

Photos by Hillary Cooper (top) & Tom Whitham (bottom)

Cottonwood mortality on Bill Williams River

National Wildlife Refuge – March 28, 2017



## Genetics- Based Restoration in Response to Climate Change and Invasive Species

**Tom Whitham, Hillary Cooper, Heather Gillette, Adrian Stone, Sean Mahoney, Jackie Parker, Catherine Gehring, Art Keith, Alicyn Gitlin, Lisa Markovchick, Julia Hull, Gery Allan, Kevin Grady, and Kevin Hultine**  
**Merriam-Powell Center for Environmental Research & Others**

NORTHERN  
ARIZONA  
UNIVERSITY

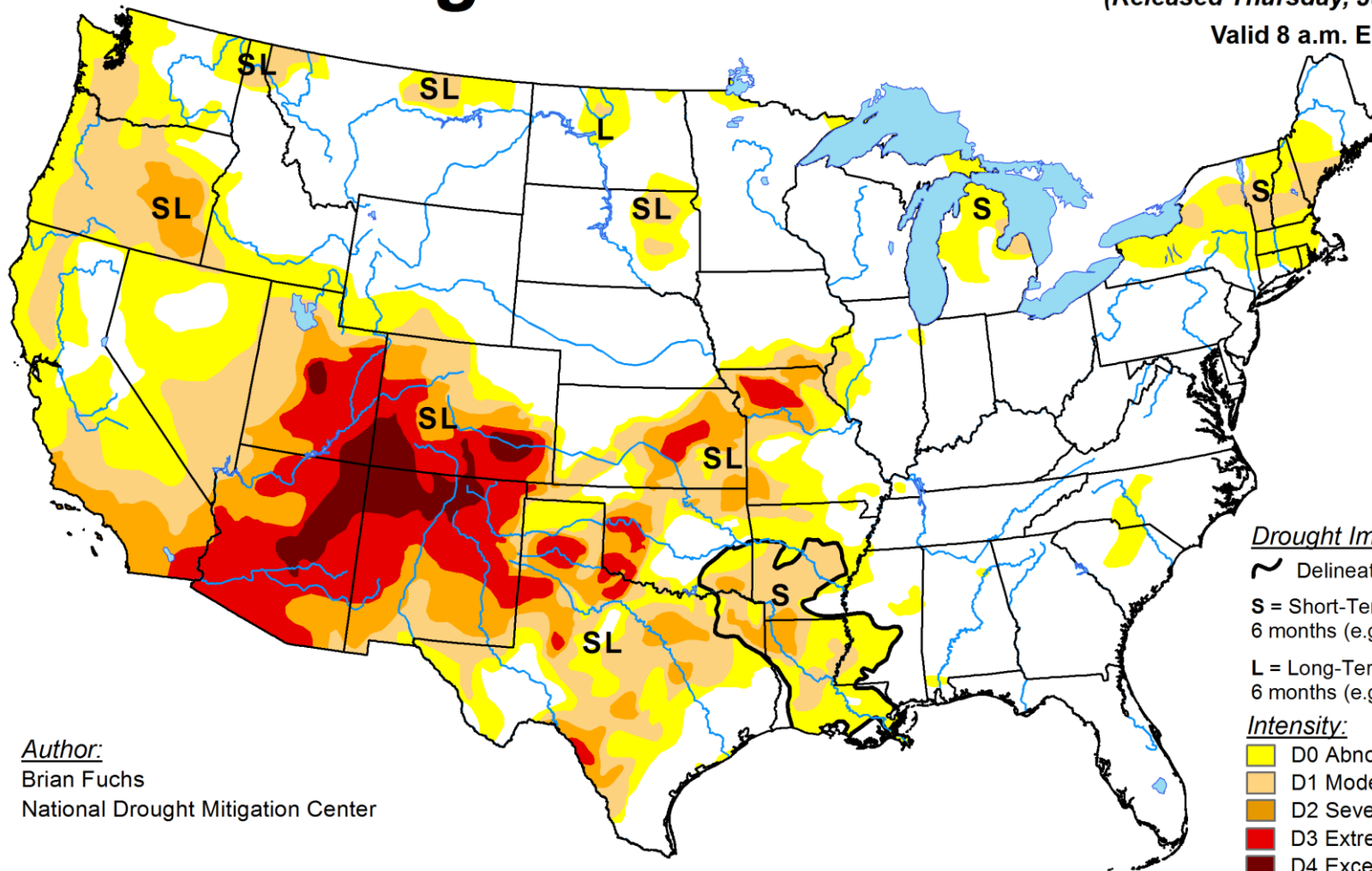


# U.S. Drought Monitor

July 10, 2018

(Released Thursday, Jul. 12, 2018)

Valid 8 a.m. EDT



## Author:

Brian Fuchs

National Drought Mitigation Center

## Drought Impact Types:

~ Delineates dominant impacts

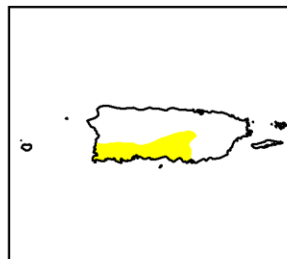
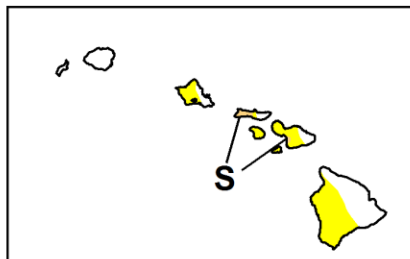
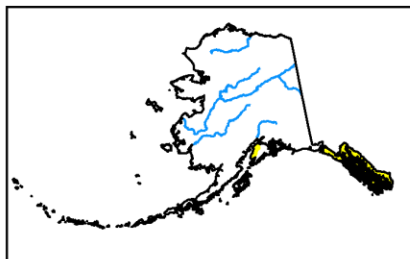
S = Short-Term, typically less than 6 months (e.g. agriculture, grasslands)

L = Long-Term, typically greater than 6 months (e.g. hydrology, ecology)

## Intensity:

- D0 Abnormally Dry
- D1 Moderate Drought
- D2 Severe Drought
- D3 Extreme Drought
- D4 Exceptional Drought

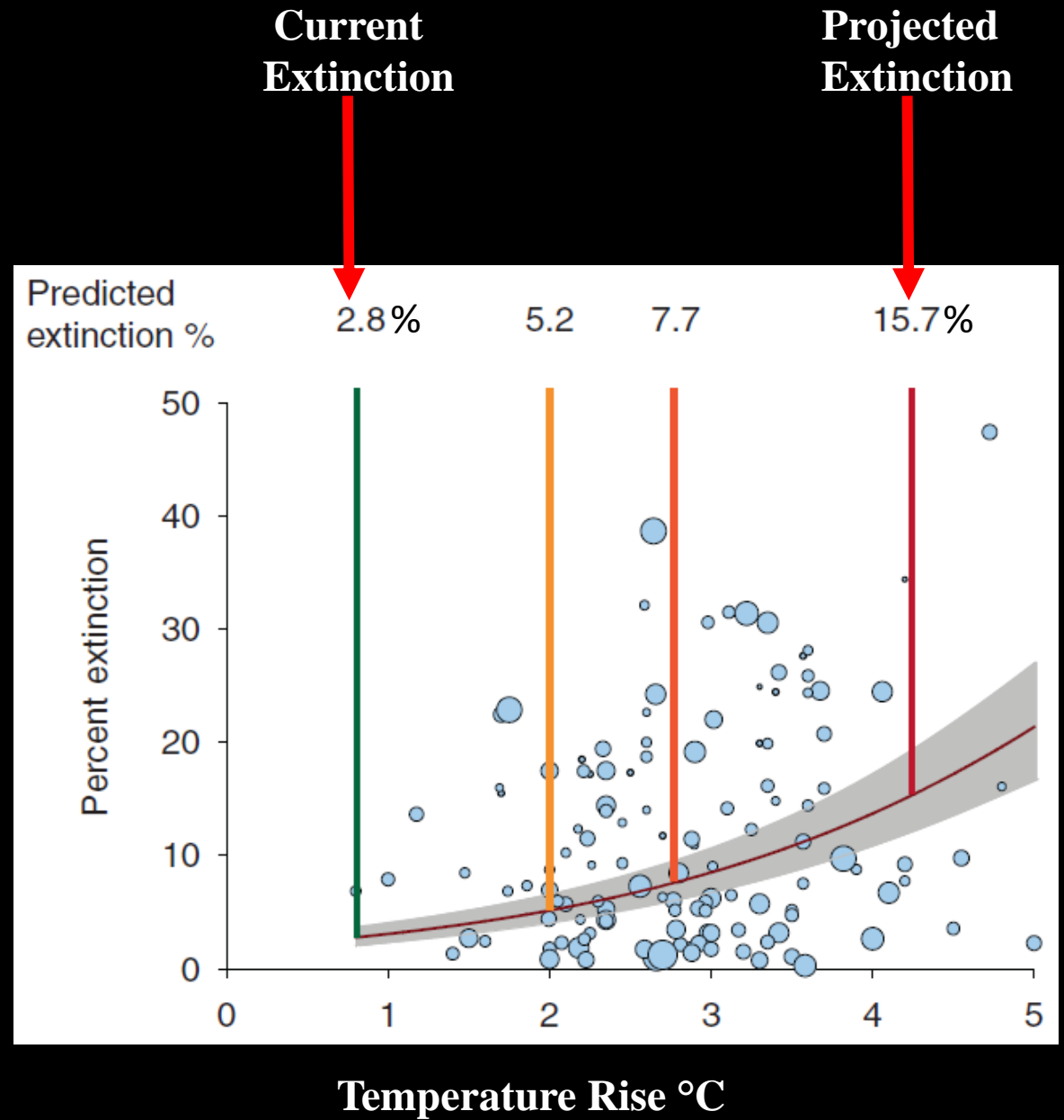
The Drought Monitor focuses on broad-scale conditions. Local conditions may vary. See accompanying text summary for forecast statements.

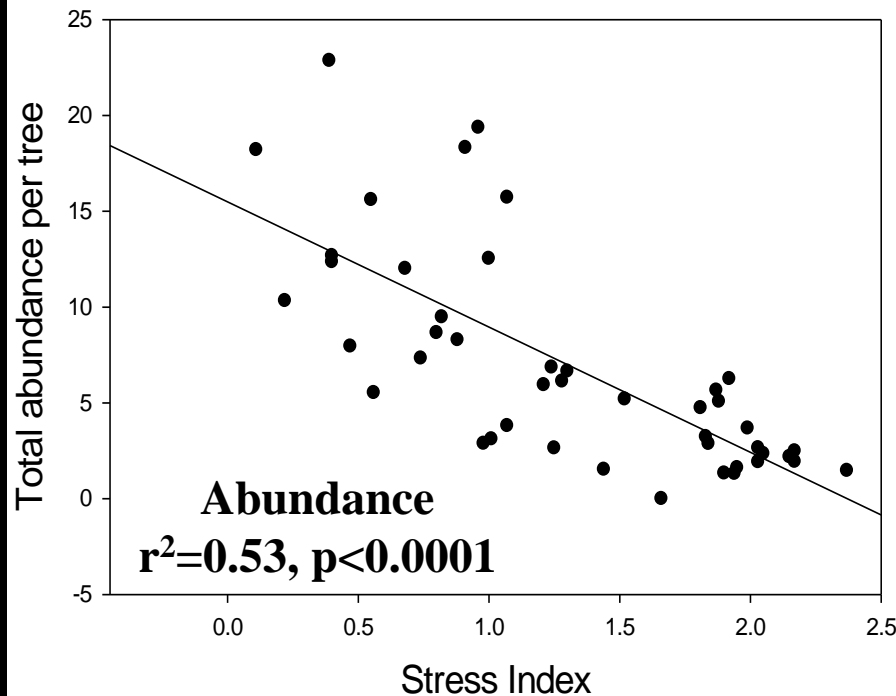
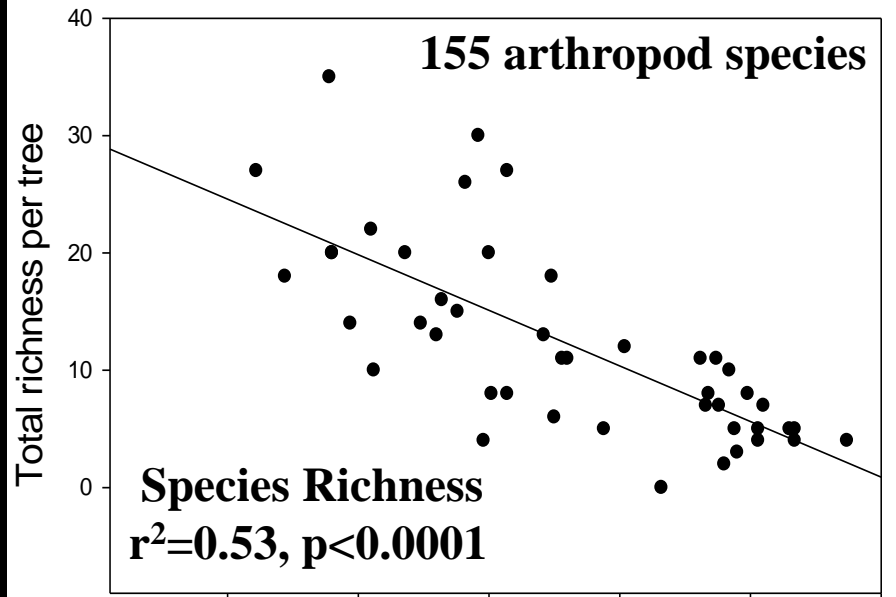


<http://droughtmonitor.unl.edu/>

# Increased extinction with global warming

Every degree  
increase in temperature  
increases extinction  
risk so planners  
need to build this into  
their designs. With  
current practices 1 in 6  
species will go extinct.



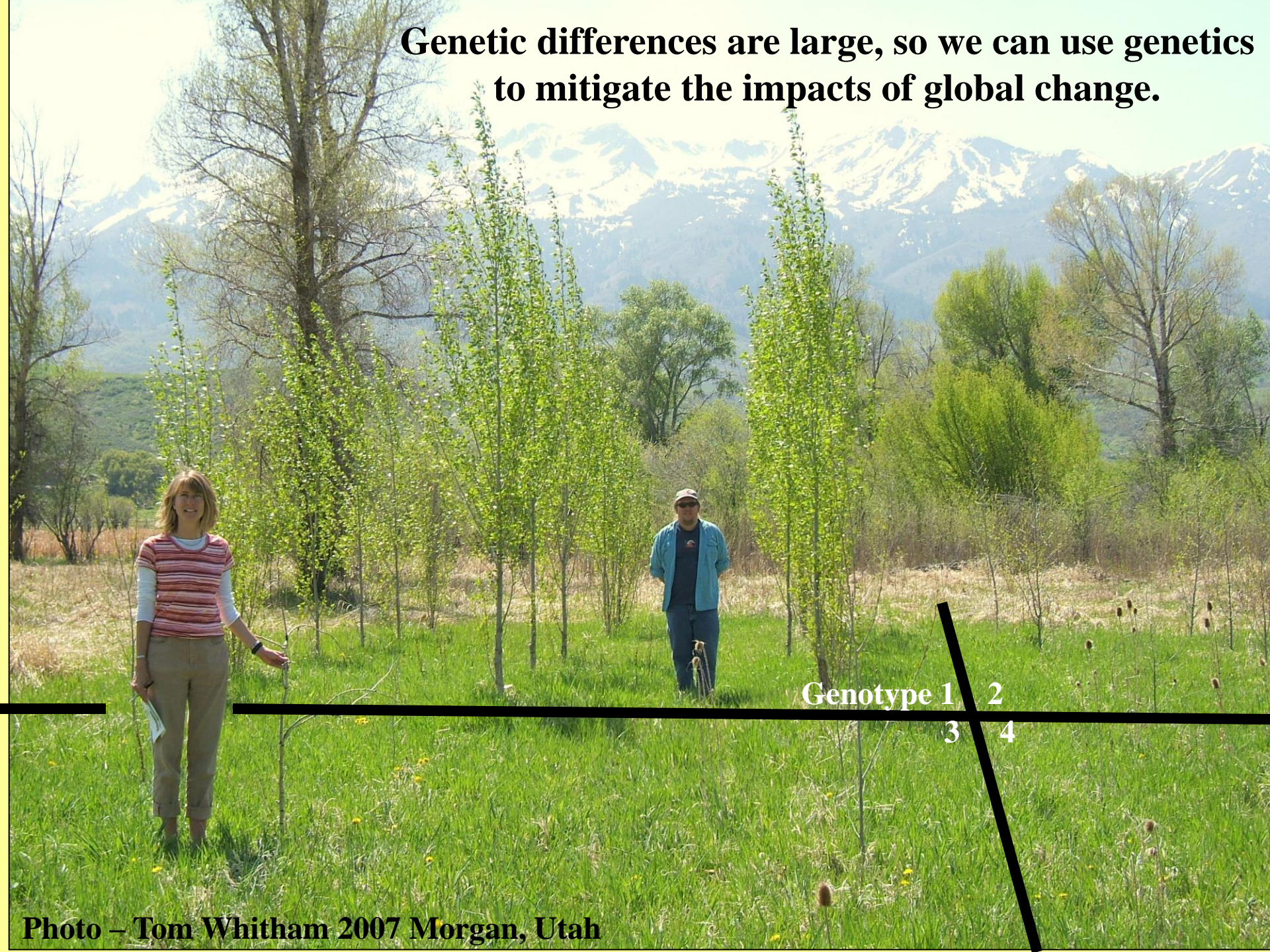


**Within a site, increased stress negatively affects arthropod diversity.**

**Stress Index = (standardized branch dieback + number of needle cohorts + radial trunk growth)**

**Stone et al. 2010 Oecologia**

**Genetic differences are large, so we can use genetics to mitigate the impacts of global change.**



**Photo – Tom Whitham 2007 Morgan, Utah**

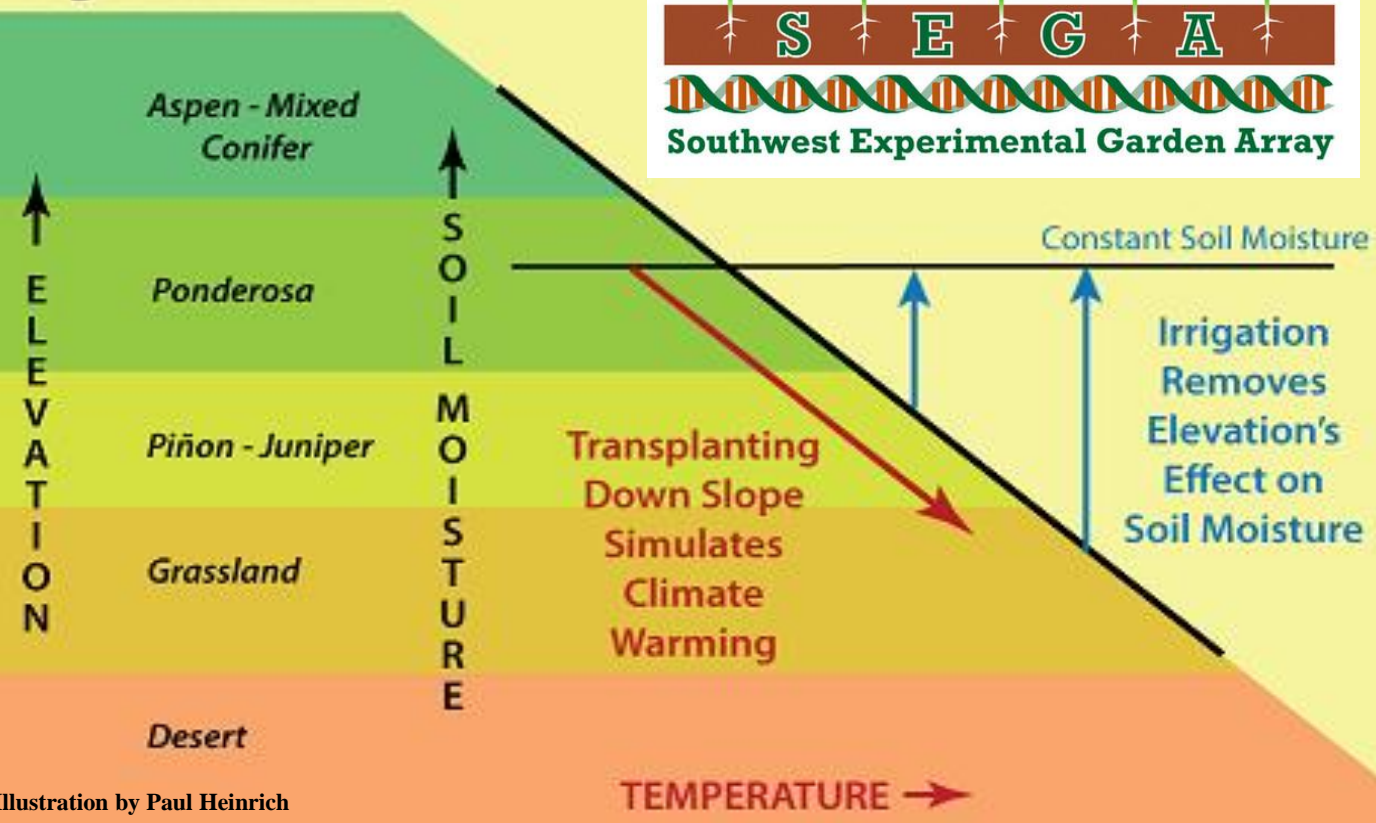
# SEGA site at The Arboretum at Flagstaff – A network of 10 common gardens along an elevation gradient to develop solutions to global change



THE ARBORETUM  
AT FLAGSTAFF

**The more locally adapted, the greater the impacts of climate change**

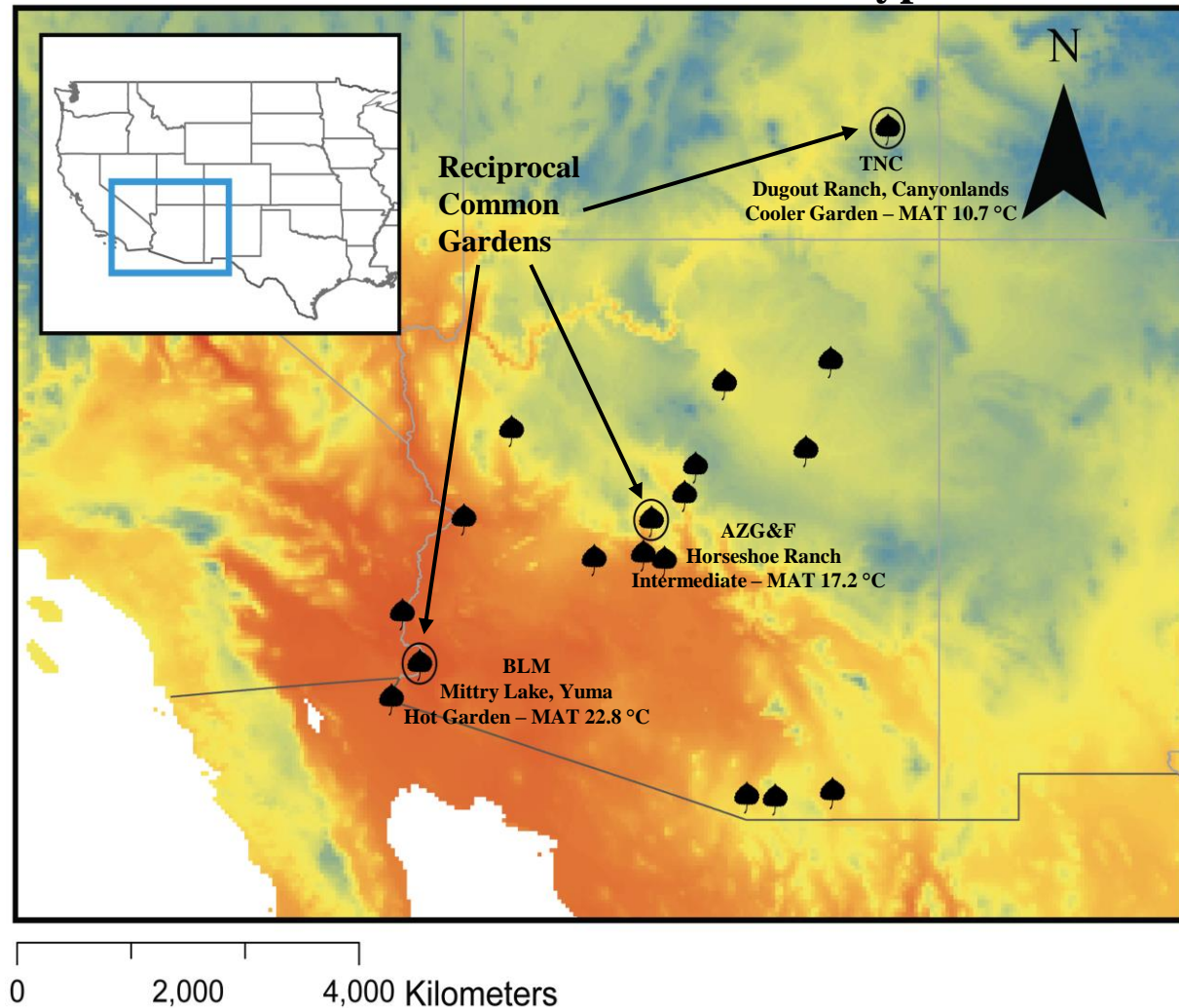
## SEGA Vegetation Zones



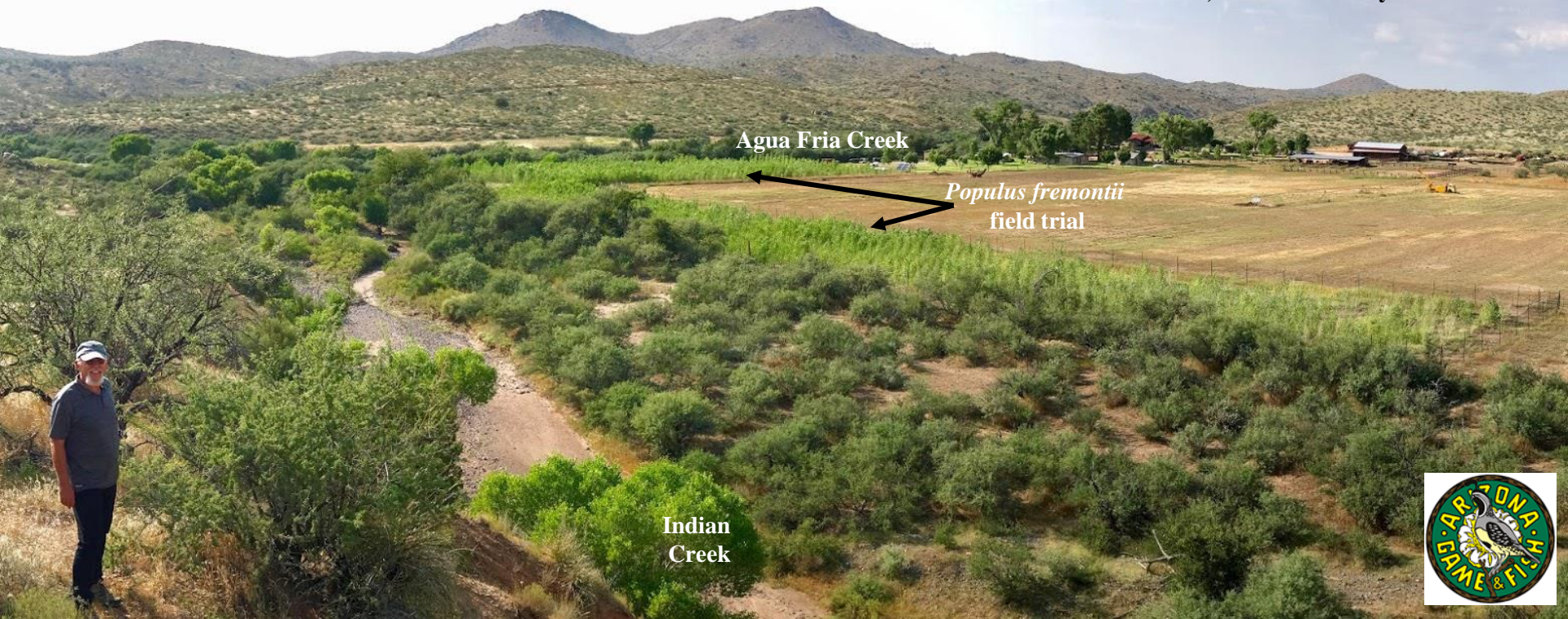
([sega.nau.edu](http://sega.nau.edu))  
\$4.5 million NSF/NAU,  
GO, and NGO  
Participants: USFS,  
NPS, BLM, BOR, TNC,  
AZ Game & Fish, Babbitt  
Ranches, Grand Canyon  
Trust, & The Arboretum  
at Flagstaff

If plants must move to survive future climate conditions, how do we decide on which ecotypes and genotypes to use in restoration projects? SEGA network provides next generation genetics-based infrastructure to scientifically make such decisions.

## Reciprocal common gardens show fine scale local adaptation within the Sonoran desert ecotype



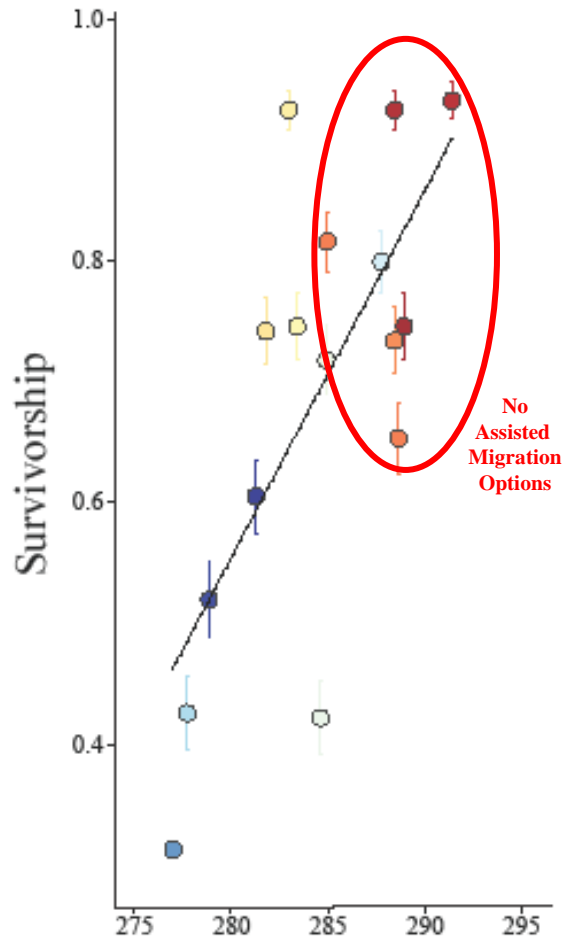
Location map of 16 provenance collection sites (leaf icon) of *Populus fremontii* and the three common garden locations (leaf with circle). The central garden is also a collection site. The shading corresponds to the degree-days above 5°C (DD5) throughout the region: red represents high DD5, blue low DD5. Cooper et al. 2019 Global Change Biology



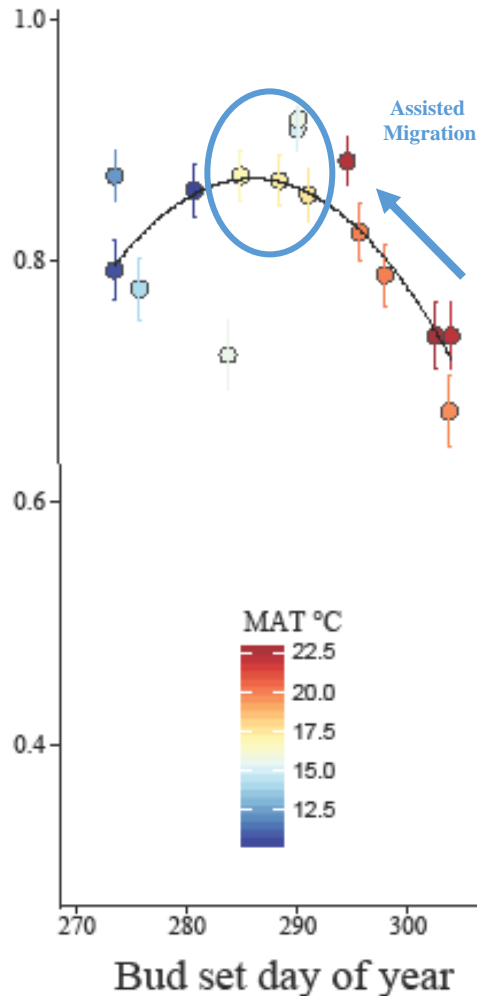
**4600 tree common garden on Arizona  
Game & Fish Dept. lands at Horseshoe  
Ranch surrounded by Agua Fria  
National Monument.**



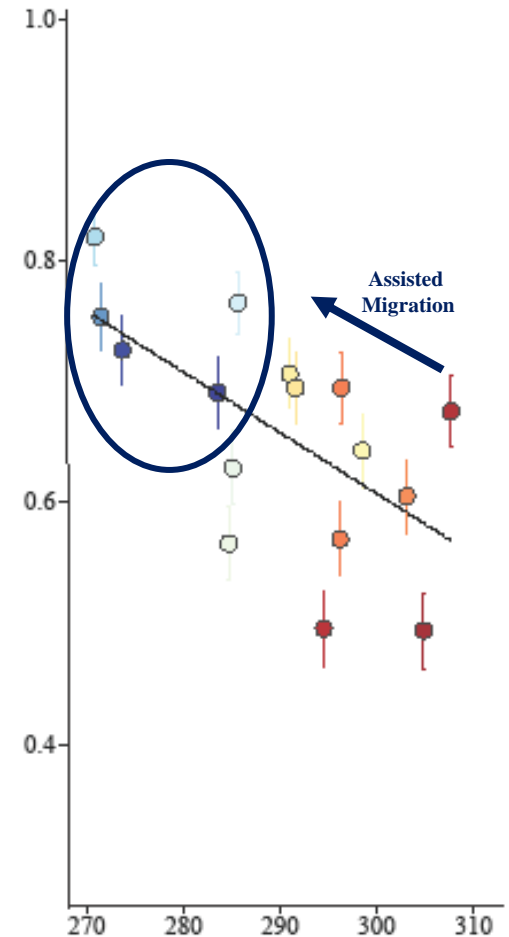
## Hot Garden Long Growing Season



## Intermediate



## Cool Garden Short Growing Season



**Solution #1 – Use populations from other sites that are already adapted to what the new environment will become.**

Population level mean ( $\pm 1$  SE) survival correlations with bud set date in each of the three common gardens. Populations are colored by the mean annual temperature (MAT °C) of their source provenance. In Yuma, survival is highest in the hotter source populations and is positively correlated with later bud set. The opposite is true in the coldest Canyonlands garden. Cooper et al. 2019 Global Change Biology.

**Genotype 146**

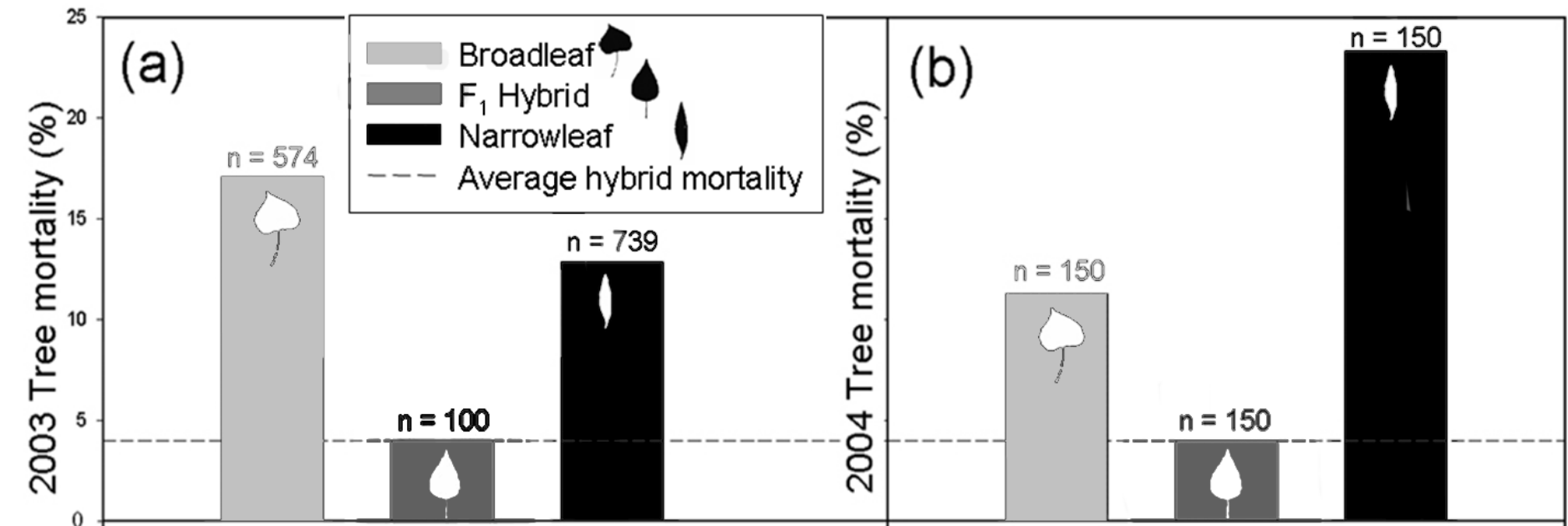


**Genotype 139**



**Solution #2 – Genetics-based differences in architecture make upright and spreading trees more competitive with invasive tamarisk**  
**Sean Mahoney et al. 2018 Restoration Ecology; Photos by Heather Gillette**

# Solution # 3 – Use naturally occurring hybrids that are more drought adapted than their parental species.



Woolbright et al. 2014 Trends in Ecology & Evolution; Gitlin et al. unpub. data



**Solution # 4 – Use  
genotypes from  
desert populations  
that root deeper and  
faster than genotypes  
from higher elevation  
populations.**

**Jackie Parker's rooting  
expt., unpub. data**

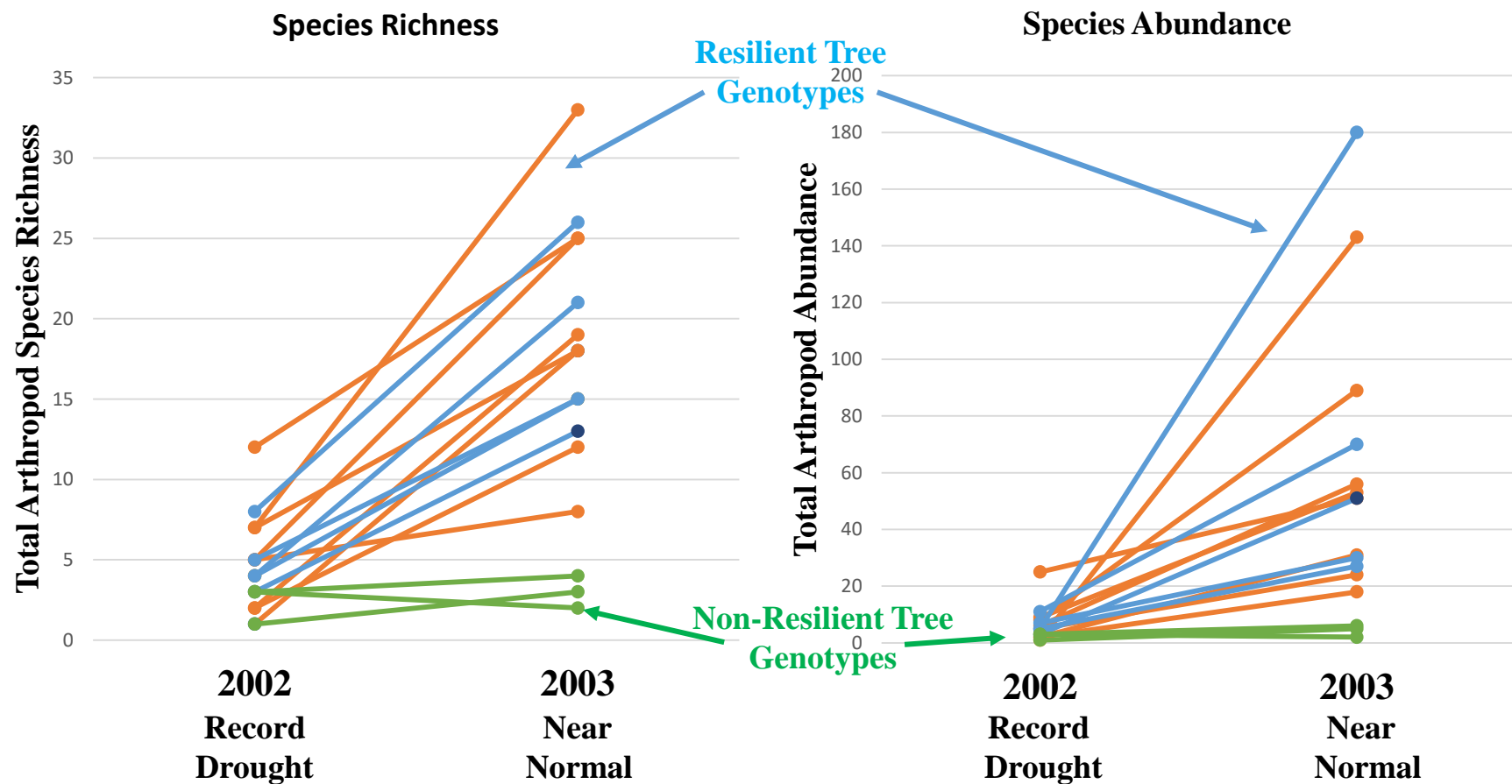
**Seedling  
from  
Drought  
Intolerant  
Mother**



**Seedling  
from  
Drought  
Tolerant  
Mother**

**Solution # 5 – Inoculate plants with better mutualists. A 2000 tree pinyon pine common garden experiment shows that drought tolerance is genetically based and the mycorrhizae on drought tolerant trees are better mutualists (Gehring et al. 2017 PNAS). Similar findings with cottonwoods (Markovchick unpub. data)**

# Solution # 6 – Use tree genotypes that are very plastic (resilient) with environmental change.



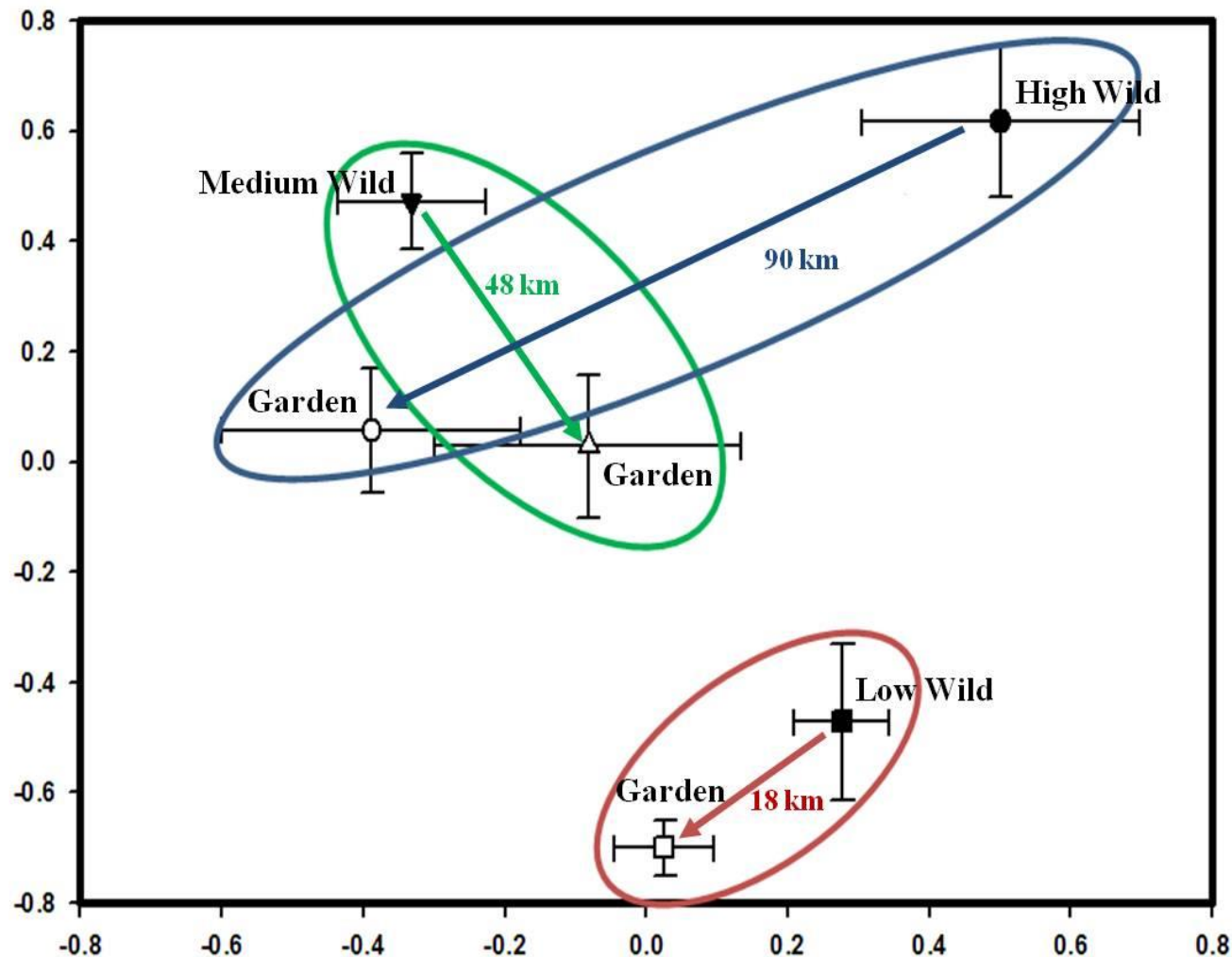
Reaction norms show significant G, E, and G x E interactions in the arthropod communities of *Pinus edulis* (Stone et al. 2019 Frontiers in Plant Science).



**Solution # 7 – Assisted migration distances should be as short as possible to assist migrating communities.**

**A fundamental issue in assisted migration is if you move plants to mitigate the impacts of climate change, will plants acquire the communities of their home sites? In other words, if you build it will they come?**

**Up to a point, if you built it they will come.**



**With transfers of 18 and 48 km, garden and wild trees support similar communities, but at 90 km they are quite different (Keith et al. unpublished data).**



**Removing invasive tamarisk and camelthorn on the Little Colorado River and restoring using new genetic guidelines with the support of Babbitt Ranches, the Nina Mason Pulliam Charitable Trust and the Wildlife Conservation Society.**



# Collaborators In Genetics-Based Restoration

Rachel Adams – plant ecology  
 Joe Bailey – community ecology  
 Davis Blasini – ecophysiology  
 Abraham Cadmus – ecophysiology  
 Hillary Cooper – phylogenetics  
 Rodolfo Dirzo – community ecology  
 Sharon Ferrier – conservation ecology  
 Kevin Floate – insect ecology  
 Kevin Grady – restoration  
 Joakim Hjältén – ecology  
 Dana Ikeda – climate modeling  
 Karl Jarvis – phylogeny  
 George Koch – ecophysiology  
 Jamie Lamit – microbial ecology  
 Rick Lindroth – chemical ecology  
 George Newcombe – plant pathology  
 Brad Potts – quantitative genetics  
 David Smith – landscape ecology  
 Amy Whipple – ecological genetics  
 Gina Wimp – community ecology  
 Scott Woolbright – molecular genetics  
 Matt Zinkgraf – molecular genetics

Gery Allan – molecular ecology  
 Randy Bangert – biogeography  
 Helen Bothwell – phylogeography  
 Aimée Classen – soil ecology  
 Sam Cushman – landscape genetics  
 Chris Doughty – remote sensing  
 Dylan Fischer – ecophysiology  
 Catherine Gehring – microbial ecology  
 Steve Hart – ecosystem/soil ecology  
 Lisa Holeski – genetics & chemistry  
 Julia Hull – fungal endophytes  
 Joann Jeplawy – aquatic ecology  
 Tom Kolb – plant physiology  
 Matthew Lau – network modeling  
 Lisa Markovchick – microbial ecology  
 Emily Palmquist – hydrology  
 Temuulen Sanki – remote sensing  
 Steve Shuster – theoretical genetics  
 Tom Whitham – community ecology  
 Todd Wojtowicz – litter arthropods

Petter Axelsson – transgenic trees  
 Rebecca Best – ecology & evolution  
 Posy Busby – ecological plant pathology  
 Zacchaeus Compson – aquatic ecology  
 Steve DiFazio – molecular ecology  
 Luke Evans – population ecology  
 Paul Flikkema – systems engineering  
 Heather Gillette – molecular ecology  
 Erika Hersch – ecological genetics  
 Kevin Hultine – invasive species  
 Nathalie Isabel – molecular ecology  
 Art Keith – insect community ecology  
 Lela Andrews – molecular ecology  
 Carri LeRoy – aquatic ecology  
 Nashelly Meneses – ecological genetics  
 Jackie Parker – plant ecology  
 Jen Schweitzer – ecosystems  
 Chris Sthultz – plant ecology

Outreach – Lara Schmit, Victor Leshyk - NAU



Macrosystems  
MRI

