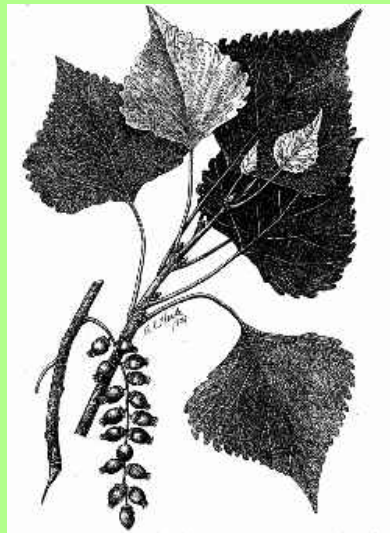


The Importance of Genetic Diversity in Riparian Restoration



Sharon Ferrier, Gery Allan, Laura Hagenauer, Karla Kennedy, Randy Bangert,
& Tom Whitham

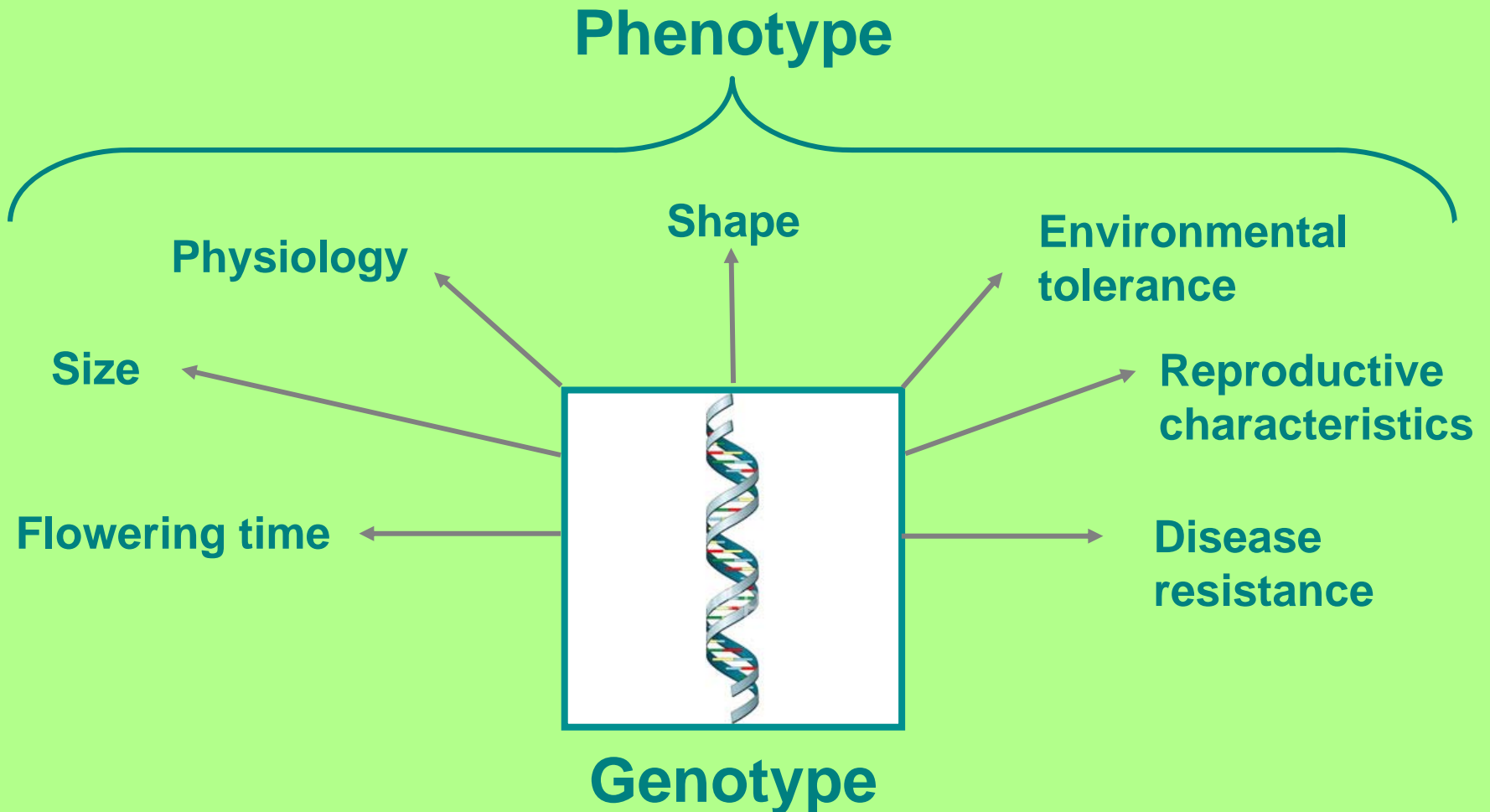
The Cottonwood Ecology Group: www.poplar.nau.edu

Northern Arizona University
Flagstaff, AZ

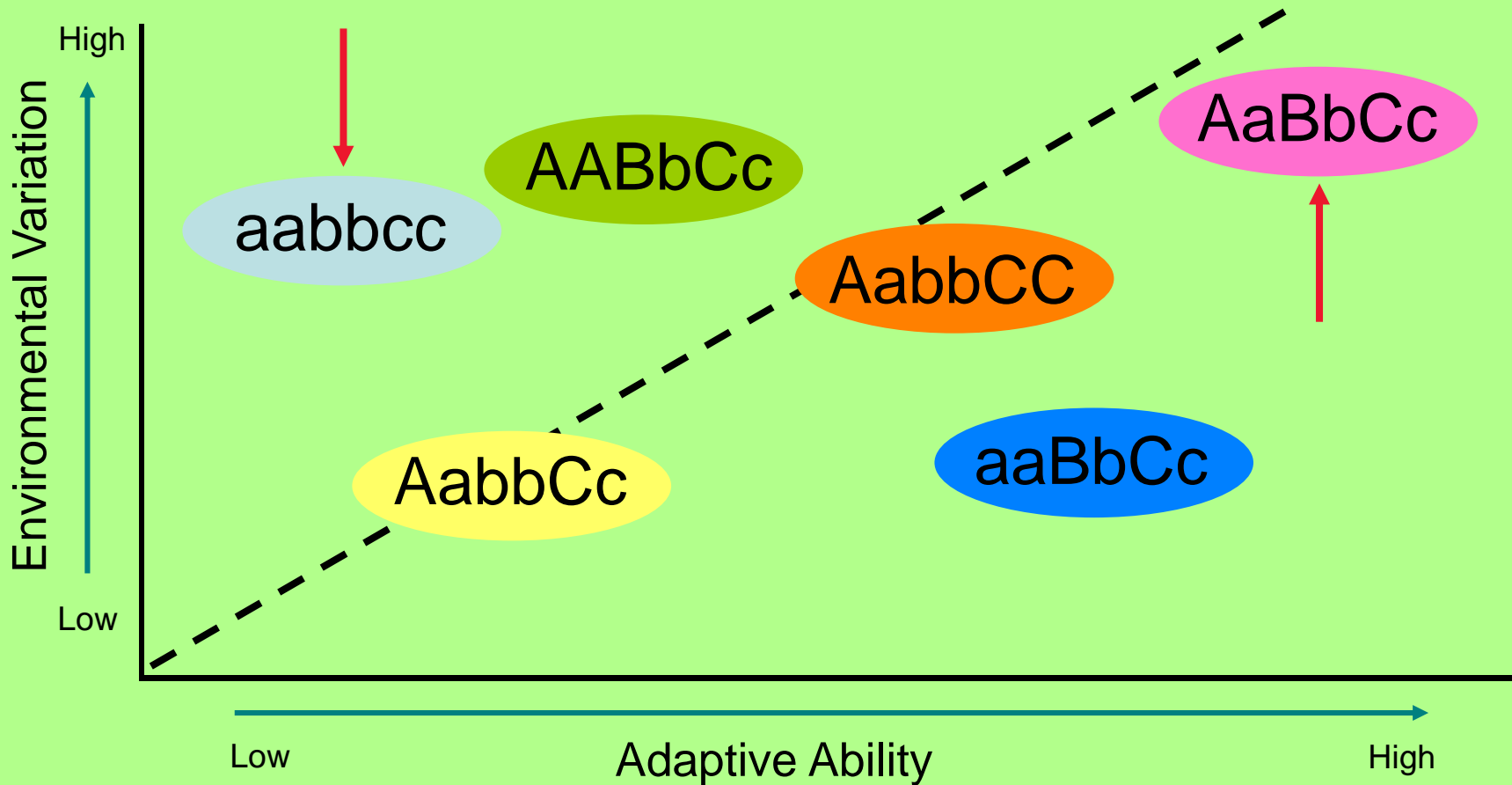


Why use genetic tools?

Genetic information determines form and function



Genotypic diversity influences individual responses to environmental variation and adaptability



Genotypic differences often affect survival and performance

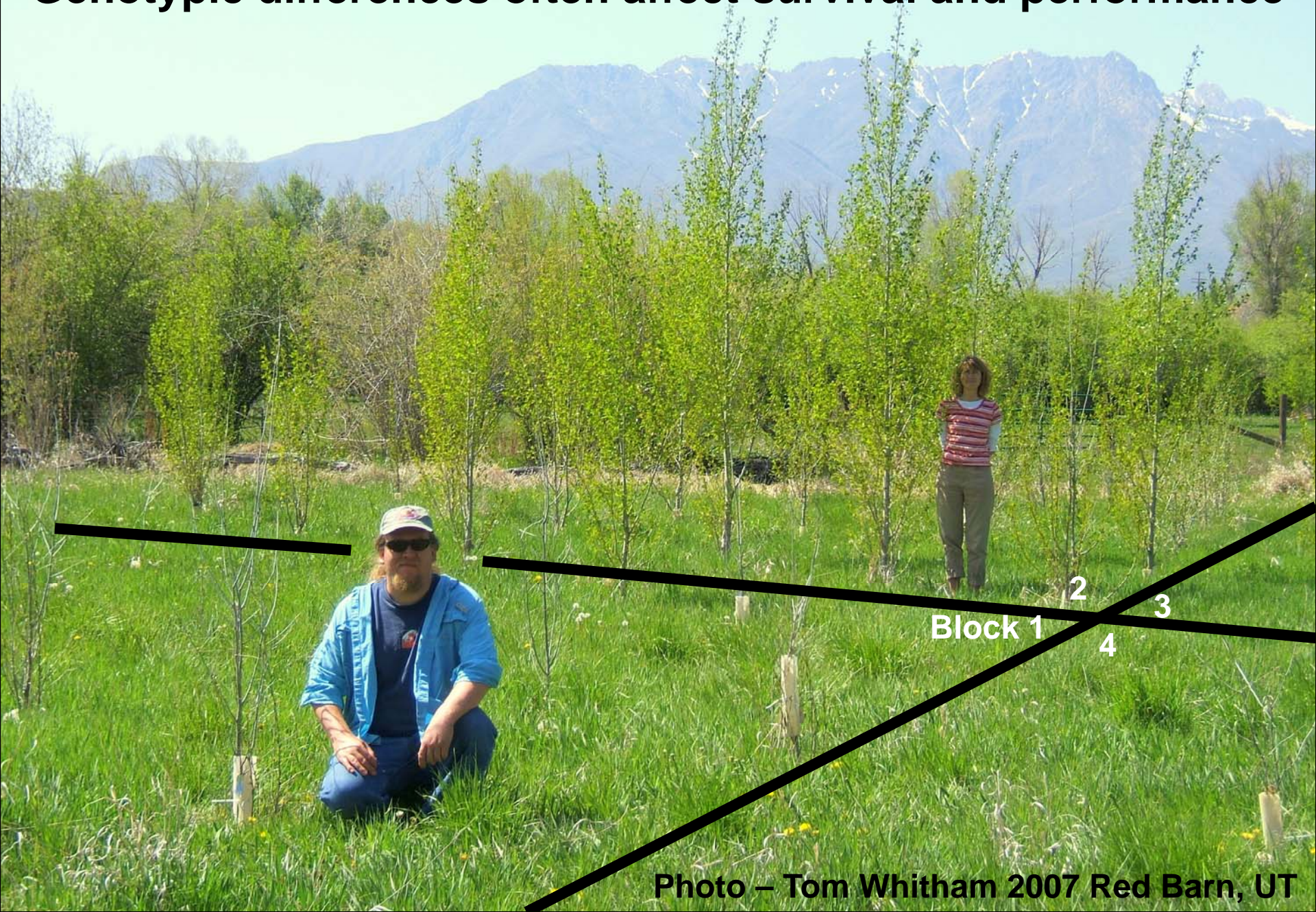
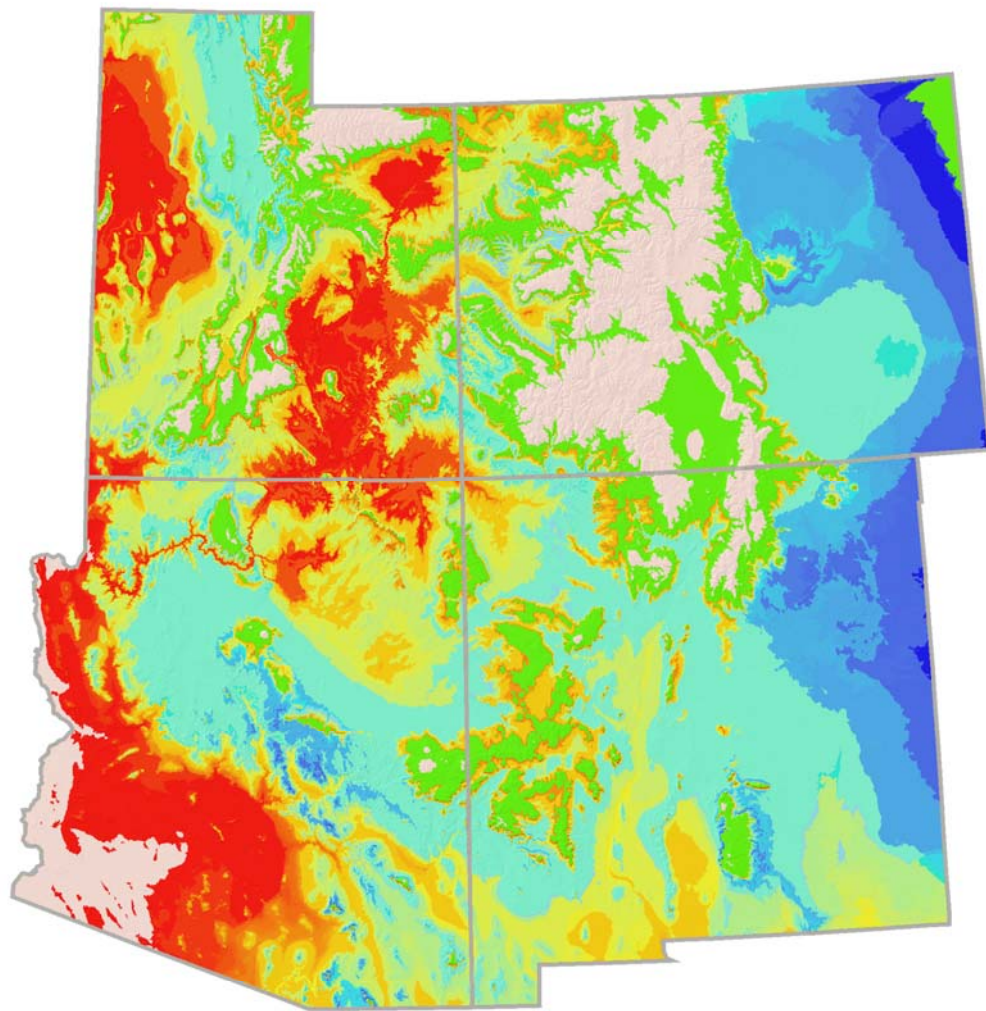


Photo – Tom Whitham 2007 Red Barn, UT

Cottonwood distributions based on GARP modeling of predicted climate change from IPCC



With predicted climate change locally adapted genotypes may no longer be locally adapted

**Gitlin & Whitham, unpub. data
Alicyn.Gitlin@nau.edu**

Plant Genetic Factors Account for ~50% of the Variation in Ecosystem Services

Plant Growth Rate
Constant 45%

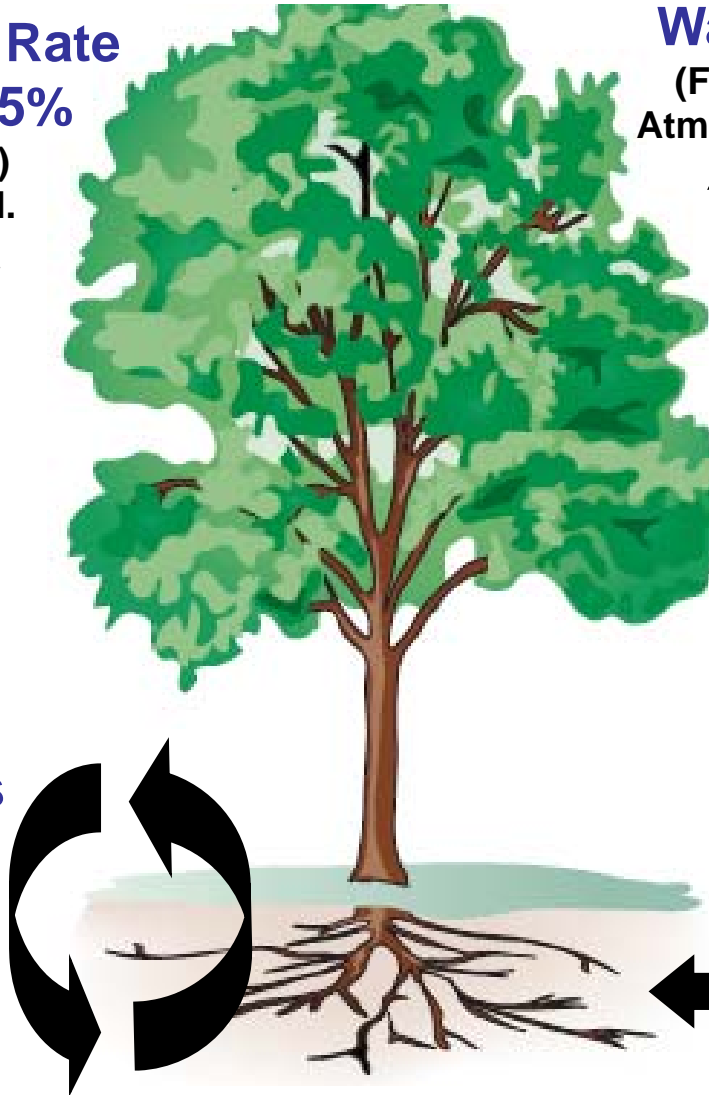
(Productivity)
Lojewski et al.
unpub. data

Community
Stability 25%

Keith et al.
unpub. data

Nutrient Cycles
34-65%

(Soil Fertility)
Schweitzer et al.
2004 Ecology Letters,
2005 Ecology, 2005
Oikos, LeRoy et al.
2006 Ecology



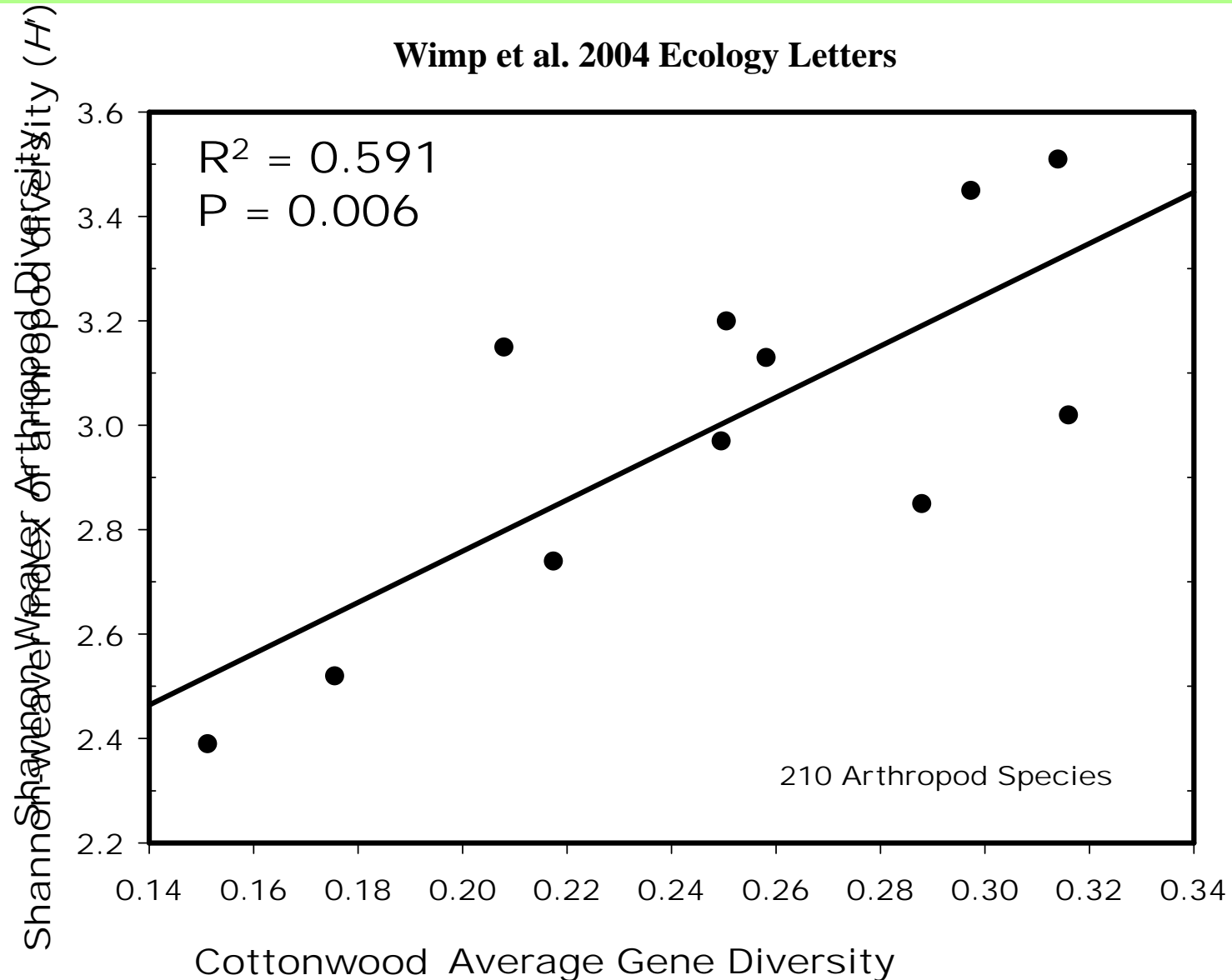
Water Cycles 35-40%
(Fluxes from Soil to Plant to
Atmosphere) Fischer et al. 2004
Oecologia

Biodiversity 43-78%
(Microorganisms, Herbivores,
Birds) Wimp et al. 2004 Ecology
Letters, Bangert et al. 2004
Conservation Biology, Shuster et
al. 2006 Evolution, Bailey et al.
Ecology Letters 2006, LeRoy et al.
2006 Ecology, Schweitzer et al.
2007 Ecology

Belowground Carbon
Storage & Root
Production 77%

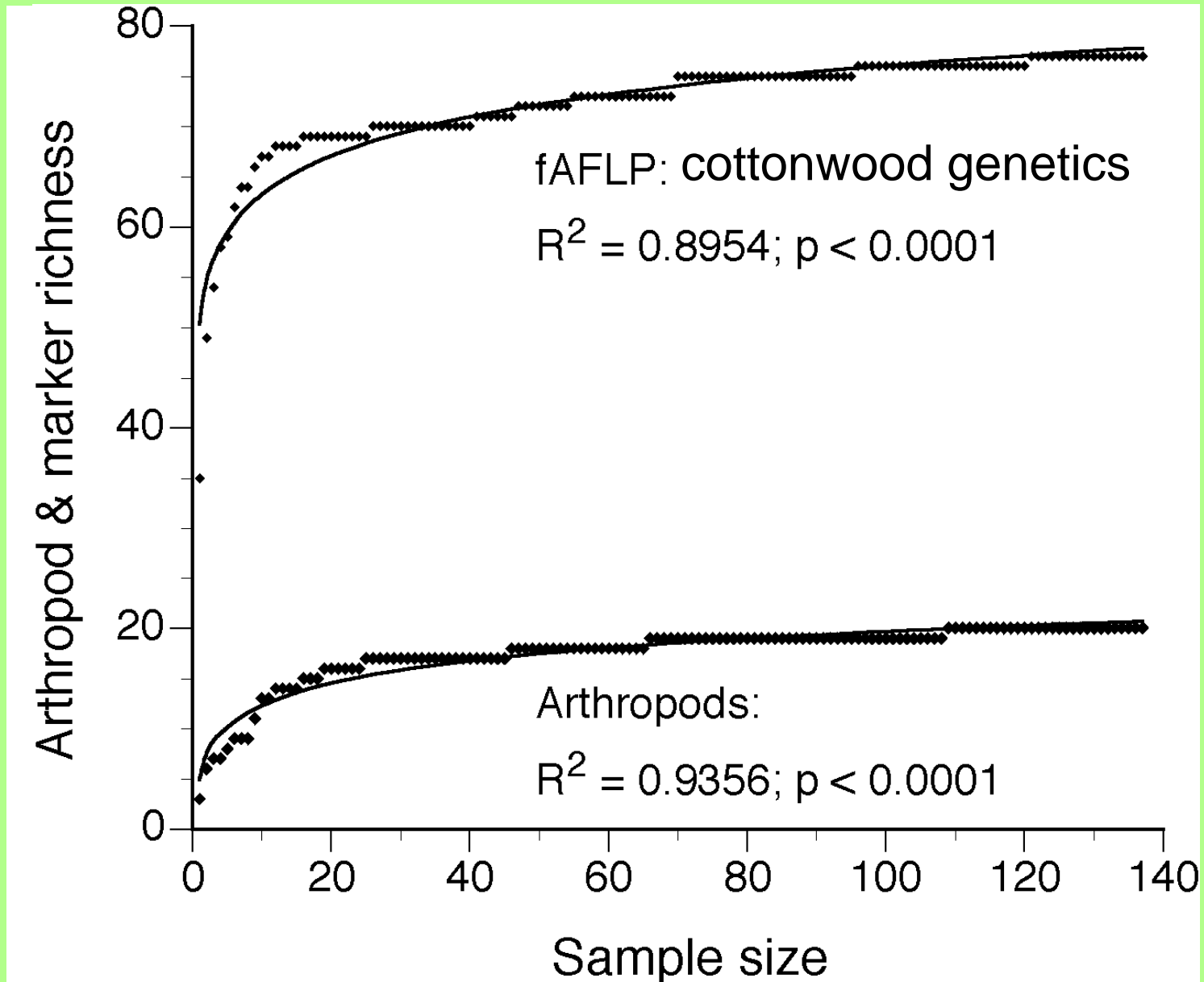
Fischer et al. 2006 Oecologia, Fischer
et al. 2007 New Phytologist

Plant genetic diversity determines arthropod diversity at a landscape level

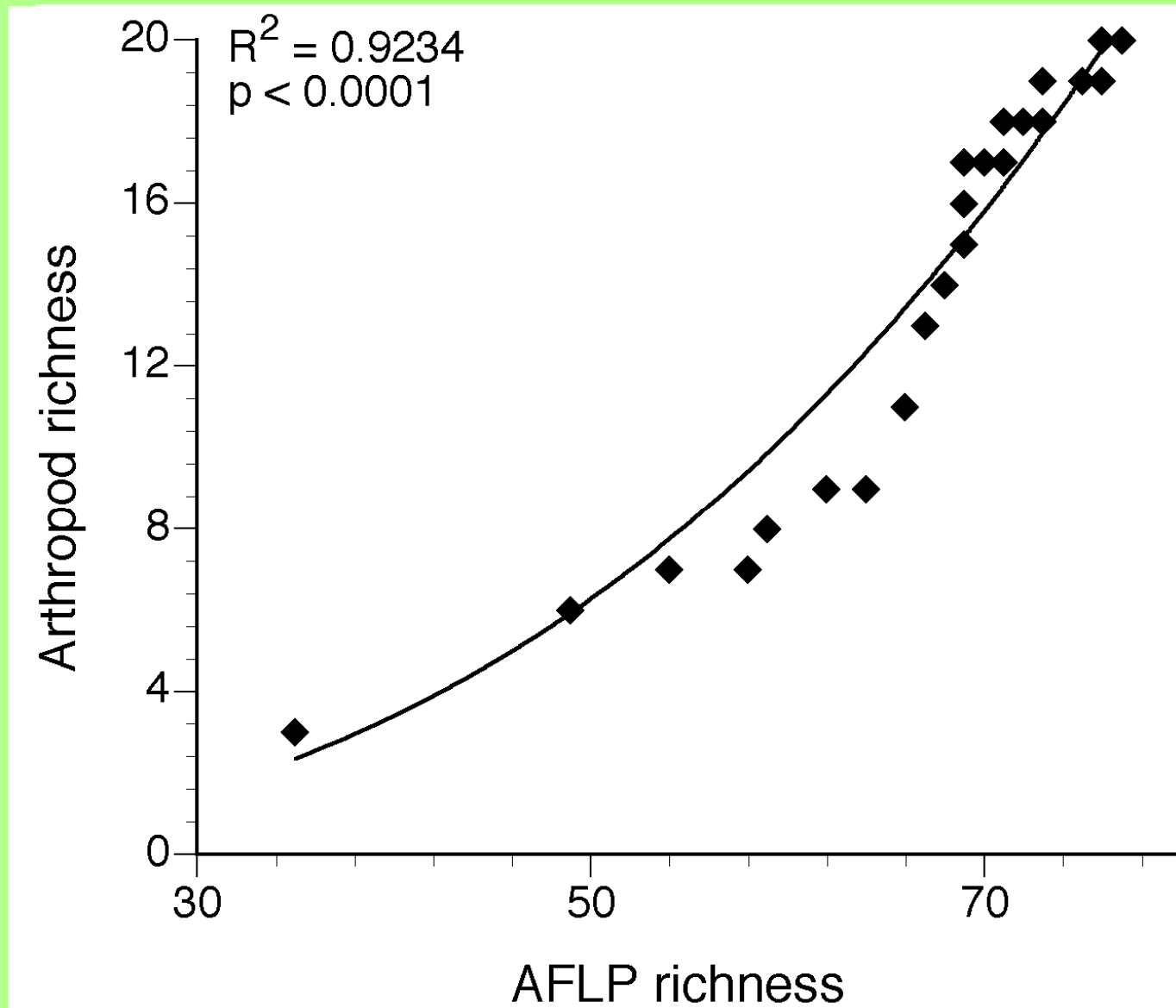


Species-area and genetic diversity-area curves are nearly identical

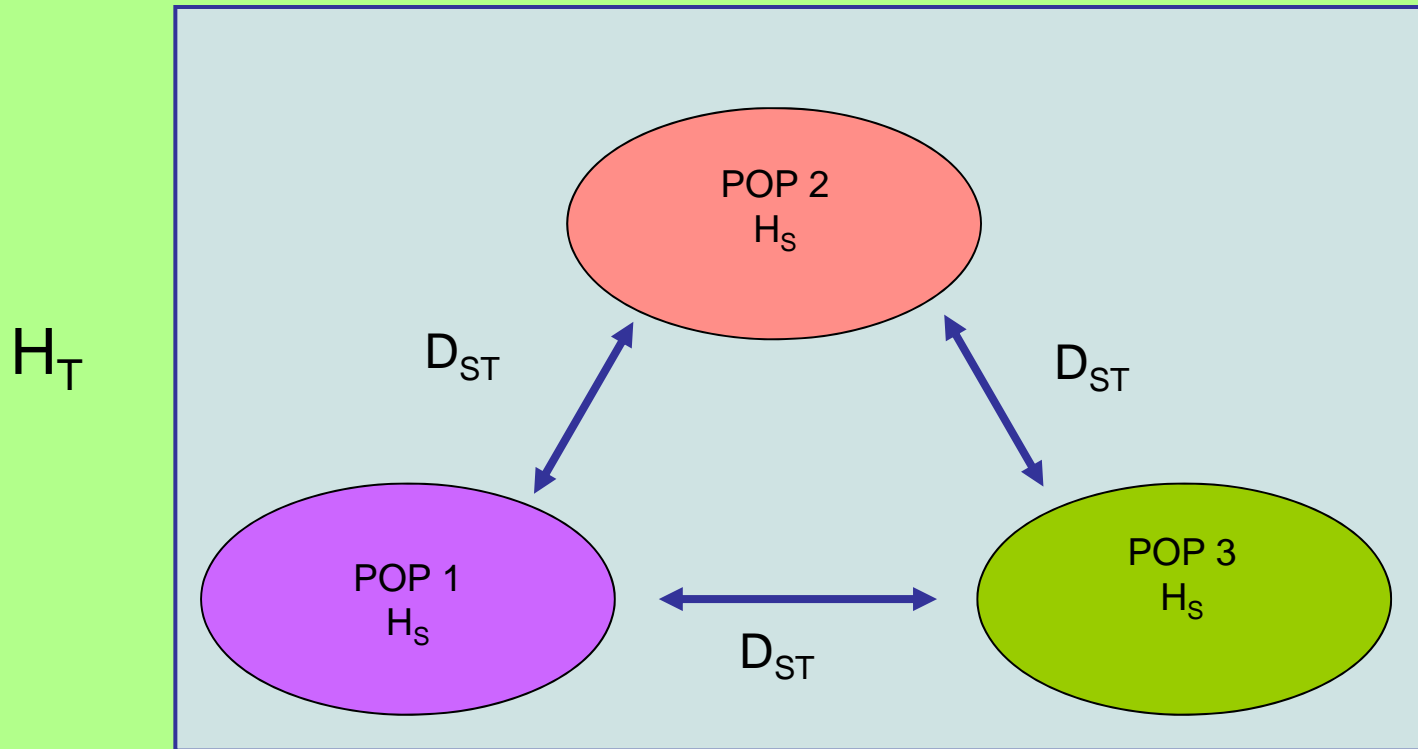
Is area important and/or is it genetic diversity?



Genetic variation in a foundation tree is highly correlated with species richness



Knowledge of population genetic diversity is critical for restoration and can be readily characterized



D_{ST} = Among population diversity

H_S = Within population diversity

H_T = Total population diversity

Restoration and Landscape Level Experiments



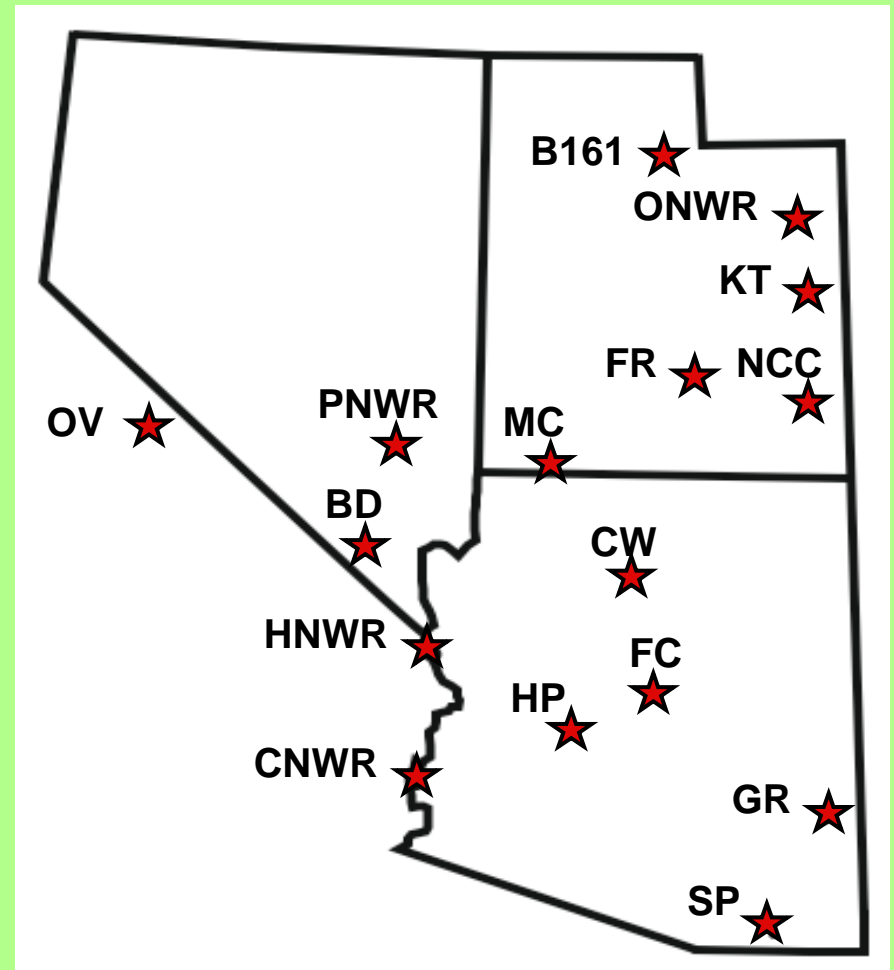
- Key Partnerships
- Two restoration projects on the Lower Colorado River (LCR)
 - Cibola National Wildlife Refuge (CNWR)
 - Palo Verde Ecological Reserve (PVER)
- The goal of these projects is to combine riparian habitat restoration with scientific investigation

Photo credit:
BOR



Collections: Cibola National Wildlife Refuge

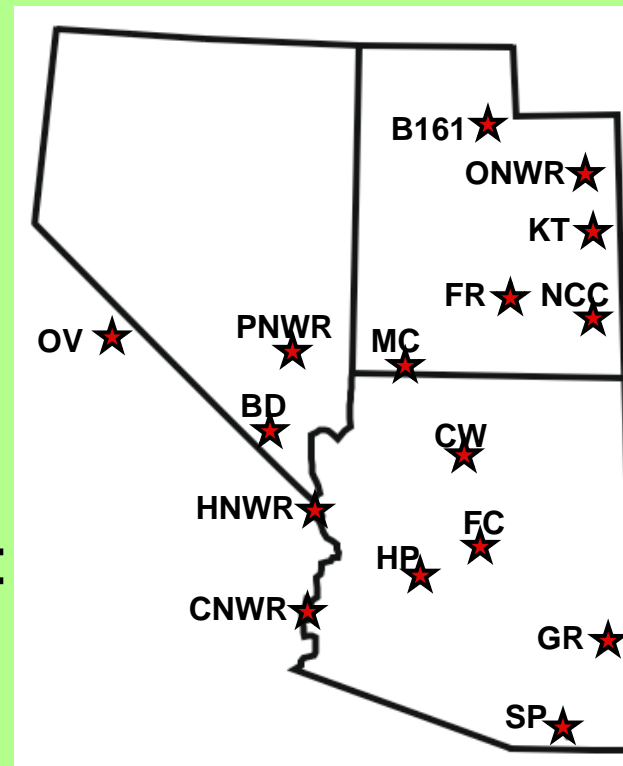
- 16 different Fremont genotypes were collected from NV, AZ, UT, and CA
- Trees propagated at NAU greenhouses
- Experimental Question: What effect does tree genetic diversity have on community diversity?





Collections: Cibola National Wildlife Refuge

- 16 different Fremont genotypes were collected from NV, AZ, UT, and CA
- Trees propagated at NAU greenhouses
- Experimental Question: What effect does tree genetic diversity have on community diversity?



Planting at Cibola NWR

6,400 Fremont Cottonwoods were planted on 4m centers in March 2007 in an ag. land conversion at Cibola NWR where they are flood irrigated.

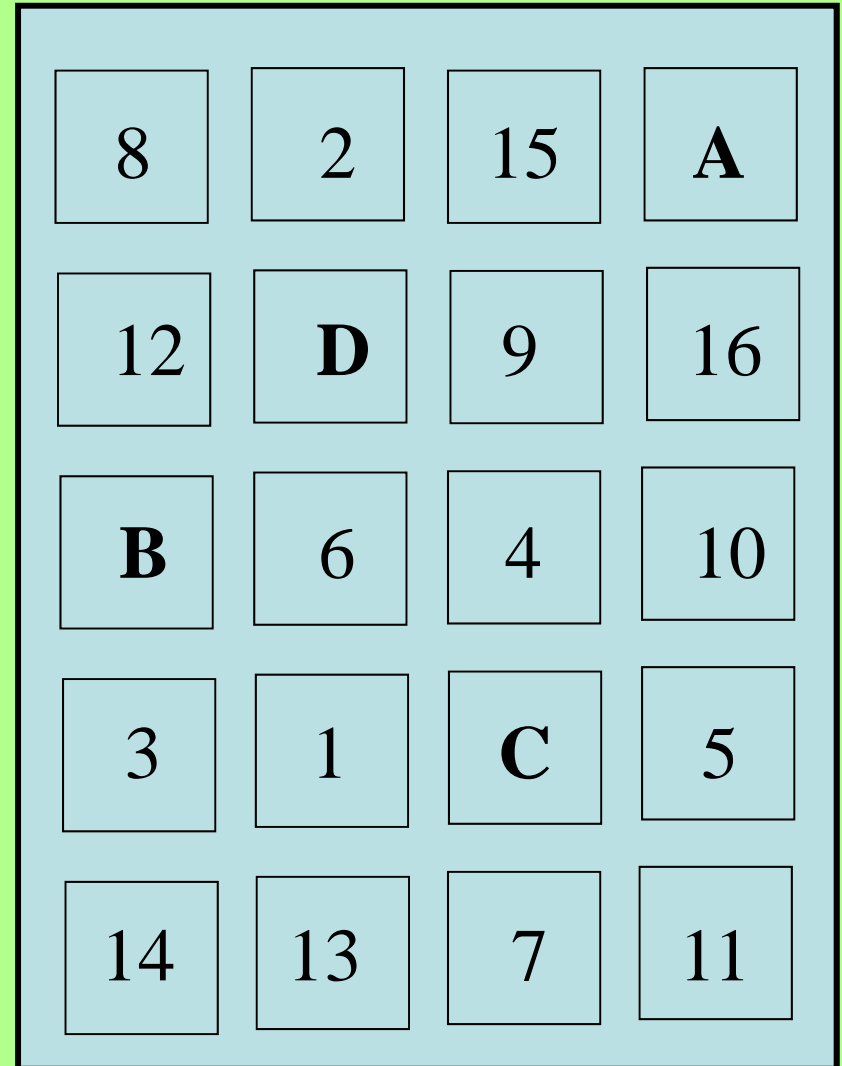


4 months later the trees are averaging 2 meters in height



Experimental Design : CNWR

- 16 different genotypes in pure and mixed stands of 16 trees each
 - Numbers represent different genotypes (e.g., genotype #8)
 - Letters represent different mixes of genotypes
 - **A** mix: 16 genotypes (1 tree/ea)
 - **B** mix: 8 genotypes (2 trees/ea)
 - **C** mix: 4 genotypes (4 trees/ea)
 - **D** mix: 2 genotypes (8 trees/ea)



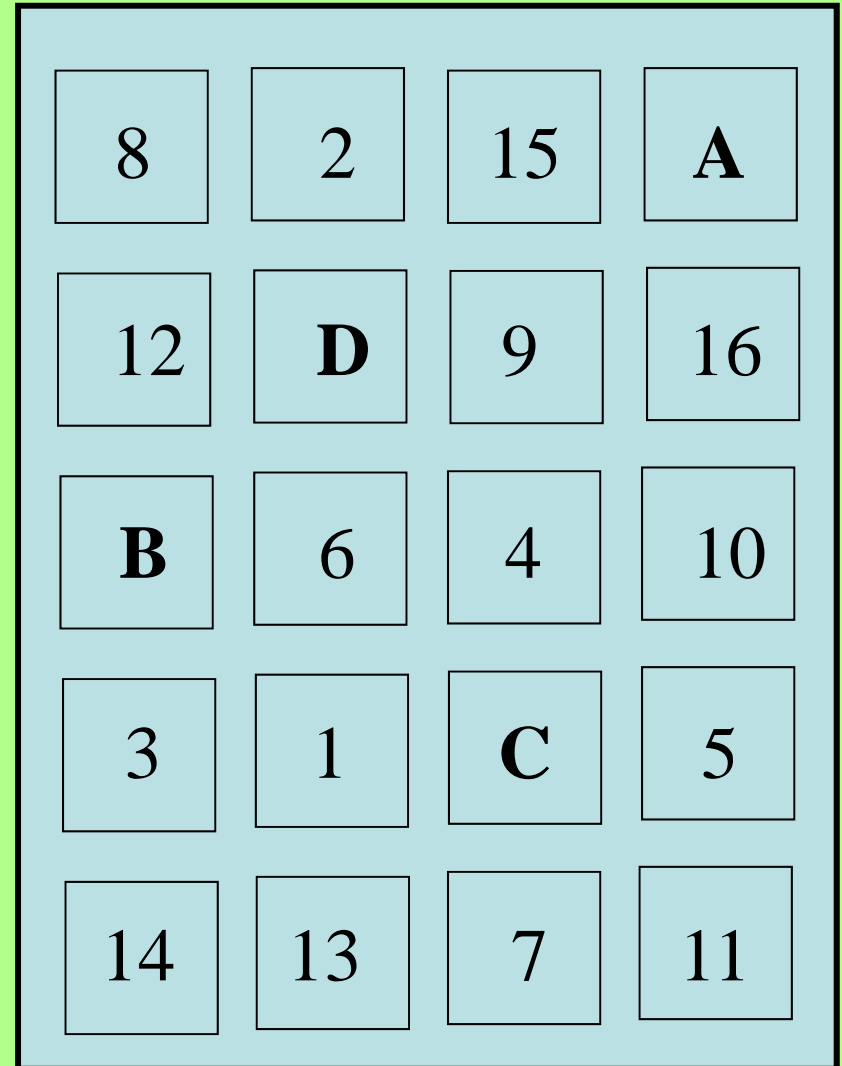
1 Block = 20 stands x 20 reps = 6400 trees

Experimental Design : CNWR

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Experimental Question:

What effect does tree genetic diversity have on community diversity?



1 Block = 20 stands x 20 reps = 6400 trees

Research at Cibola NWR

7 years old



5 years old



2 years old



**Do different aged stands support
different arthropod communities
and increased biodiversity?**

Laura Hagenauer

Ph.D. Student

Northern Arizona University



Research at Cibola NWR

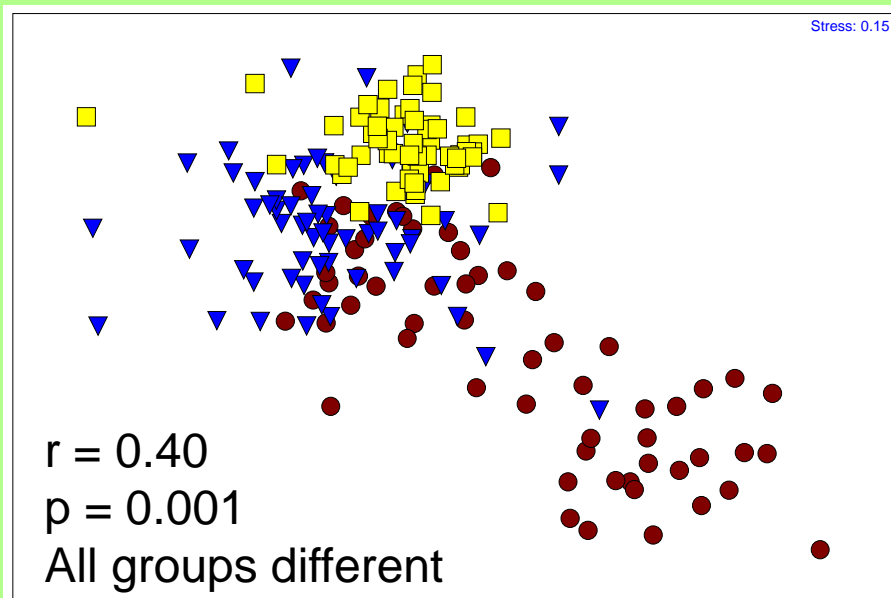
7 years old ▼



5 years old ●



2 years old ■



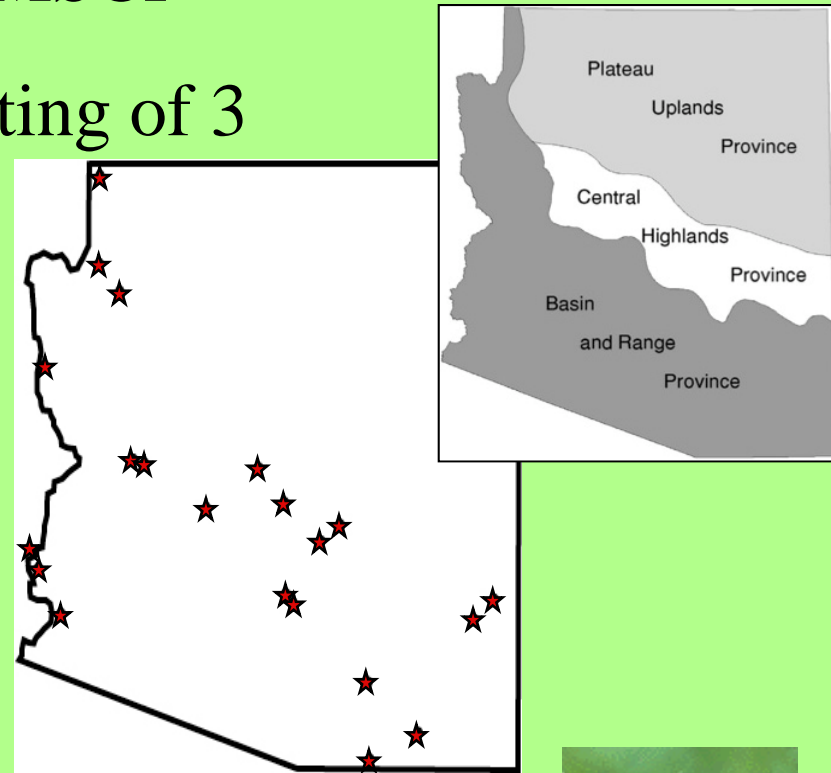
Different aged stands DO support different arthropod communities

Laura Hagenauer
Ph.D. Student
Northern Arizona University



Palo Verde Ecological Reserve

- Restoration of approximately 5,940 acres of riparian habitat along the LCR: MSCP
- Collection, propagation and planting of 3 riparian species:
 - Fremont cottonwood, Goodding's willow, and Coyote willow
 - 20 collection locations
 - 10 genotypes/ species/ location
 - 200 genotypes/ species
 - 600 total genotypes
- Examine how differences in vegetation density and genetic diversity affect recovery of the southwestern willow flycatcher



Planting at PVER

16,896 individuals were planted on 2m centers in March 2007 in an ag. land conversion at PVER where they are flood irrigated.

March 2007



May 2007 - 2 months



August 2007 - 5 months



March 2007



Experimental Design: PVER

4	1	3	1	2	6
6	2	6	3	3	6
2	4	6	5	6	5
5	3	4	5	5	2
3	1	1	5	4	6
2	4	5	6	6	5
2	4	5	6	5	6
2	5	5	3	3	3
1	6	5	6	6	6
2	4	3	1	1	4
6	1	5	5	6	5

Different genotypes planted in different proportions

Treatment	Density Treatment Pf / Sg	Raw Numbers Pf / Sg / Se	Proportion Pf / Sg / Se
1	L / L	22/22/212	.087/.087/.83
2	H / L	43/22/189	.17/.087/.74
3	L / H	22/43/189	.087/.17/.74
4	H / H	43/43/169	.17/.17/.66
5	M / M	33/33/189	.13/.13/.74
6	M / M	33/33/189	.13/.13/.74

20 acre ag land
conversion

66 blocks

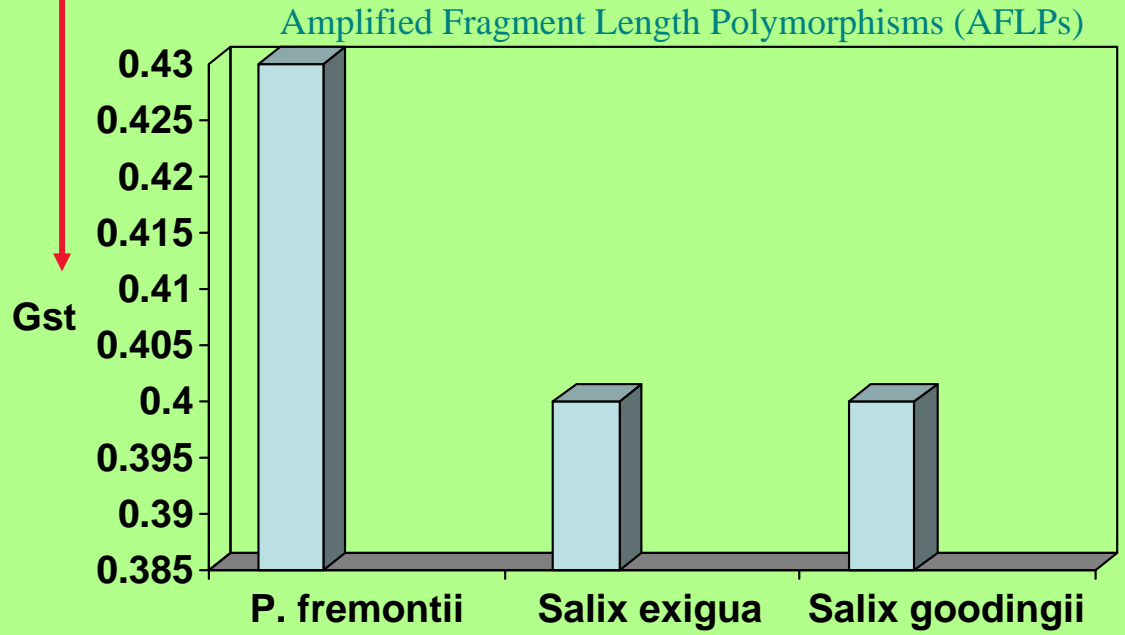
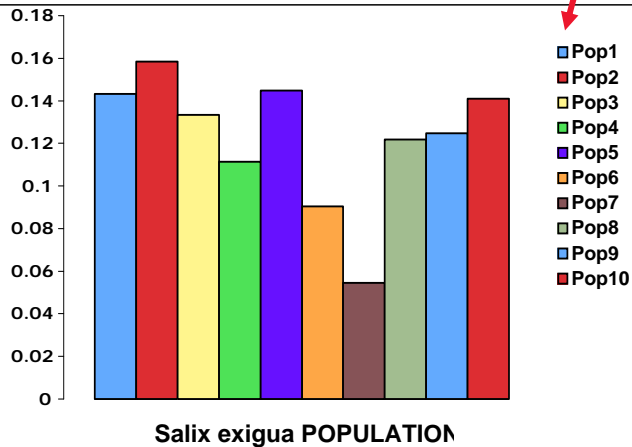
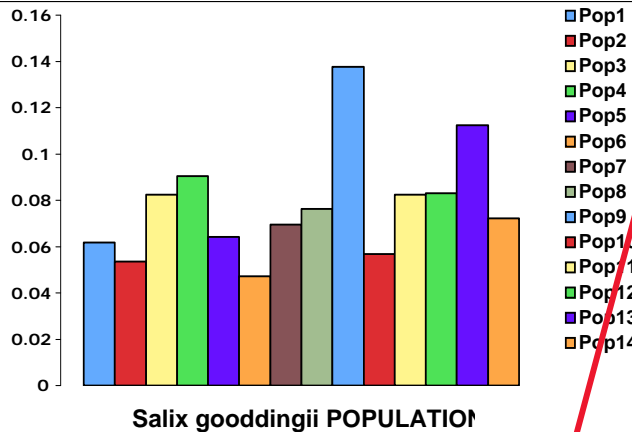
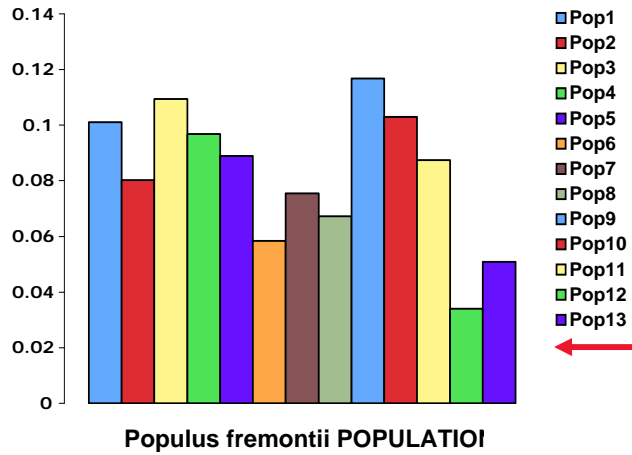
Pf = *Populus fremontii* **Sg** = *Salix gooddingii*

Se = *Salix exigua*

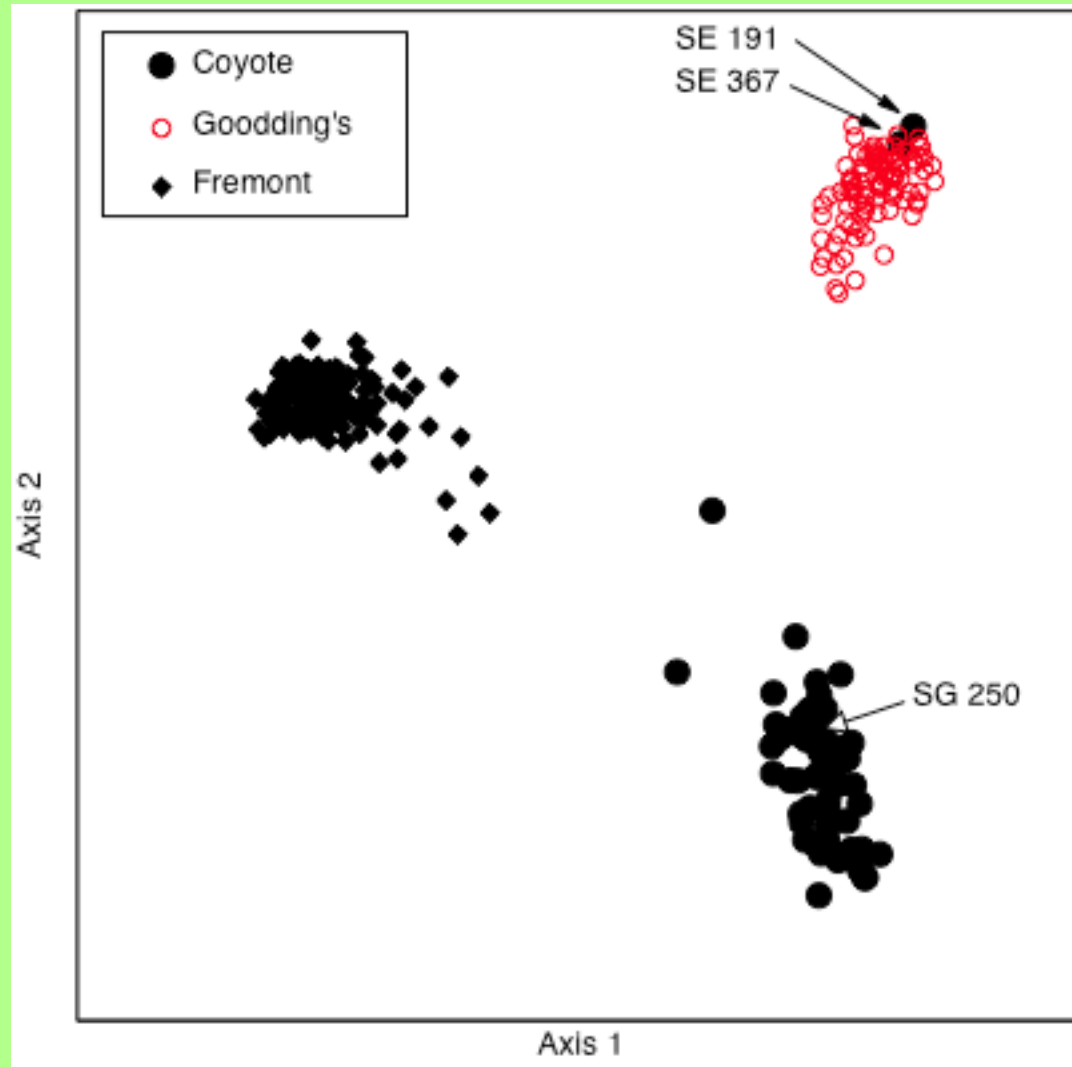
Using genetic diversity as a tool for restoration

H_s : Within population diversity

G_{ST} : Among population diversity



Genetic differentiation of 3 riparian species used in the PVER garden



Mis-labeled samples are easily identified using the AFLP method

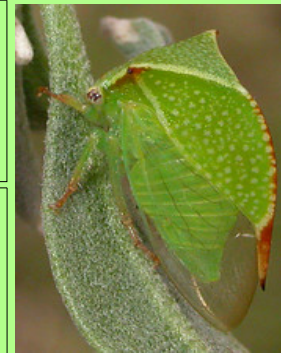
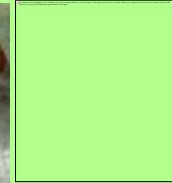
Arthropod Surveys at PVER



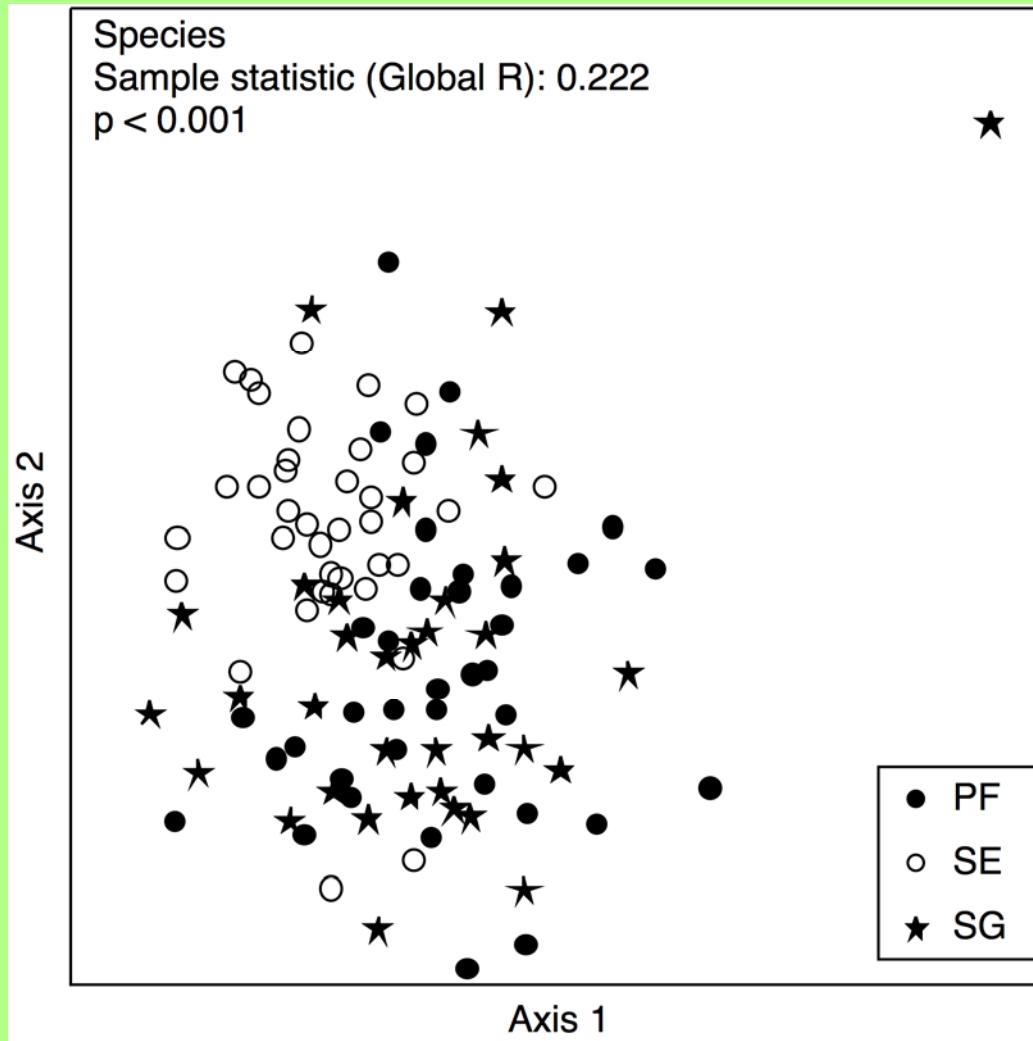
Timed Visual Census Sampling: Sampling the specific arthropods associated with individual genotypes of each of the three plant species in all density treatments.



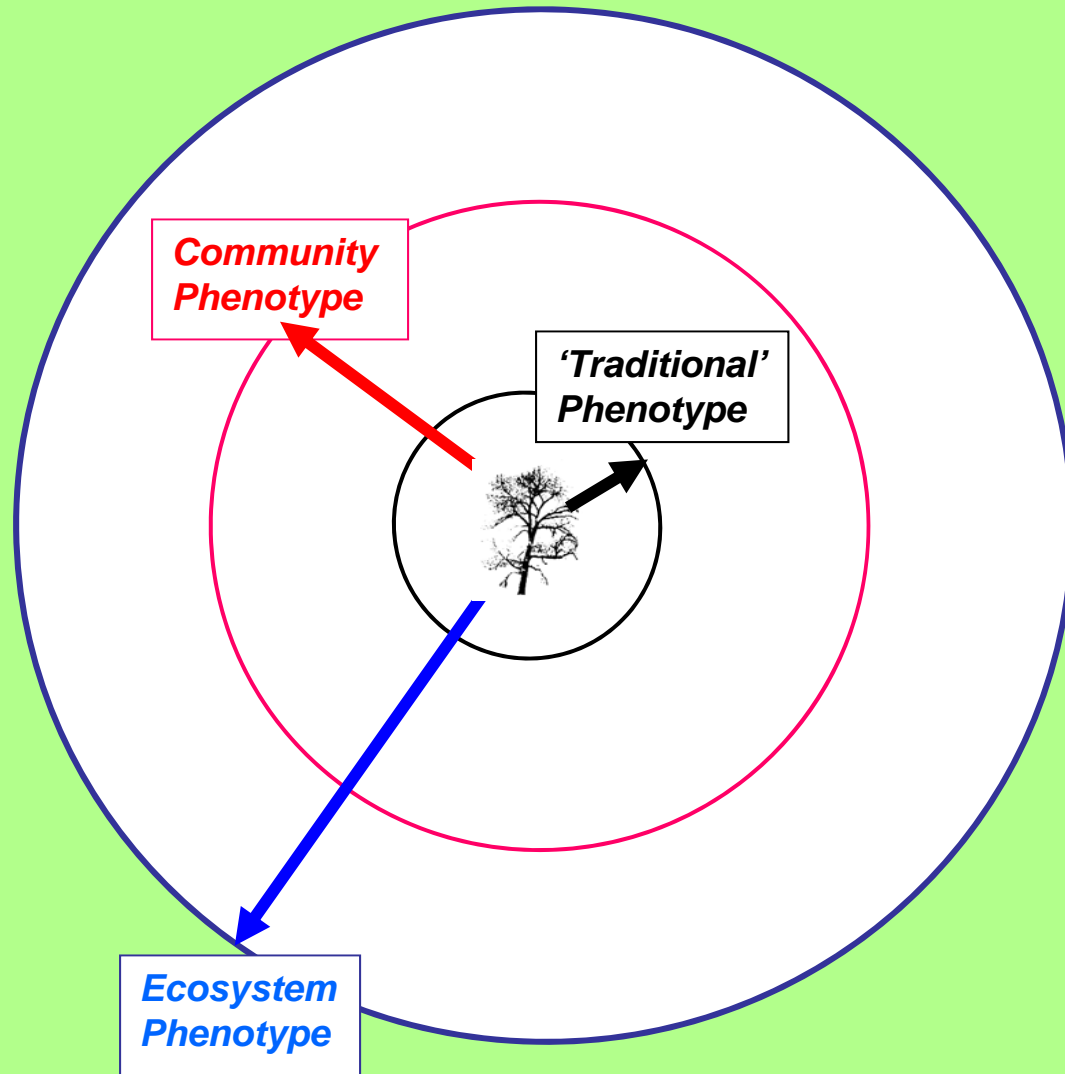
Malaise Sampling: Sampling the general arthropod prey base available in each of the density treatments



Arthropod communities are statistically different among plant species



Ecological Genetic Considerations for Riparian Habitat Restoration



1. Host **plant genetics** directly **influences** the structure and diversification of **ecological communities**.

2. **Foundation tree species** can have a profound **impact** on **dependent communities and ecosystem processes** (Whitham et al. 2006 Nature Reviews Genetics).

3. **Genetic variation** and its extended effects should be a **major consideration** when considering management strategies for **habitat restoration**

Science meets Restoration



NAU ECOLOGY GROUP & COLLABORATORS

Gery Allan – molecular systematics

Brad Blake – greenhouse manager

Sam Chapman – nutrient cycling

Zacchaeus Compson – aquatic ecology

ecology

Sharon Ferrier – conservation ecology

Robert Footitt – molecular systematics

Laura Hagenauer - biodiversity

Paul Heinrich – public outreach

Paul Keim – microbial genetics

George Koch – physiological ecology

Carri LeRoy – aquatic ecology

Eric Lonsdorf – genetic modeling

Brad Potts – quantitative genetics

Paul Selmants - soil ecology

Adrian Stone – community ecology

Talbot Trotter – dendrochronology

Gina Wimp – community ecology

Scott Woolbright - molecular genetics

Hao Ma – molecular genetics

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Helen Bothwell – phylogeography

Aimee Classen – ecosystem dynamic

Steve DiFazio – molecular ecology

Dylan Fischer – ecophysiology

Catherine Gehring – microbial ecology

Steve Hart – ecosystem/soil ecology

Lisa Holeski – genetics & chemistry

Art Keith – insect community ecology

Zsuzsi Kovacs – mycorrhizal ecology

Rick Lindroth – chemical ecology

Jane Marks – aquatic ecology

Brian Rehill – chemical ecology

David Smith – spatial genetics

Chris Sthultz – plant ecology

Amy Whipple – ecological genetics

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Randy Bangert – biogeography

Bill Bridgeland – avian ecology

Neil Cobb – insect ecology

Luke Evans – population

Kevin Floate – insect ecology

Alicyn Gitlin – climate modeling

Kris Haskins – mycorrhizal ecology

Barbara Honchak – ecological genetics

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Nathan Lojewski – productivity

Nashelly Meneses – ecological genetics

Jen Schweitzer – ecosystems

Steve Shuster – theoretical genetics

Richard Turek – statistics

Tom Whitham – ecology

Stuart Wooley – phytochemistry

Gancho Slavov – population genetics

Gail Iglitz - Bureau of Reclamation

Conrad Jones - CA Fish and Game

