

Fall 2014 MI SAF Meeting Update on Breeding for BBD Resistant American beech And Selection & Characterization of "Lingering Ash"

> HOST RESISTANCE AS A TOOL TO COMBAT INVASIVE THREATS

> > Jennifer Koch & Dave Carey Northern Research Station US Forest Service

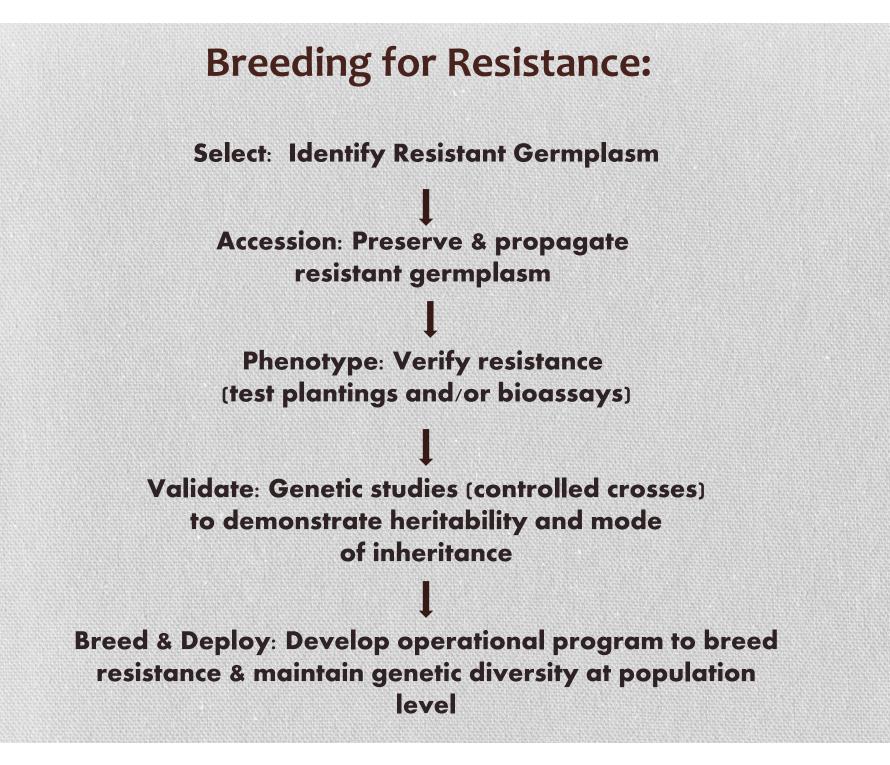
Genetic Variation is the Basis for Tree Improvement

Natural stands have LOTS of genetic diversity

- Rare alleles may confer resistance (or tolerance) even when there is no co-evolution
- Rare because no selective advantage prior to invasive threat

Careful selection and breeding are required to maintain genetic diversity



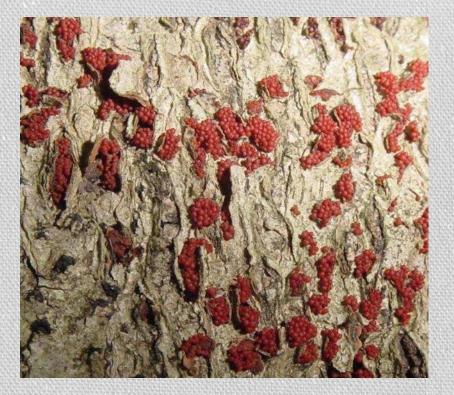


Beech Bark Disease: The Causal Complex



Cryptococcus fagisuga: the beech scale insect

(fruiting bodies photograph from David R. Houston and James T. O'Brien FID Leaflet #75)



Sexual fruiting bodies, or perithecia Neonectria faginata –possibly derived from introduced species Neonectria ditissima-native species

***Resistance is to the scale insect!!!**

Genetic Studies



Full- and Half-Sib Families				
Family	Source			
1505(R) x 1504(R)	Controlled cross			
ME(R) OP	Managed stand No S			
1506(S) x 1504(R)	Controlled cross			
1506(S) OP	Open Pollinated			
1504(R) OP	Open Pollinated			
1510(S) OP	Open Pollinated			





Using the Artificial Inoculation Technique to Screen Seedlings for Beech Scale Resistance



Collect eggs from an infested tree (look up!)



Beech scale adults & eggs.

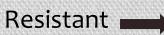
(Derived from D Houston 1982 Res. Paper NE-506)



Scale eggs are placed on foam and tied to the stems of seedlings.

Eggs hatch & colonize susceptible trees.

Artificial Inoculation Technique-One Year Later:







Highly Susceptible

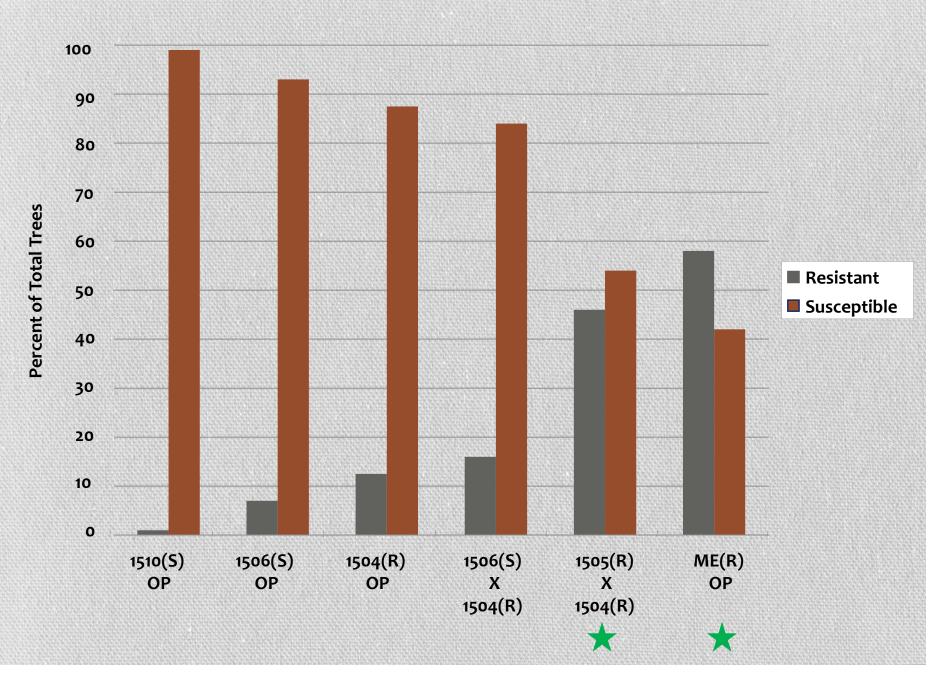
Juvenile nymphs

Susceptible

Adult Scale Insect



Scale Challenge Results



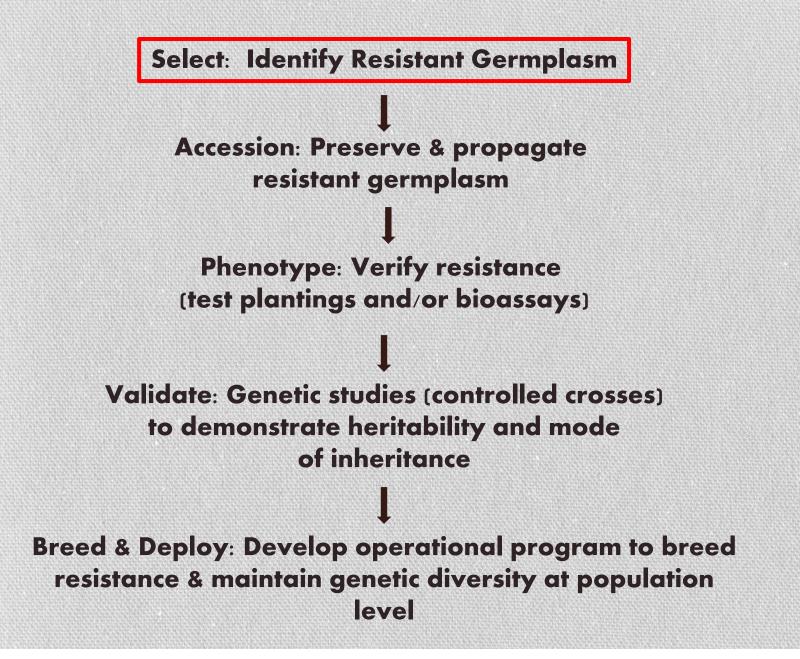


What we have learned about beech scale resistance: (in collaboration with Dana Nelson, SRS)

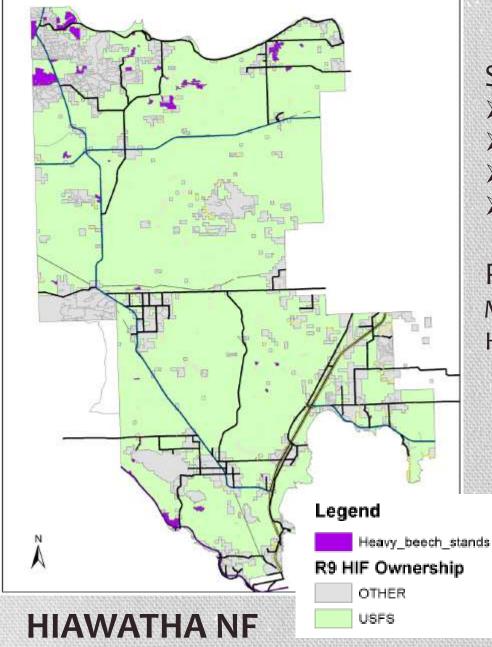
- 1. Confirmed that scale resistance is inherited
- 2. Resistance can be increased through breeding
- 3. Approx. 50 % resistant progeny when both parents are resistant
- 4. Basis for development of American beech seed orchards



Breeding for Resistance:



MI BEECH TREE IMPROVEMENT PROGRAM: SELECTION



SURVEYING

- Focus on areas with high % beech
- Focus on heavily BBD impacted areas
- Look for trees with healthy crowns
- Lack of any insect/disease symptoms

PARTNERS Michigan DNR Hiawatha NF

> Mention Mi DNR resistant beech Reporting form & bring copies

BEECH TREE IMPROVEMENT PROGRAM: SELECTION Field Screen for Scale-Resistance



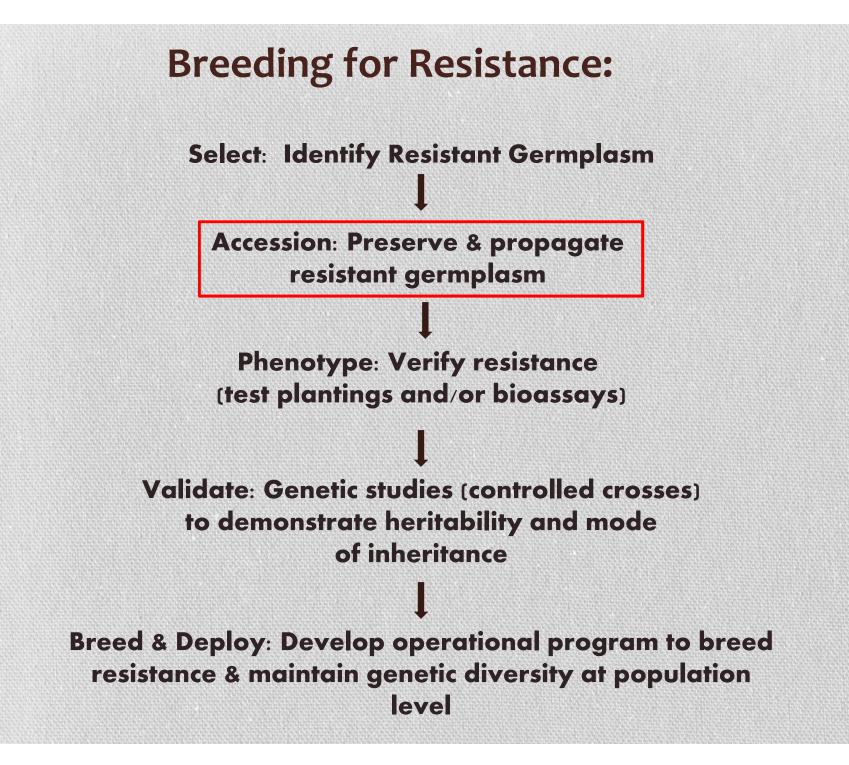
Collect scale eggs



Foam pad with eggs placed against bark and covered with Tyvek

a year, reveals susceptible tree

Koch, Jennifer L.; Carey, David W. 2014. A technique to screen American beech for resistance to the beech scale insect (*Cryptococcus fagisuga* Lind). Journal of Visualized Experiments. e51515. http://www.fs.fed.us/nrs/video/46347/



BEECH TREE IMPROVEMENT PROGRAM: PROPAGATION

- 1. SCION COLLECTION (Partners: MI DNR, Region 9, ORSO)
- 2. SEED COLLECTION & ROOTSTOCK GERMINATION (Partners: NRS, Region 9, ORSO, MI DNR, MSU/MI Tree Cooperative)
- 3. GRAFTING (Partners: NRS, ORSO)



Photo: P.Berrang

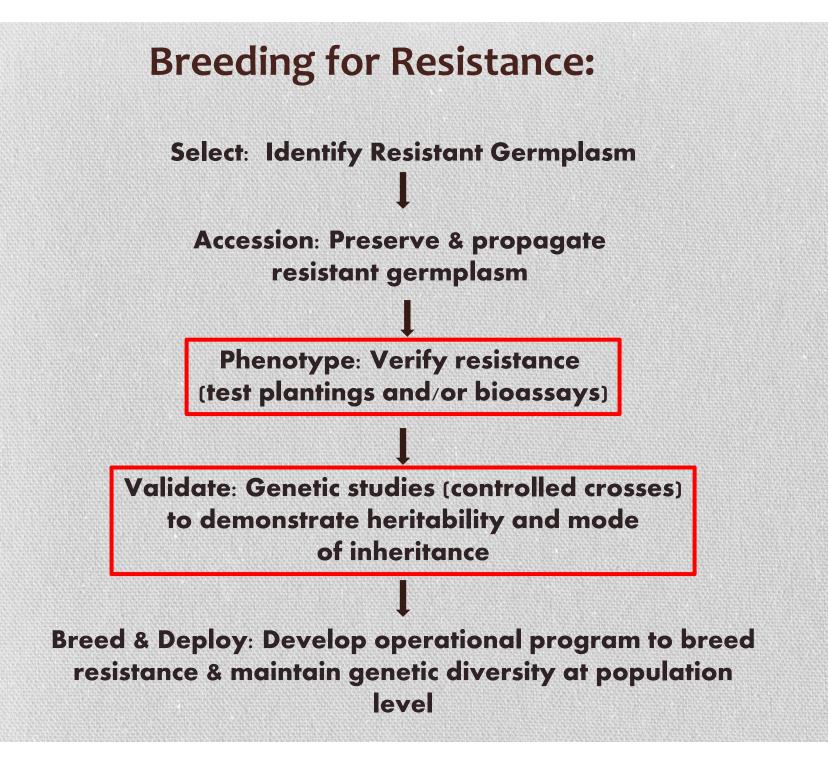


Beech rootstock germinating



Hot-callus beech grafting

Carey, David W.; Mason, Mary E.; Bloese, Paul; Koch, Jennifer L. 2013. Hot callusing for propagation of American beech by grafting. HortScience. 48(5): 620-624.



BEECH TREE IMPROVEMENT: BREEDING & TESTING POTTED SEEDLING/GRAFTED RAMET SCREEN (Northern Research Station, The Holden Arboretum)





Beech scale screening facility The Holden Arboretum



Containerized crosses on grafted ramets

Resistant Susceptible

Containerized Seed Orchard: More Families!

Maternal Parent	Paternal Parent	Family Size	Number of Resistant Progeny	Number of Susceptible Progeny	Proportion of Resistant Progeny
1201	1208	24	12	12	50 %
1202	1208	13	7	6	54 %
1209	1219	21	8	13	38 %
1211	1228	44	23	21	52 %
1228	1211	43	25	18	58 %
1505	ME-85	11	5	11	45 %
1505	ME-23	22	18	4	81 %

Confirms initial results
Possibly different combining abilities
Pilot restoration plantings

BEECH TREE IMPROVEMENT PROGRAM: PROPAGATION

SEED ORCHARD ESTBLISHMENT (Partners: NRS, MI DNR, MSU, HTRIC)

22 BBD Resistant Genotypes

- > 16 from UP State Land
- > 5 from UP Hiawatha NF
- 1 from LP Ludington SP
- 8 ramets per genotype per site = 176

Two Replicate Planting Sites

- Hardwood Tree Regeneration & Improvement Center Purdue University & USFS Northern Research Station West Lafayette, IN Installation fall 2015 (97% of propagation complete)
- Kellogg Experimental Forest
 Michigan State Univerity
 Augusta, MI
 Installation fall 2016 (57 % of propagation complete)

We've continued to perform controlled crosses as opportunities arise.

American beech is monoecious and self sterile.

The goal is to have each parent in at least 2 crosses (daisy chain strategy).

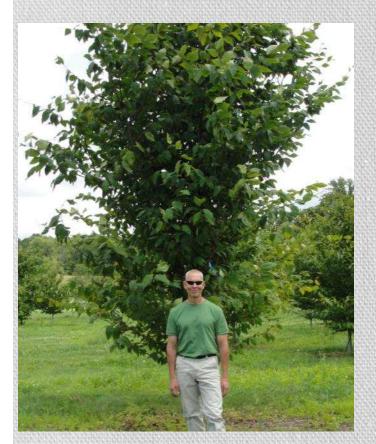




To estimate orchard output we're looking at the proportion of resistant progeny as well as the distribution of the "S" progeny.

Past present and on deck from MI 25 parents 24 mating combinations 470 progeny tested 900 progeny (pending funding) new parents and pairs Families are a resource (mechanisms?)

BEECH TREE IMPROVEMENT: BREEDING & TESTING

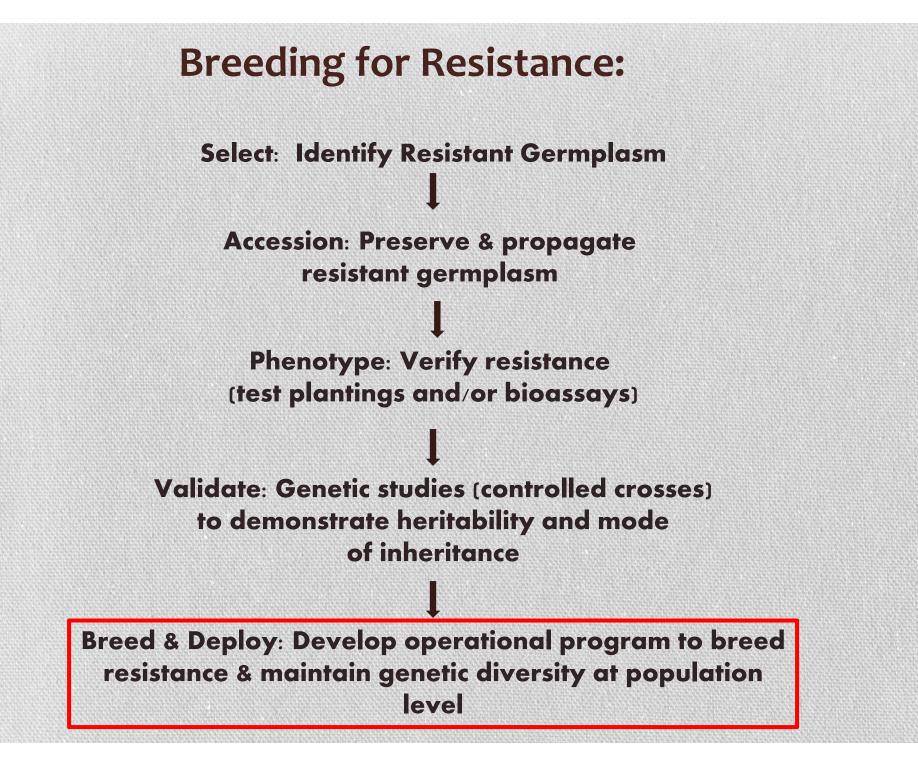




Beech seedling evaluation research planting at The Holden Arboretum



- Evaluate age-age correlation of scale resistance
- Evaluate durability of resistance
- Holden committed to culling susceptible trees and maintaining as a seed orchard



BEECH TREE IMPROVEMENT: OUTPLANTING

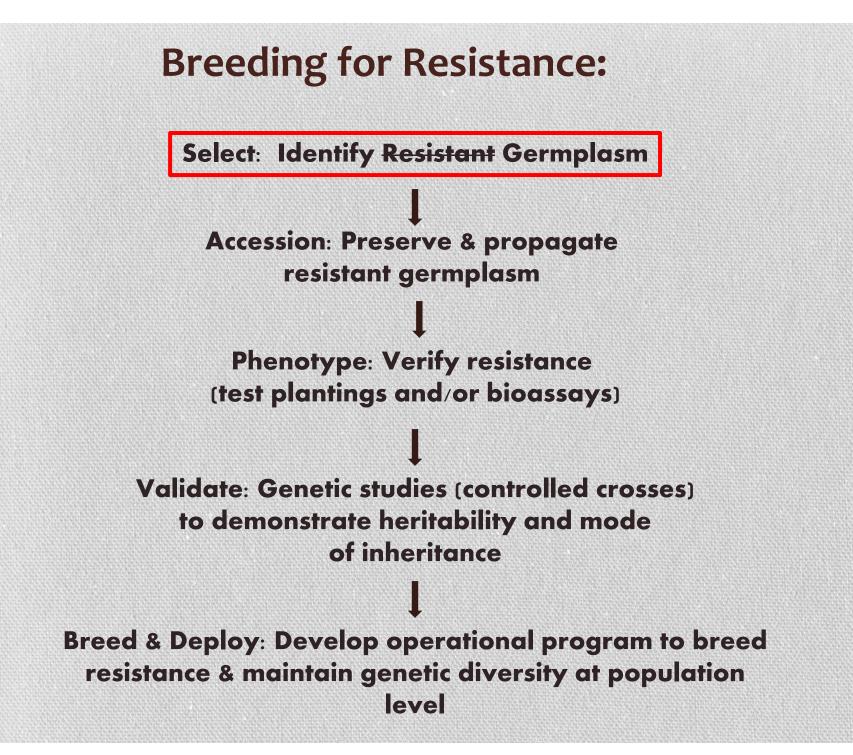
JUST BEGINNING

MI DNR, NRS
Planted 138 seedling in 2011
Over half are BBD-resistant
Survival summer 2013 = 80 %



American Beech Seedling Underplanting, Naubinway, MI





Selecting Lingering Ash in Natural Forest Areas



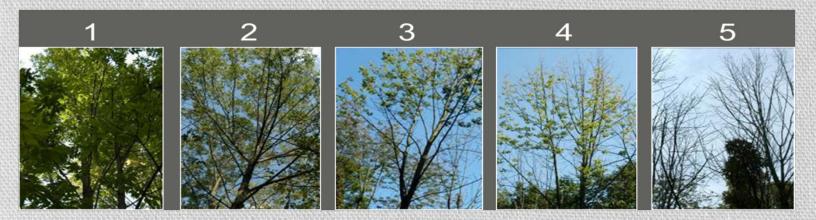
Kathleen Knight, US Forest Service Dan Herms, The Ohio State Univ.

Three main areas:

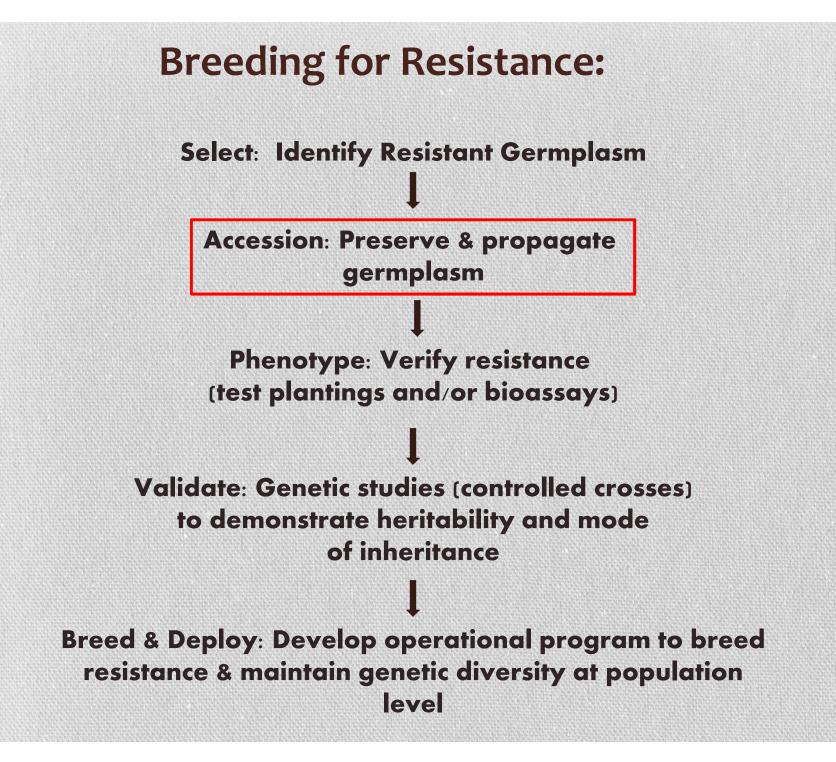
- 1. Long term monitoring plots:
 - Michigan: 33 sites w/ 99 plots
 - Ohio: 50 sites, 165 plots
- 2. FHP sponsored fly-over (MTU study)
- 1. Two Clusters:
 - Oak Openings + Metropark
 - Indian Springs Metropark

SELECTION CRITERIA FOR "LINGERING" ASH

- >95% mortality of mature ash trees has occurred at least 2 years ago
- Tree was large enough(4" dbh) to be infested during the peak EAB infestation
- 3. Tree currently has a healthy canopy (1 or 2) and is at least 10" dbh.



Submit a survivor ash: www.nrs.fs.fed.us/SurvivorAsh



Scion Collection





Lingering Ash Grafting



Grafted clones of a lingering ash selection



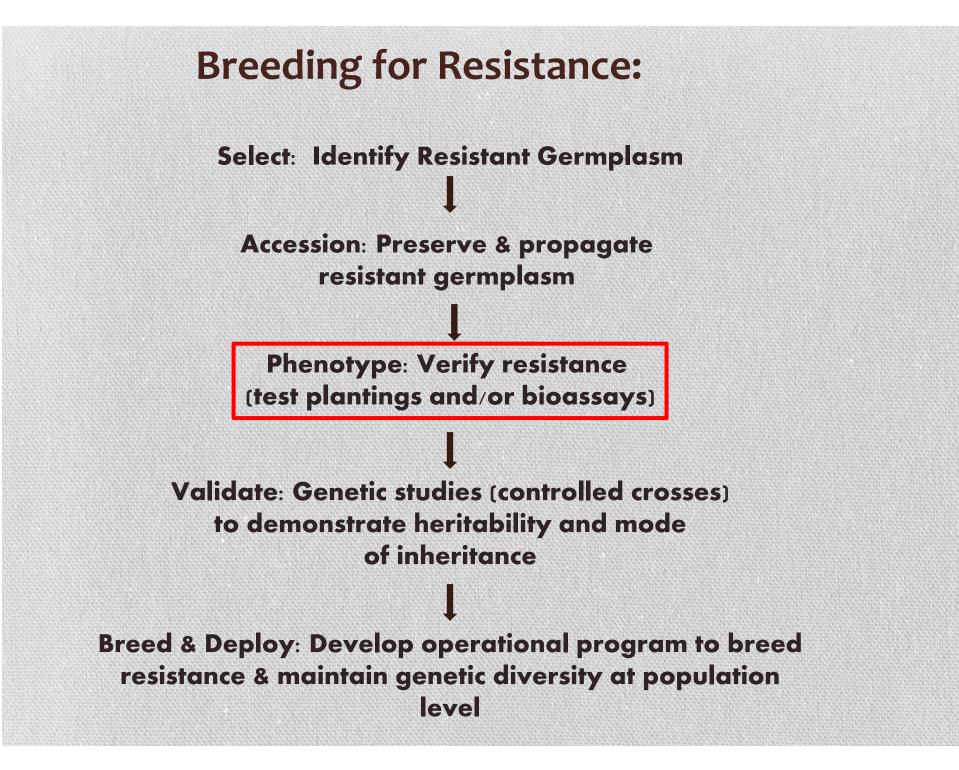
Controlled cross on newly grafted lingering ash

Identification of Resistant Germplasm: "Lingering Ash"



Multiple grafts of each selection will be installed in research plantings

Species	Number of genotypes
Green Ash	40
White Ash	8
Black Ash	3
Unknown (green or white?)	6
TOTAL	57* (21 sites)

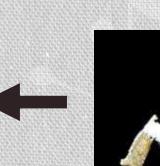


Bioassay Development

- Many points of interaction
- One point may break cycle
- Or better at several points may mean tolerance
- Multiple bioassays may be needed to find and measure resistance







EAB Life Cycle*



Photo by David Cappaert. Reprinted with permission.

Adult Landing and Feeding Preference Collaborator Therese Poland, NRS, East Lansing, MI



experimental cage with leaves and EAB

Development of egg bioassay to confirm resistance:

Destructive sampling 8 weeks After estimated hatch date





Coffee filter with eggs affixed to bark



Healthy larva



Host-killed larva

Development of Egg Bioassay to Confirm Resistance:



Data Collected: -Egg hatched -Larval outcome L1, L2, L3, L4, host-killed

-Larval weight

dead-other

Healthy larva

Host-killed larva

Variation in "Host-Killed" Larvae Phenotypes: EAB killed at entry point



Manchurian Ash



White Ash

Calloused dead-end gallery





"Host-Killed" Larvae Phenotypes: Raised surface galleries

Asian species



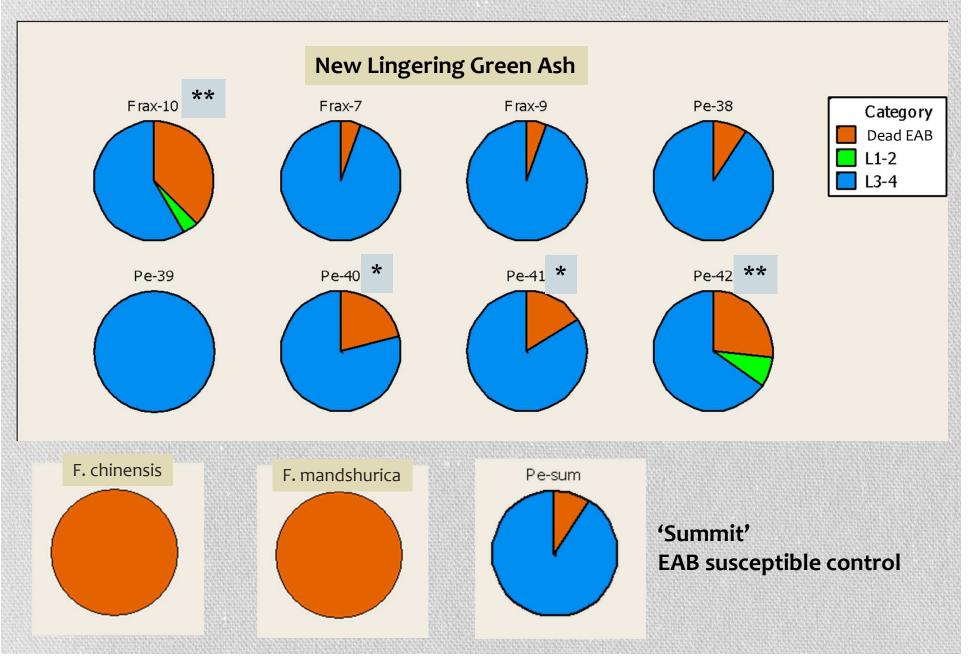
North American species



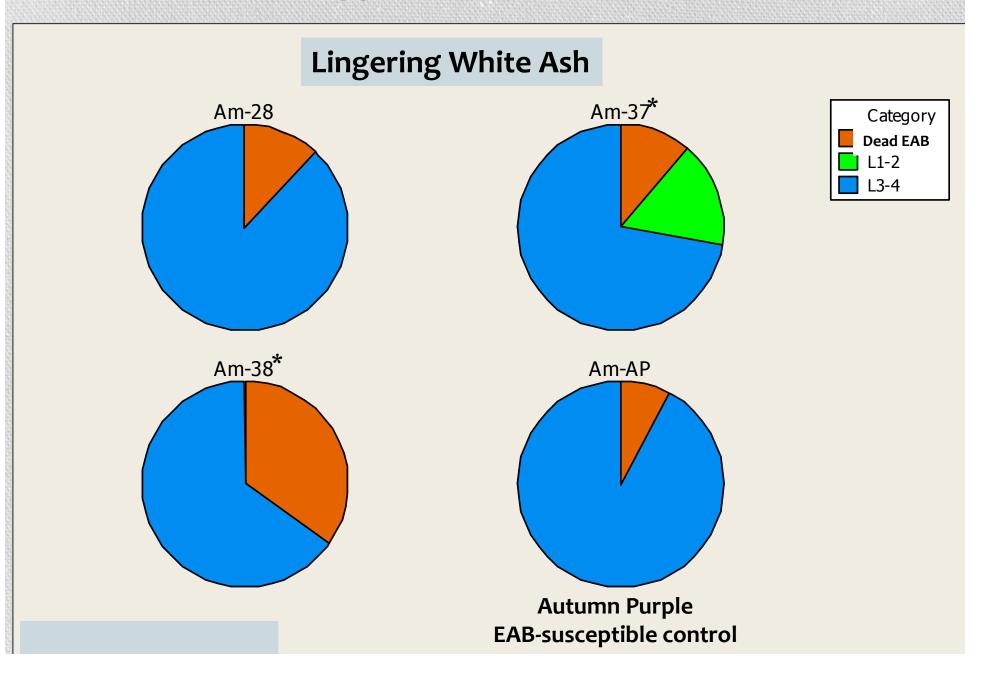
Green ash



2012 Egg Bioassay Results Summary



2012 Egg Bioassay Results Summary



Li	Lingering Ash: Multiple Phenotypes/Mechanisms? 2011 Data Statistical Analysis								
	Genotype	Adult Feeding Preference	Egg Bioassay: Larval Outcome (Dead, L1-4)	Egg Bioassay: Larval Weight	% Stand/Plot Mortality 2012				
	PE-15	SIG*	NA	NA	100 %				
	PE-19	NS†	SIG*	NS†	99 %				
	PE-20	NS	NA	NA	100 %				
	PE-21	NS	NS	NS	100 %				
	PE-22	SIG*	NS†	SIG*	100 %				
	PE-24	SIG*	NS <mark>†</mark>	NS	96 %				

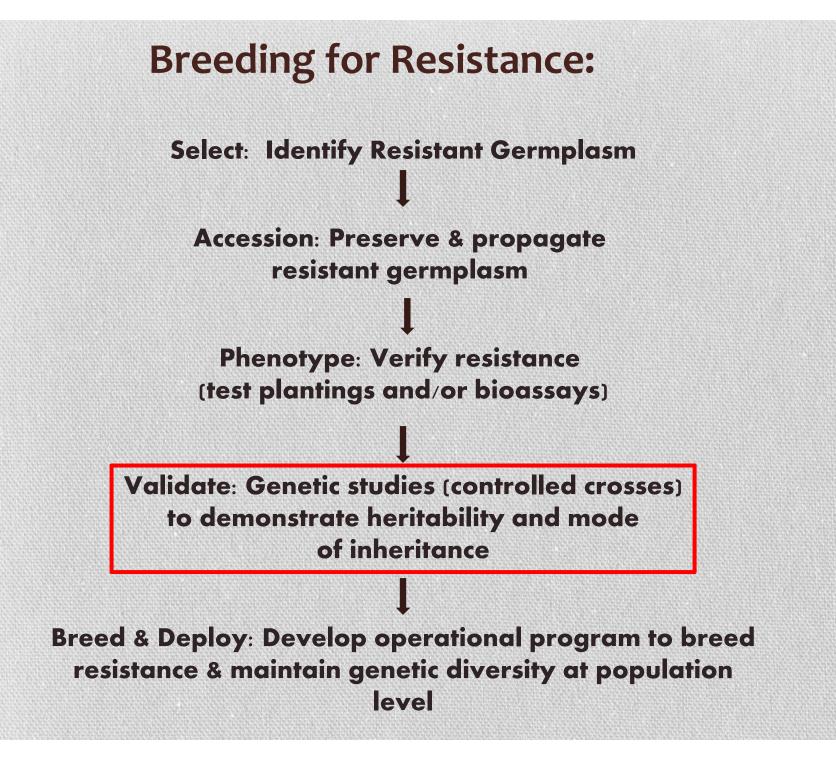
NA=not available NS=not significant

*significantly different relative to EAB-susceptible control

+p-value low but not significant – additional replicates may push toward significance

Field Plantings

- 40 Green + 8 White "Lingering Ash" + controls
 - Dawes Arboretum, Newark OH
 - Holden Arboretum, Kirtland, OH
- Validate initial selection phenotype
- Validate bioassay rankings
- Integrates performance over life cycle
- Analyze genetics x environment interaction



Green & White Ash Genetic Families

> Mapping families for both green & white ash

- Genetic Linkage Maps John Carlson, Penn State University
- Genetic Markers Jeanne Romero-Severson, University of Notre Dame
- NSF Hardwood Genomics Project (hardwoodgenomics.org)
- Lingering x Lingering families
 - Determine if lingering trait is heritable
 - Stack multiple tolerance mechanisms





Koch Group



Ryan Mark Jennifer Mary Dave Reynolds Miller Koch Mason Carey

Funding USDA APHIS USDA NRI Competitive Grants US Forest Service American Recovery and Reinvestment Act (ARRA) Forest Health Management Special Technology Development Program

Thank You!

BBD RESISTANCE WORK:

Region 9, ORSO Paul Berrang, Scott Rogers HNF Majorie Allmaras, Sam Barnes, Tim Baker MI DNR Bob Heyd, Rich Mergerner, Ron Murray, Roger Mech

MSU Paul Bloese

LINGERING ASH WORK:

Toledo Metroparks Huron-Clinton Metroparks MI DNR

Kathleen Knight Lab (USFS, NRS) Britton Flash, Rachel Kappler Dan Herms Lab (OSU) Diane Hartzler Therese Poland Lab (USFS, NRS) Andrew Storer Lab (MTU) Jordan Marshall (Purdue, Fort Wayne)