


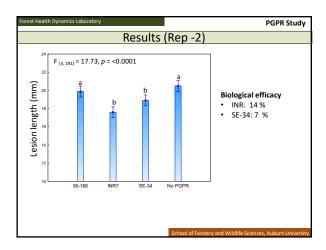








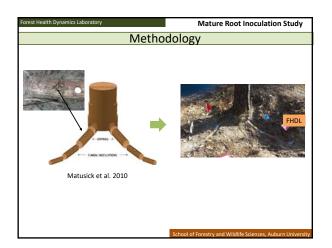


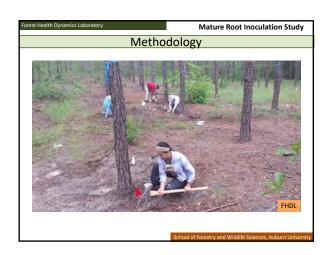


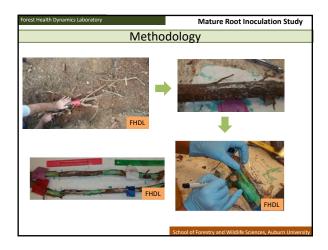
Forest Health Dynamics Laboratory	PGPR Study
Concl	usion
 Most of the studied PGPR strain growth of the fungi <i>in vitro</i> Fungal sporulation is inhibited 	ns have the ability to inhibit the
 PGPR may produce some meta growth of the fungi 	bolites which inhibit the
 Specific strains of PGPR promote PGPR strains may reduce the full 	1 0
	School of Forestry and Wildlife Sciences, Auburn University

Faced Haalib Dunamics I shareton	1
Forest Health Dynamics Laboratory Disease Tolerance Study	
Intra-Species Variation in Response of	
Mature Loblolly Pine to Root-Infecting	
Ophiostomatoid Fungi	
	-
School of Forestry and Wildlife Sciences, Auburn University	
Forest Health Dynamics Laboratory Mature Root Inoculation Study	
Motivation	-
> Previous studies	
 Intra-specific variation in tolerance of P. taeda to ophiostomatoid fungi (Singh et al. 2014) 	
• Considerable variation in susceptibility to nitch confer within	
Considerable variation in susceptibility to pitch canker within mature <i>Pinus</i> species (Rice et al. 2007)	
Decearch gan	
Research gap Screening conducted in seedling stage	
Performance of mature trees in an ecological scenario	
renormance of mature trees in an ecological sections	
	-
School of Forestry and Wildlife Sciences, Auburn University	
Forest Health Dynamics Laboratory Mature Root Inoculation Study	
Objective and Hypothesis	
Objective To determine the intra-species variation in tolerance of mature <i>P</i> .	
taeda families to root-infecting ophiostomatoid fungi	
Hypothesis	
Intra-species variation in tolerance of <i>P. taeda</i> to ophiostomatoid fungi is independent of the plant growth stage	
Calcad of Favorious and William Colons as Assessed to the	
School of Forestry and Wildlife Sciences, Auburn University	

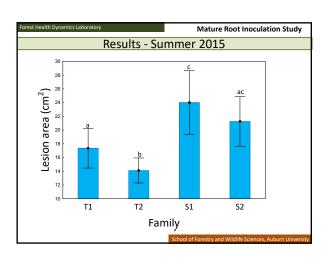
Mature Root Inoculation Study Methodology Summer 2015 and Spring 2016 in 3 locations • Eufaula AL (2 sites), Georgetown GA, and Cordele GA Four families of *P. taeda*• Two susceptible and two tolerant to ophiostomatoid fungi 25 *P. taeda* trees per family (17 year old)



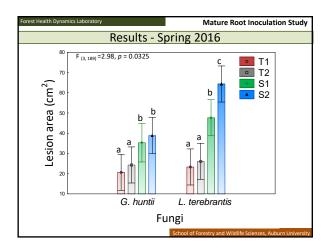




Health Dynamics Laboratory			Mature Root Inoculation S		
Results					
Time	Variable	DF	Lesion Area	Lesion Length	
Summer	Root Diameter	1	0.01	0.01	
2015	Family	3	<0.0001	0.31	
	Treatment	1	0.08	0.1	
	Family * Treatment	3	0.85	0.46	
Spring	Root Diameter	1	0.03	0.86	
2016	Family	3	<0.0001	<0.0001	
	Treatment	1	0.002	<0.0001	
	Family * Treatment	3	0.03	0.32	







Conclusion Intra-specific variation in tolerance of mature families to ophiostomatoid fungi Infection is proportionally larger in mature trees than seedlings Families had same patterns of tolerance independent of the tree growth stage Results support seedling screening studies Shows reliability of seedling screening studies Great potential for selecting fungi tolerant P. taeda families from current planting stock

