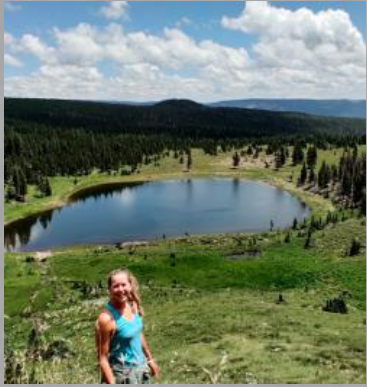


Landscape
Stewardship
Collective

Maintaining genetic diversity during the agricultural increase of native seeds: insights from workhorse grass species

Rob Massatti & Trevor Faske
Landscape Stewardship Collective
February 25th, 2026

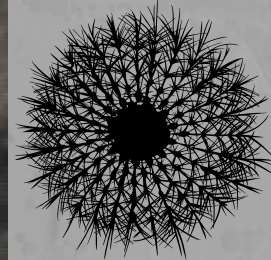




Daniel Winkler
Sasha Reed



Sophia Lasche
Akasha Faist



Landscape
Stewardship
Collective

Trevor Faske



Sara Fuentes-Soriano



Ashlee Wolf

Restoration – an inherently difficult process

Site specific challenges

- Land use history
- Invasive species
- Ongoing disturbance
- Weather
- Climate (aridlands notoriously difficult)

Procedural challenges

- How do we restore?
 - Seeding rate, seed coatings, weather forecasting, seed application, etc...
- What should we use for restoration?
 - Local adaptation, species mixes, composite sources vs. cultivars, etc...
- How do we account for changing climates?
 - Present climate, future climate, or both?

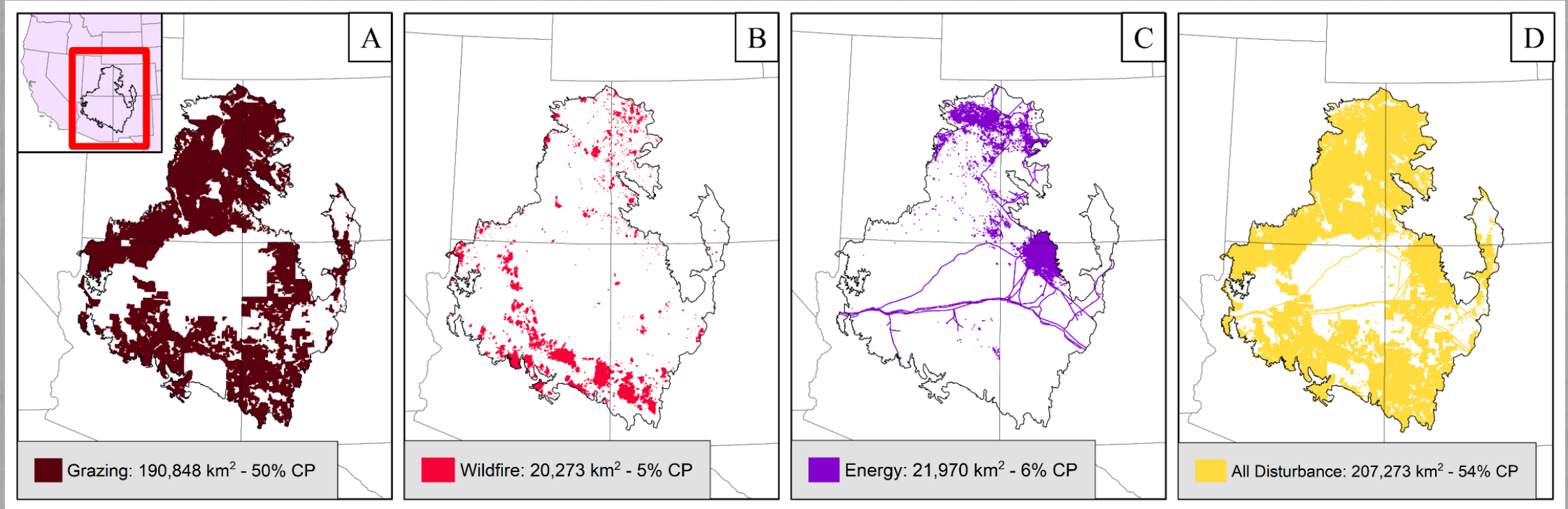


Cameron Peak Fire. Courtesy photo USDA Forest Service.



Milford Flat Fire. Photo: M.E. Miller, NPS. Public domain.

Disturbances are common!



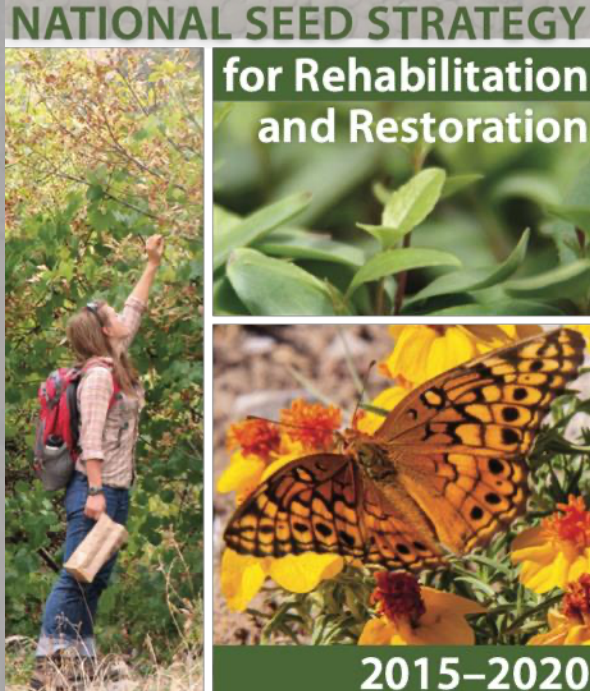
Winkler et al., 2025

- On average, 2.4 million pounds of seed are purchased annually for use on BLM land
- Average annual cost: \$20 million

The National Seed Strategy

Mission: To ensure the availability of *genetically appropriate* seed to restore viable and productive plant communities and sustainable ecosystems.

Vision: The right seed in the right place at the right time.



www.blm.gov/programs/natural-resources/native-plant-communities/national-seed-strategy

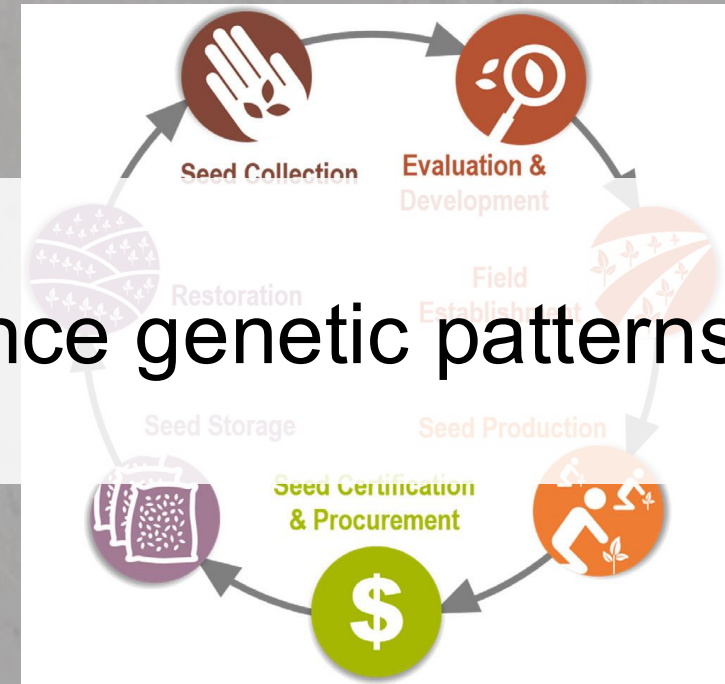


McCormick et al. 2021

The National Seed Strategy

Mission: *To ensure the availability of **genetically appropriate** seed to restore viable and productive plant communities and sustainable ecosystems.*

Vision: *The right seed in the right place at the right time.*

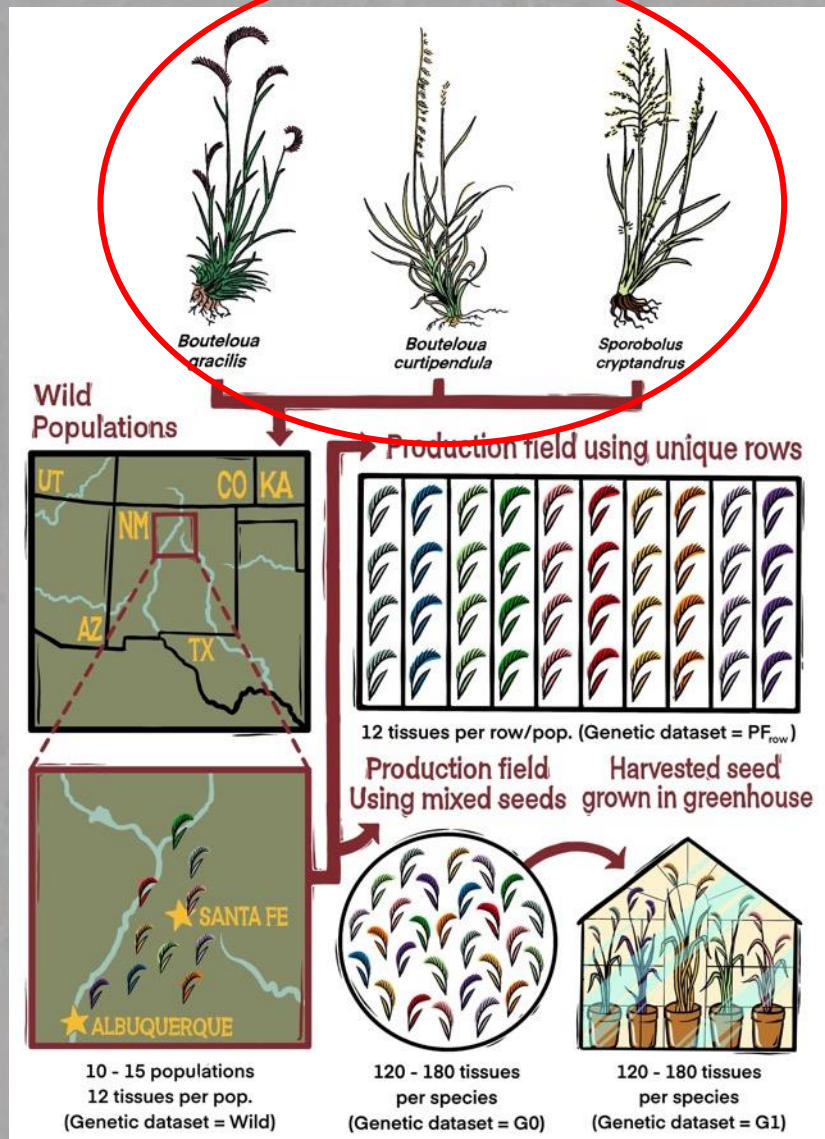


How does seed production influence genetic patterns?

www.blm.gov/programs/natural-resources/native-plant-communities/national-seed-strategy

McCormick et al. 2021

Investigating the effects of seed increase



3 species:

- *Bouteloua gracilis* (blue grama grass)
- *Bouteloua curtipendula* (sideoats grama grass)
- *Sporobolus cryptandrus* (sand dropseed)



Bouteloua gracilis

Polyploidy (2N = 20, 21, 24,
28, 35, 40, 42, 60, 61, 77, 84)



Bouteloua curtipendula

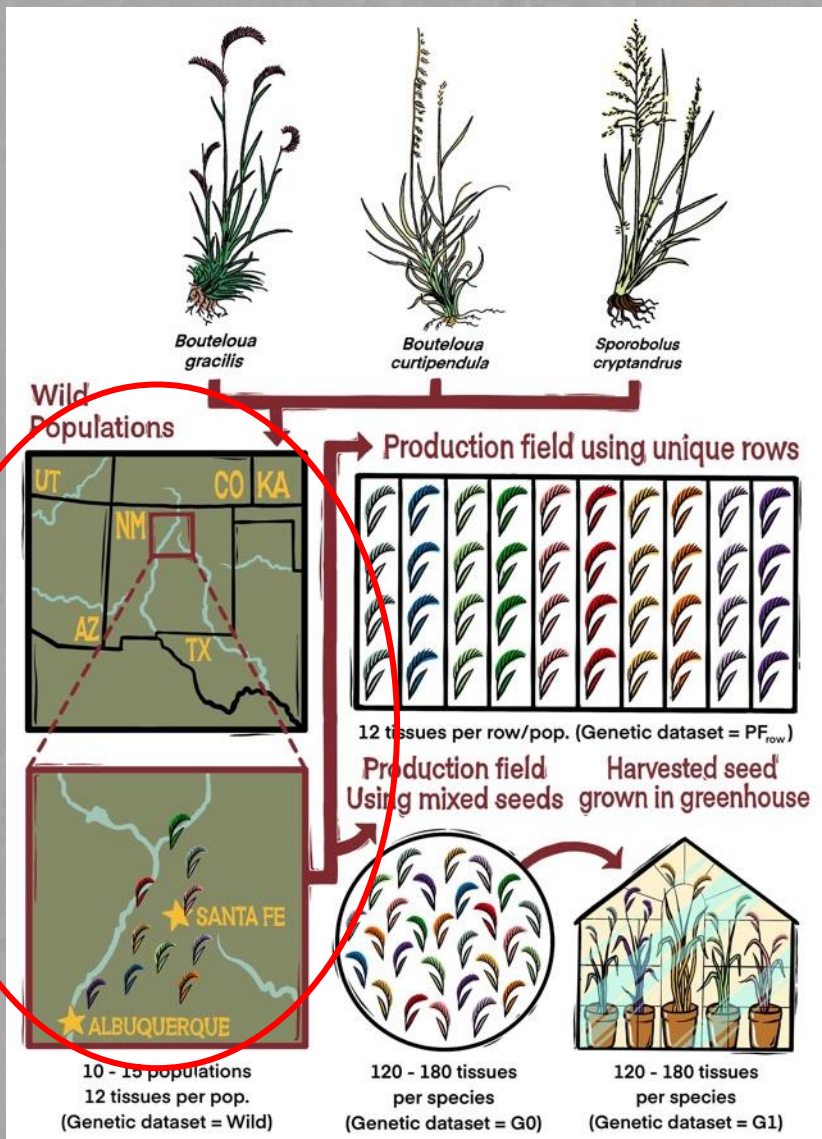
Polyploidy (2N = 20, 28, 35, 40-43, 41-
64, 42, 45, 52, 56, 58-64, 69-103, 70,
74, 80, 82, 85-101, 86, 96, 98)



Sporobolus cryptandrus

Self-fertilizing

Investigating the effects of seed increase

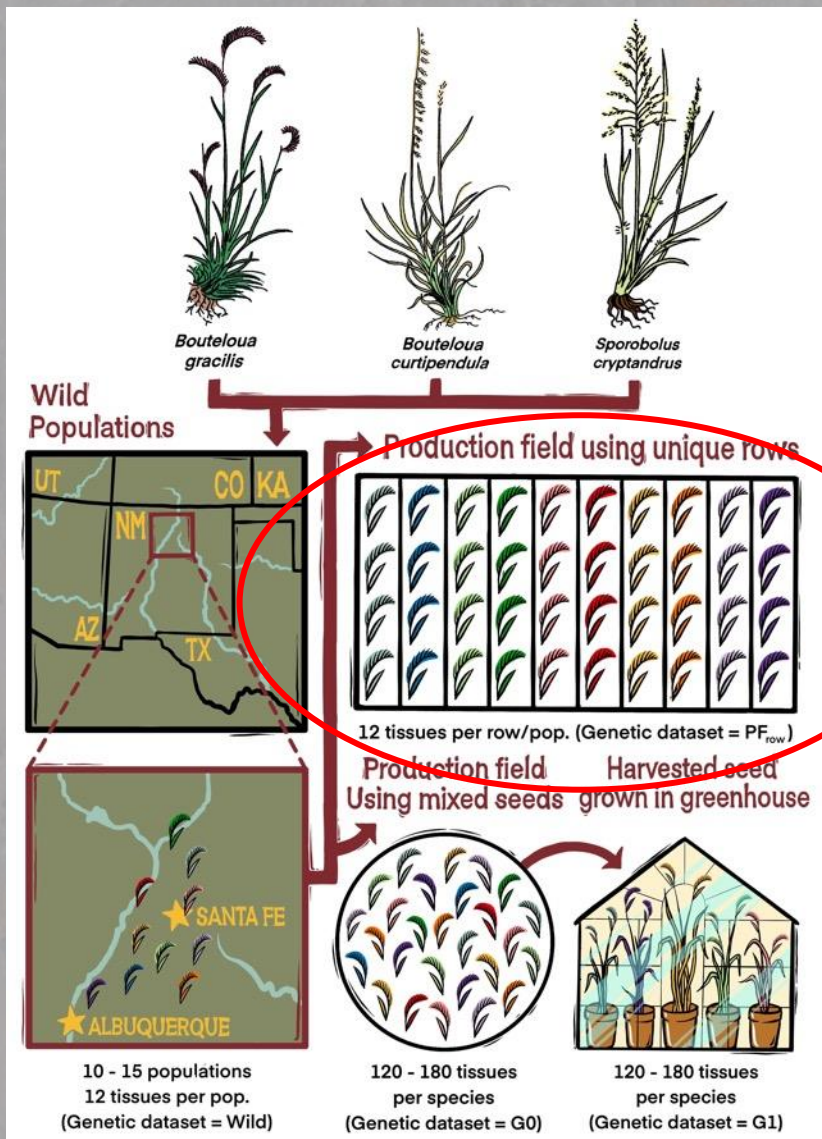


Seed collected from 10-15 wild populations

- Seeds of Success protocols
- Institute for Applied Ecology & NM BLM
- Environmentally similar collection sites

Genetic data from wild collected tissues = **Wild**
Collected seed = G0

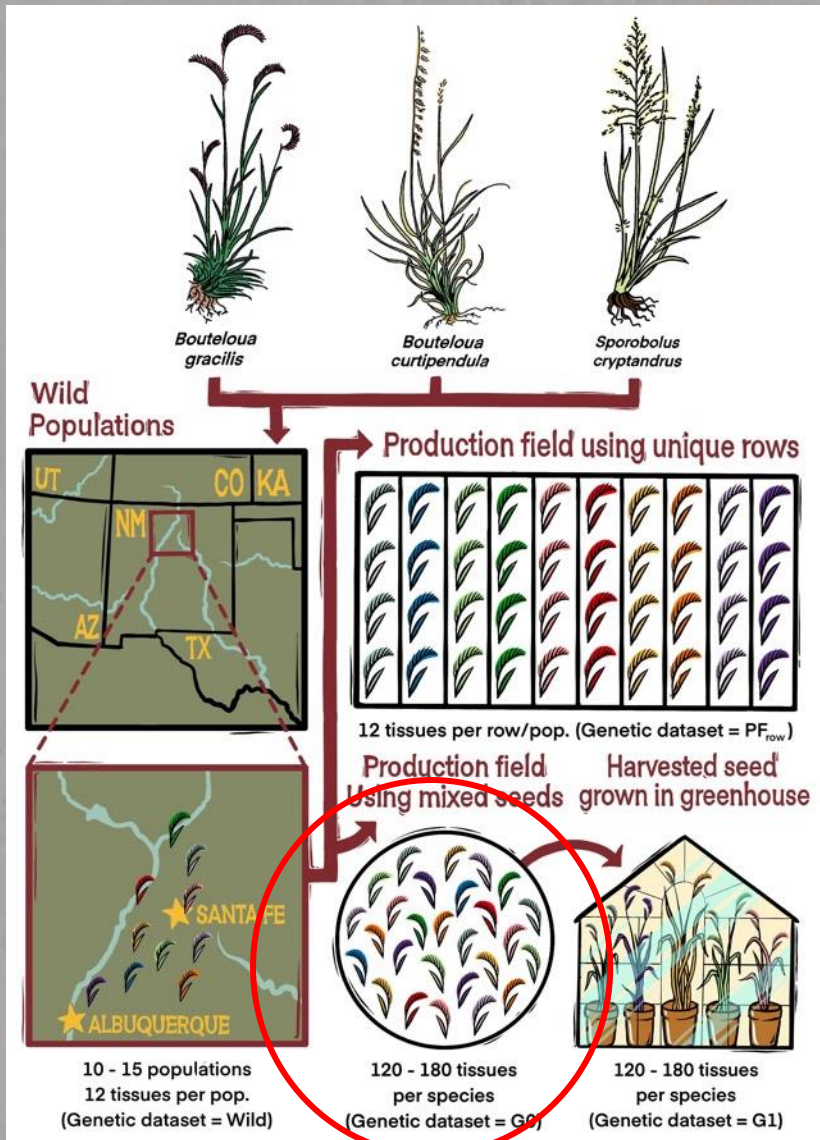
Investigating the effects of seed increase



G0 sources grown separately

- Genetic dataset = **G0 (rows)**

Investigating the effects of seed increase



Faske et al., *In review*

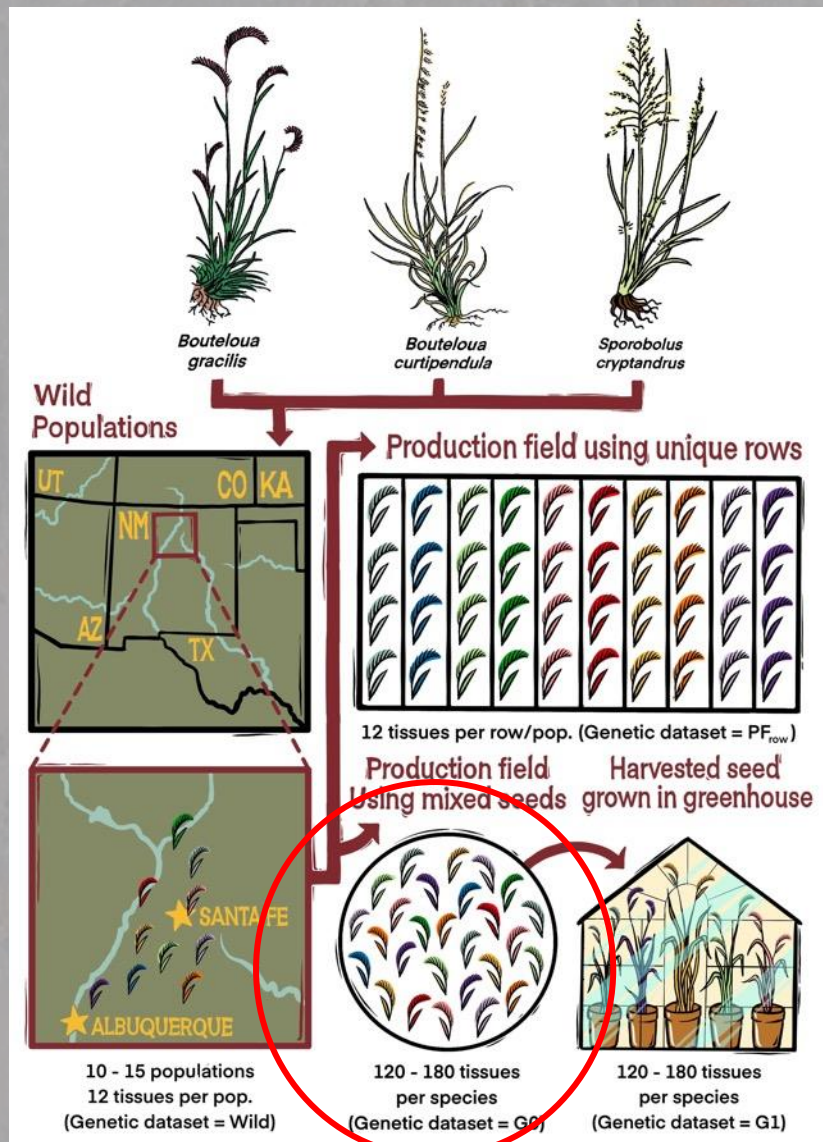
G0 sources grown separately

- Genetic dataset = **G0 (rows)**

G0 sources pooled

- Genetic dataset = **G0 (pooled)**
- **THIS IS THE COMMON SCENARIO**

Investigating the effects of seed increase



***All fields established from greenhouse grown transplants**

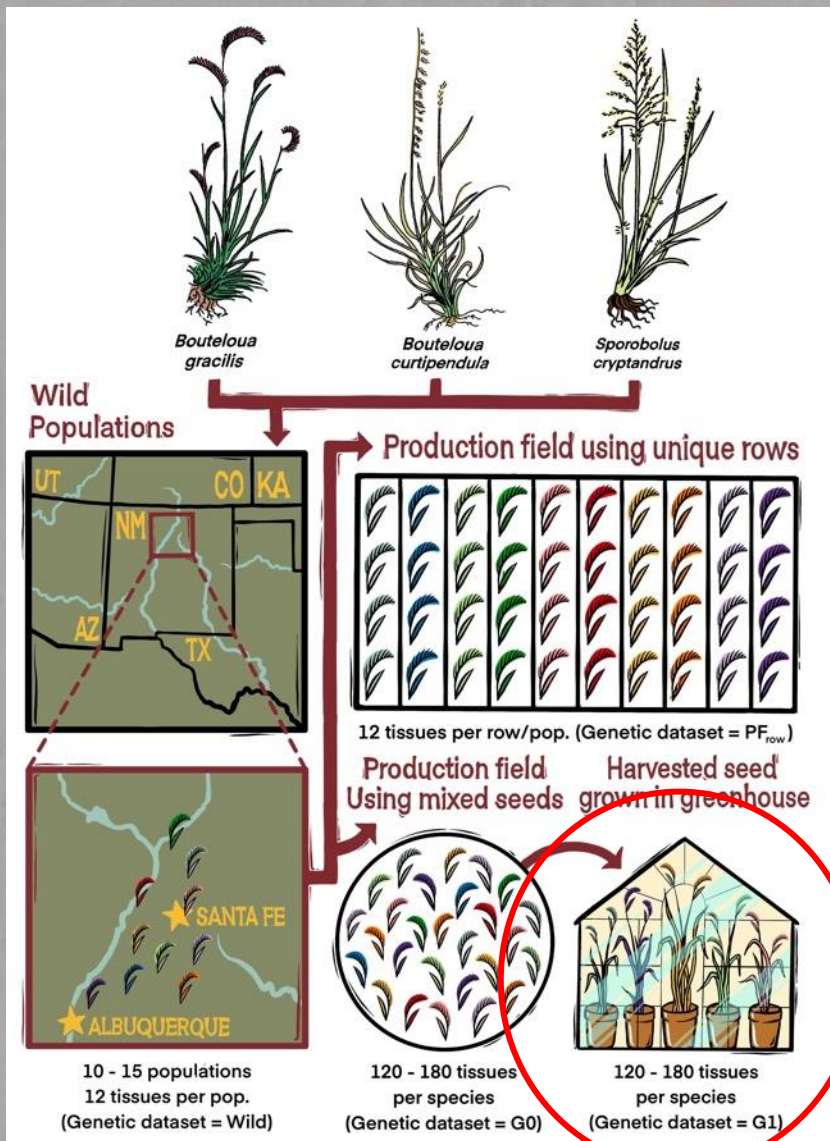
G0 sources grown separately

- Genetic dataset = **G0 (rows)**

G0 sources pooled

- Genetic dataset = **G0 (pooled)**
- **THIS IS THE COMMON SCENARIO**

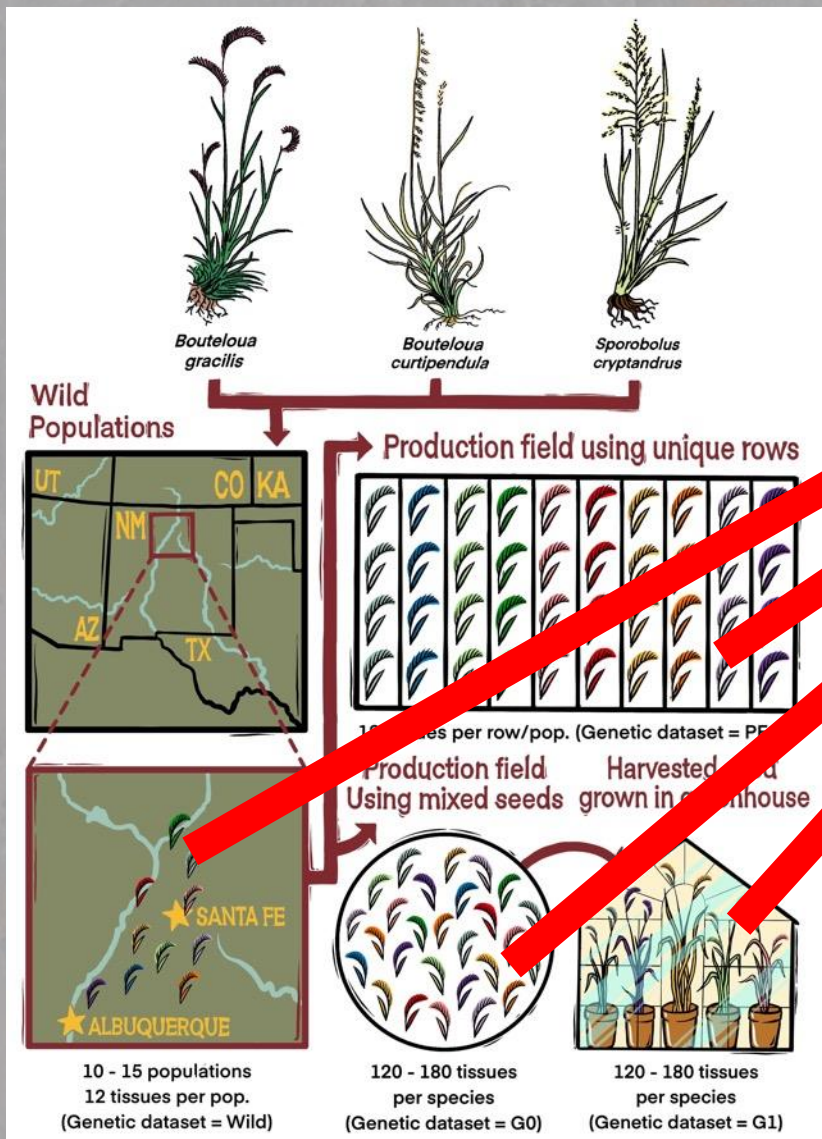
Investigating the effects of seed increase



Faske et al., in review

- G1 seed harvested from G0 pooled
- Genetic dataset = **G1**

Investigating the effects of seed increase



4 datasets per species

- Wild
- G0 (rows)
- G0 (pooled)
- G1

Questions:

How do genetic patterns shift between Wild and G1?
Are there differences among species?



Bouteloua gracilis

15 sources, 555 plants
19,324 genetic markers



Bouteloua curtipendula

10 sources, 557 plants
18,881 genetic markers

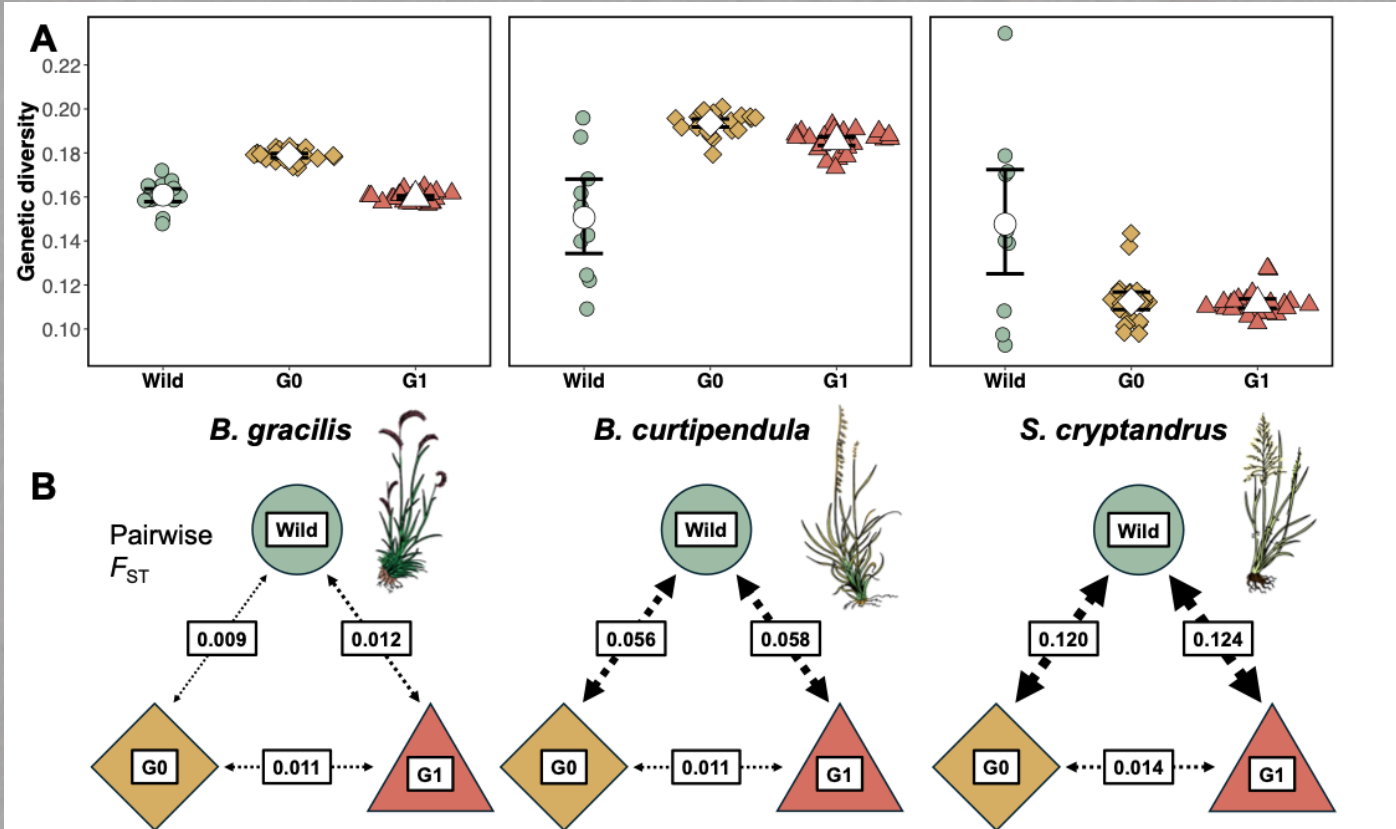


Sporobolus cryptandrus

10 sources, 594 plants
27,925 genetic markers

Results: How do genetic patterns shift from wild sources to harvested seed?

Genetic diversity and differentiation among Wild, G0, and G1 generations



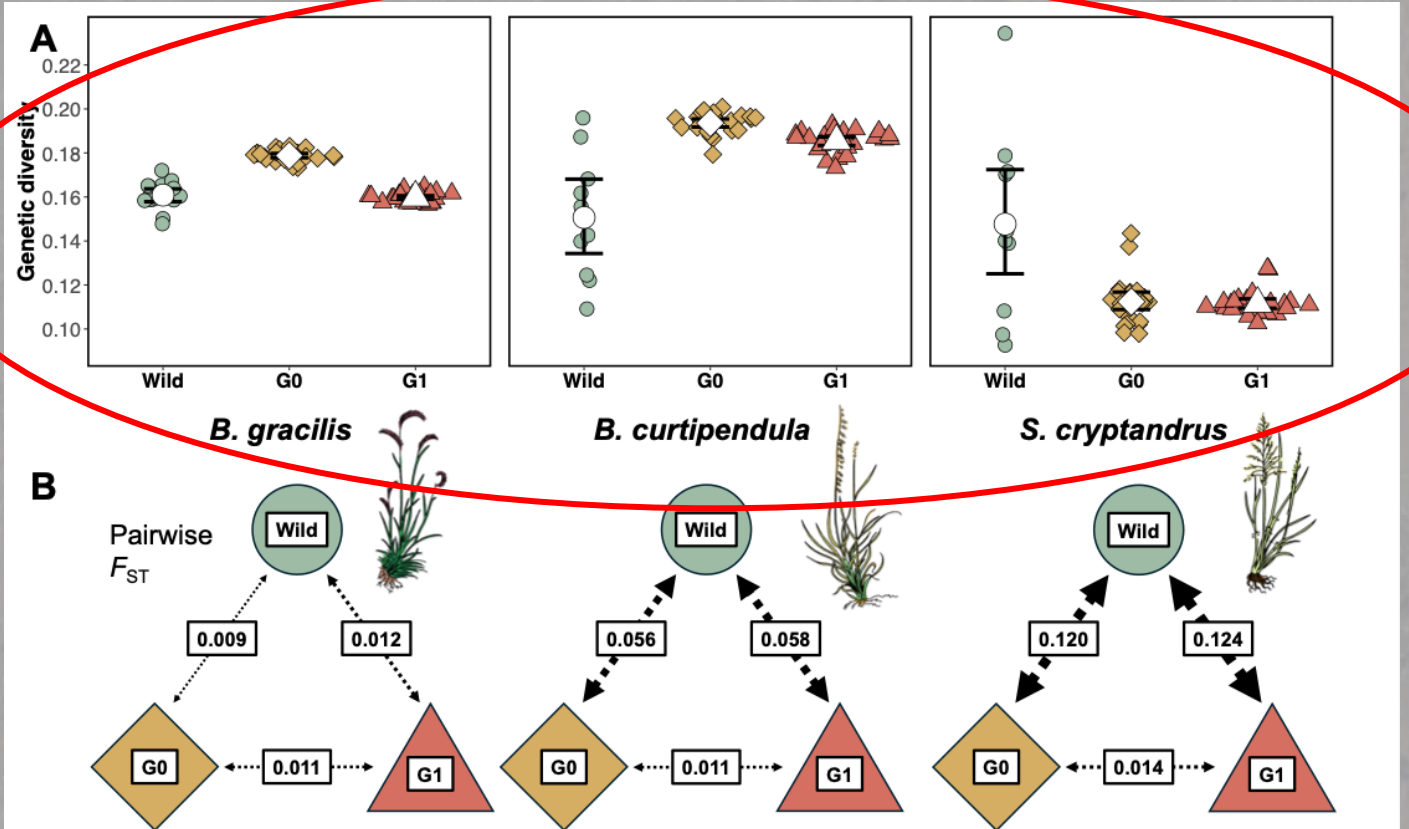
Genetic diversity

Genetic differentiation

Faske et al., *In review*

Results: How do genetic patterns shift from wild sources to harvested seed?

Genetic diversity and differentiation

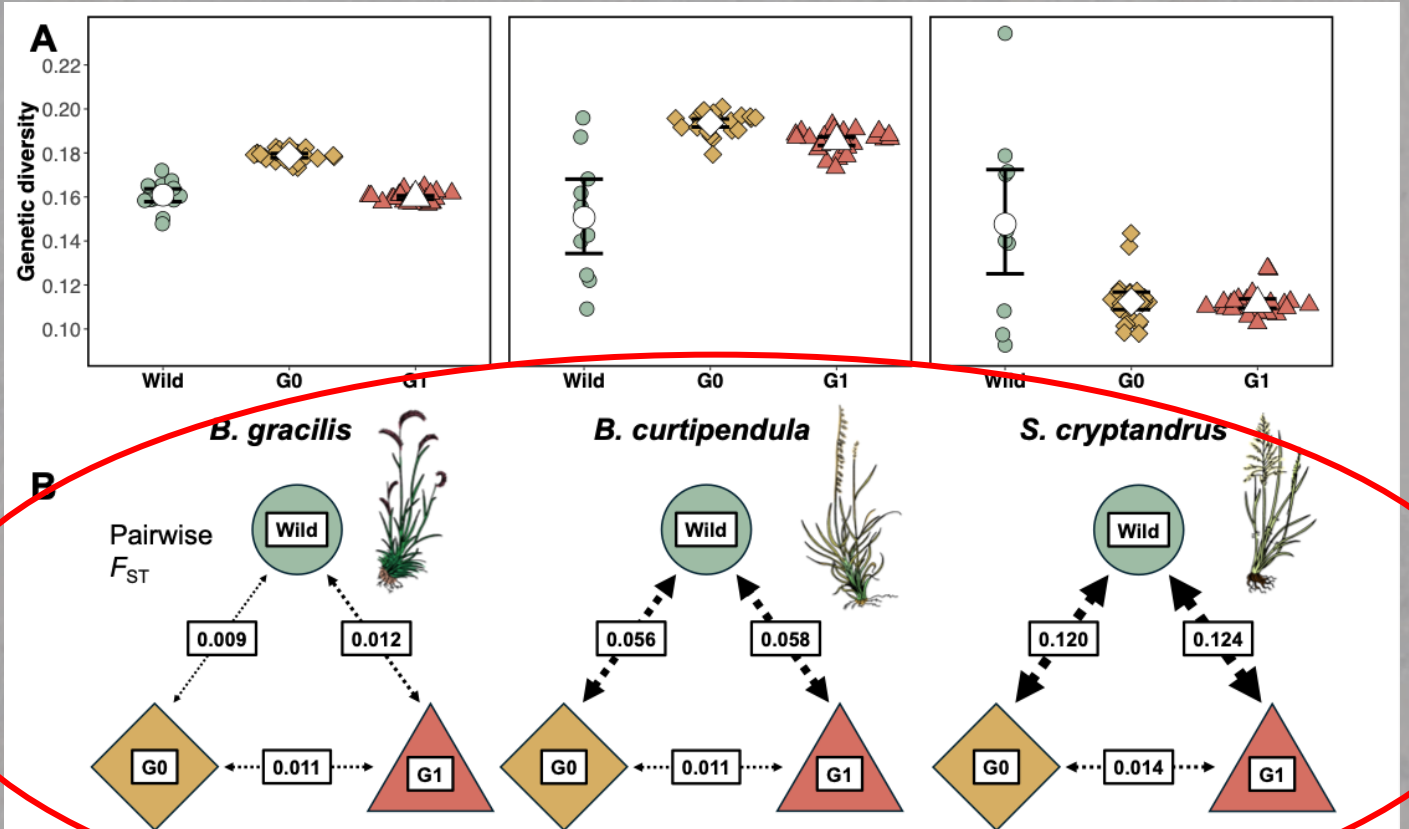


- Genetic diversity patterns across generations are unique

Faske et al., *In review*

Results: How do genetic patterns shift from wild sources to harvested seed?

Genetic diversity and differentiation

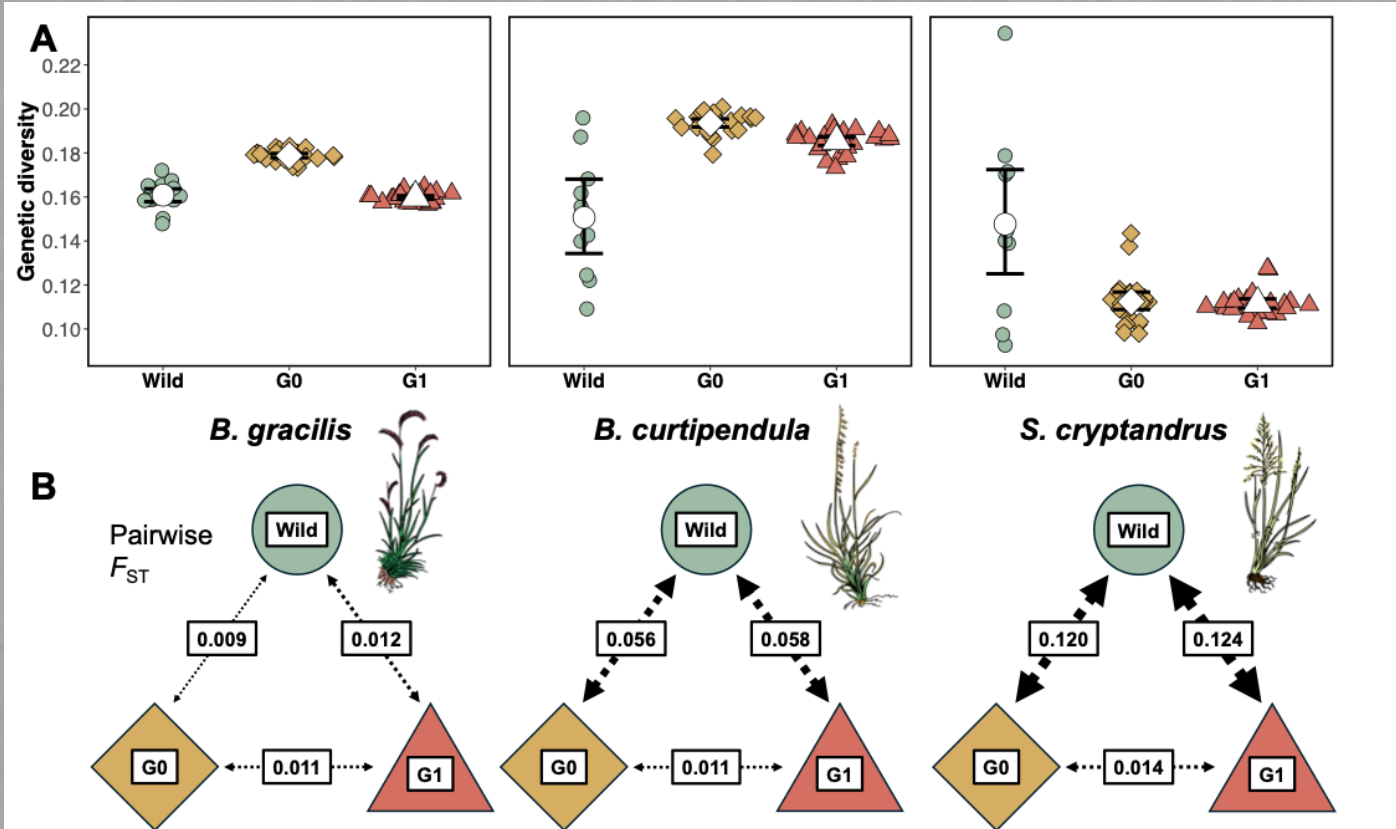


- Genetic diversity patterns across generations are unique
- Genetic differentiation varies across species between Wild and G0/G1
 - 0 = completely the same
 - 1 = completely different

Faske et al., *In review*

Results: How do genetic patterns shift from wild sources to harvested seed?

Genetic diversity and differentiation

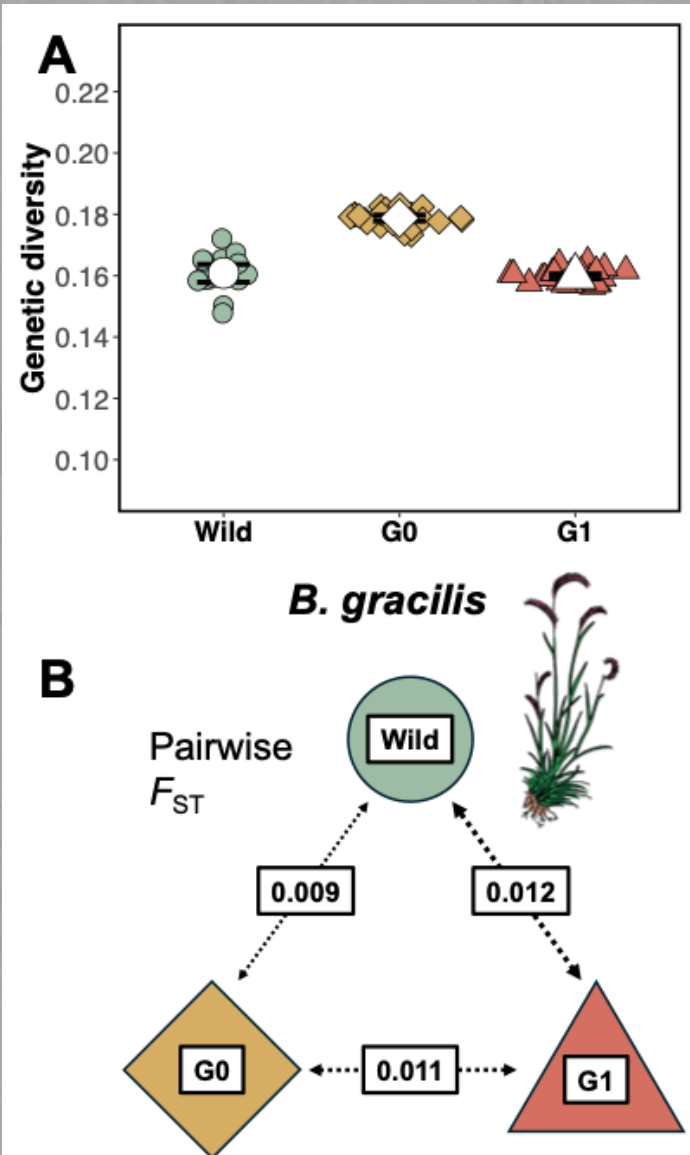


Faske et al., *In review*

- Genetic diversity patterns across generations are unique
- Genetic differentiation varies across species between Wild and G0/G1

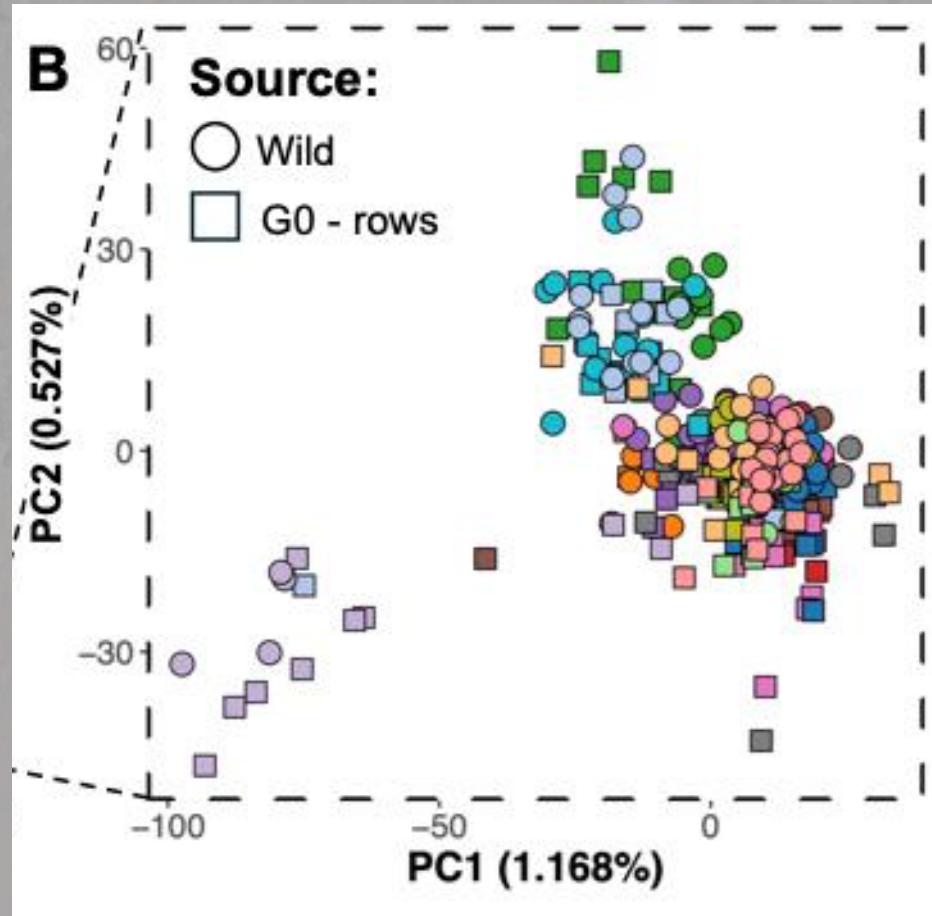
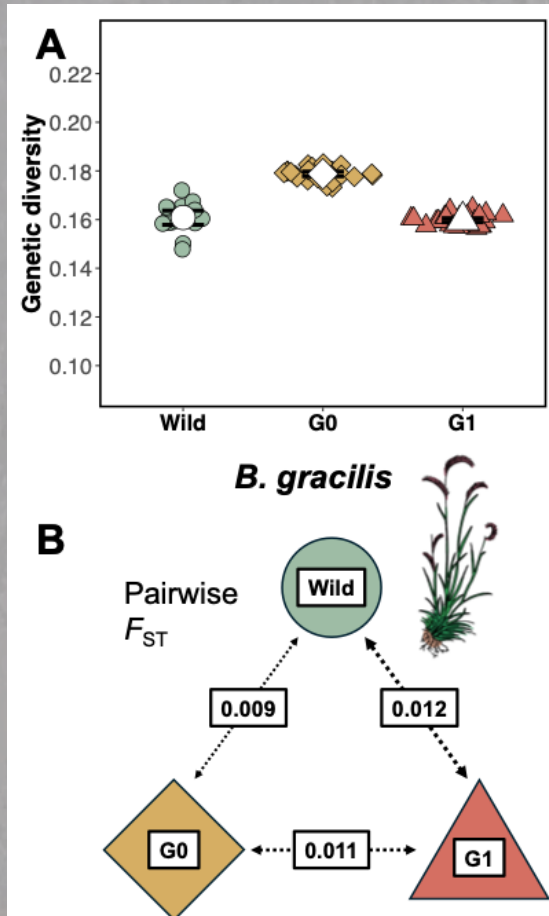
What's driving differences among species?

Results: *Bouteloua gracilis*



- Small changes in genetic diversity among Wild, G0, G1
- Genetic differentiation low

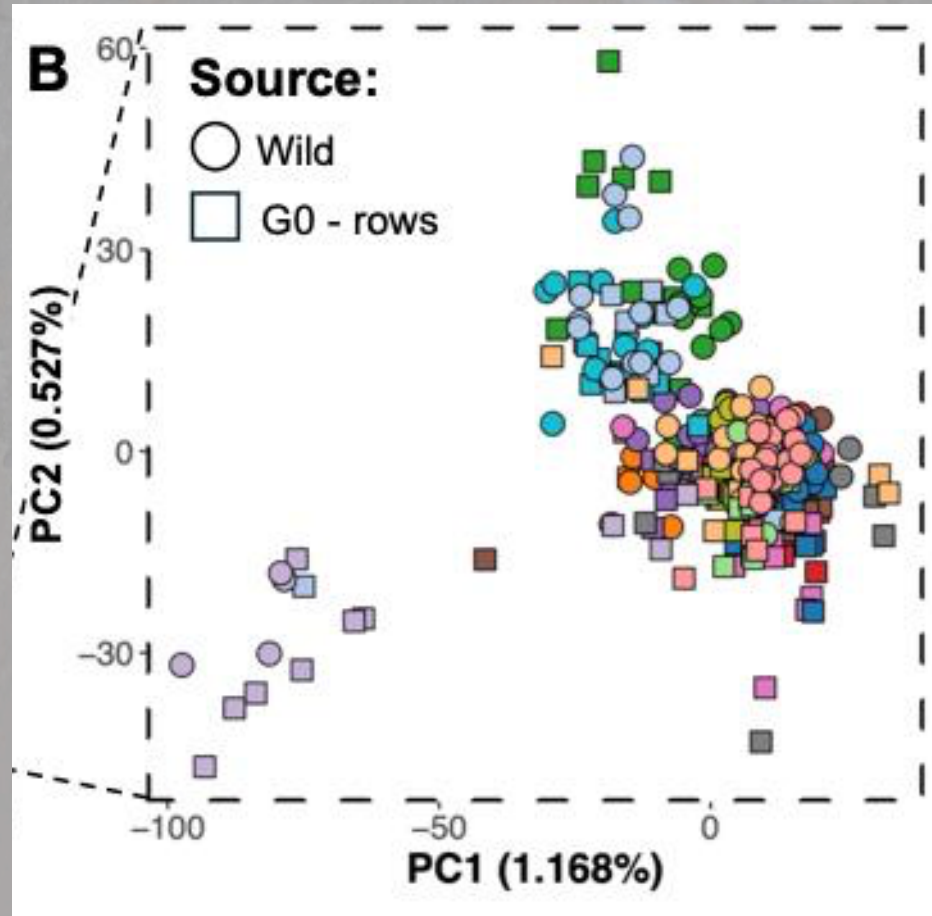
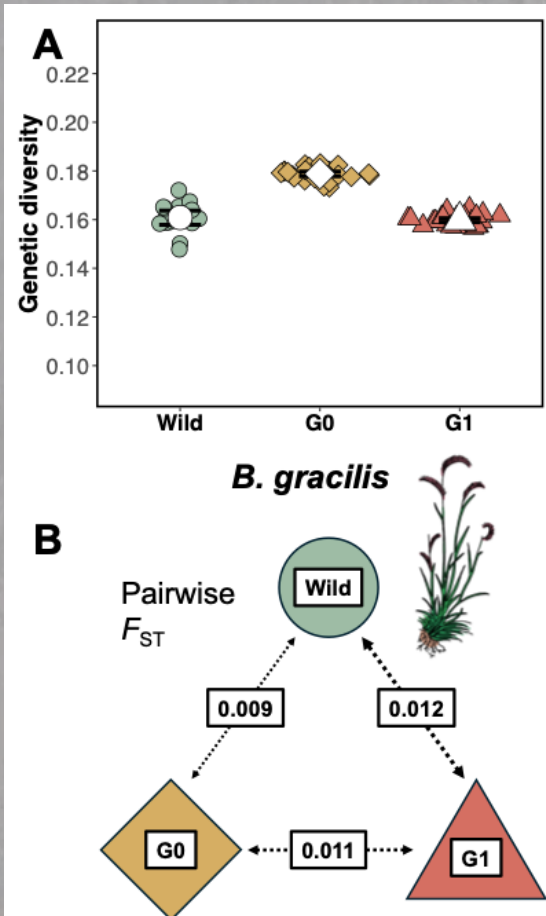
Results: *Bouteloua gracilis*



- Individuals from Wild and G0 overlap according to their source identity

Colors represent source population

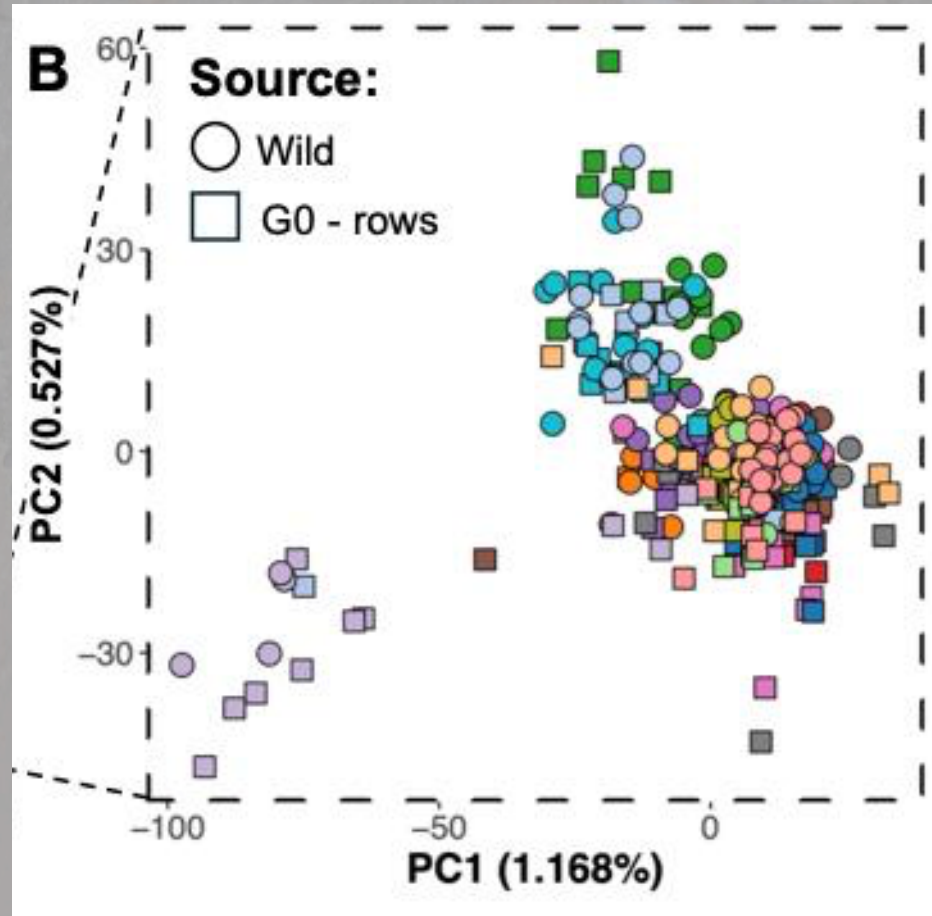
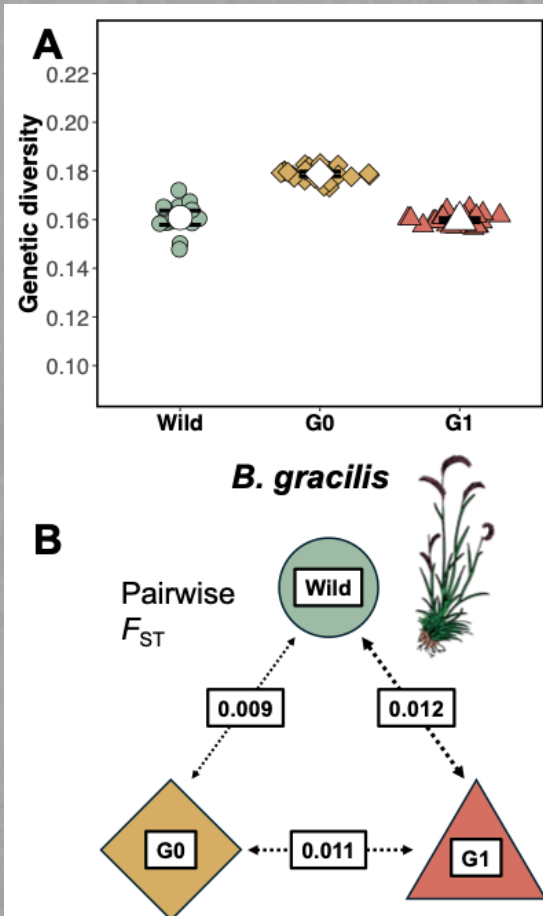
Results: *Bouteloua gracilis*



- Individuals from Wild and G0 overlap according to their source identity
- Meets expectations for an outcrossing species

Colors represent source population

Results: *Bouteloua gracilis*

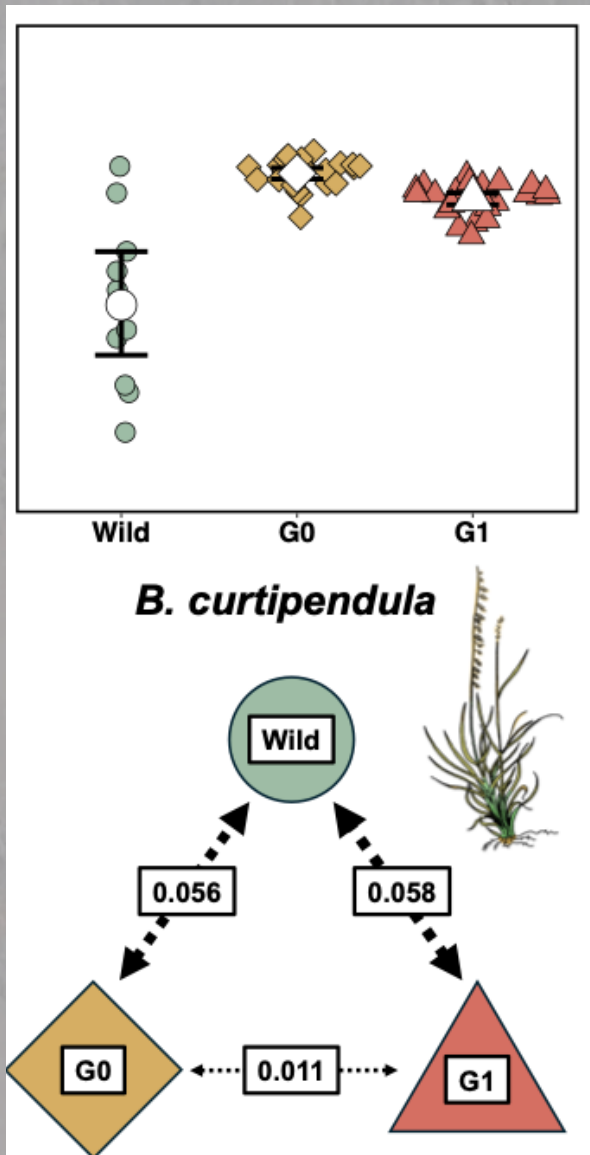


- Individuals from Wild and G0 overlap according to their source identity
- Meets expectations for an outcrossing species

Wild and G0 represent the same genetic pool, hence:

- consistent genetic diversity
- low genetic differentiation

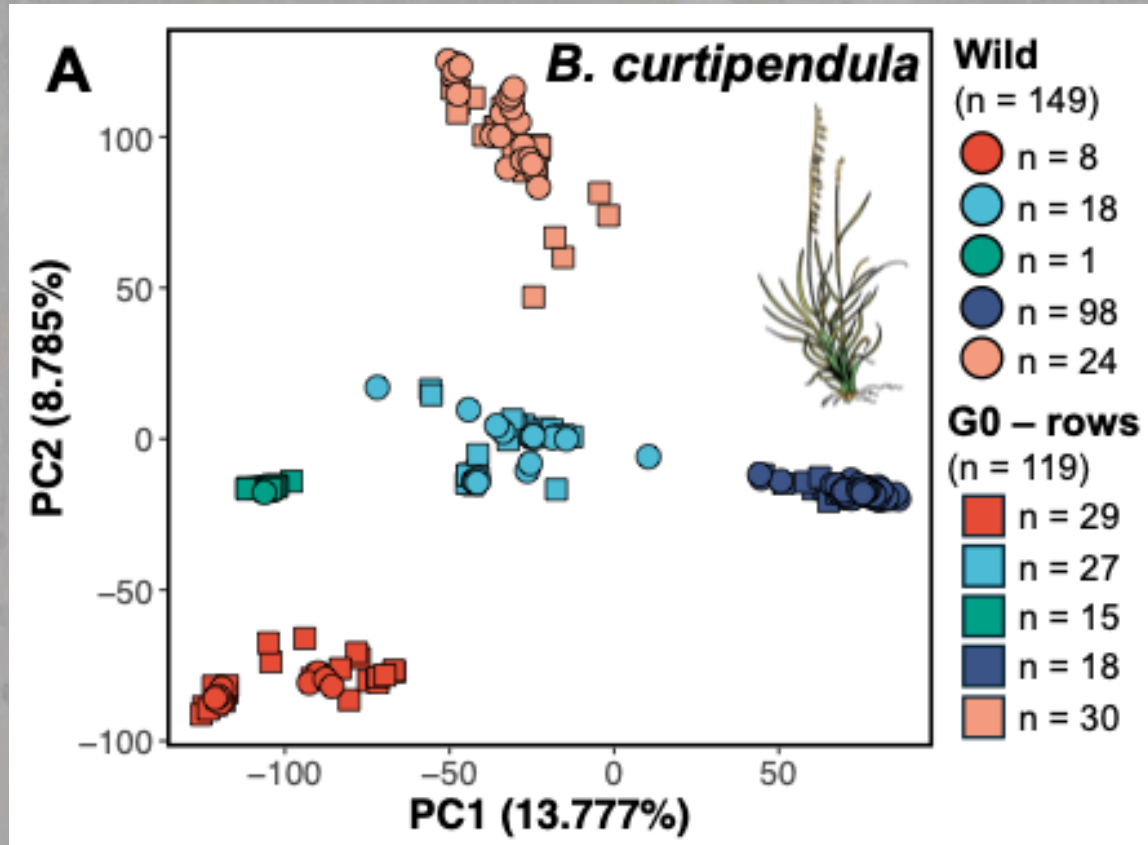
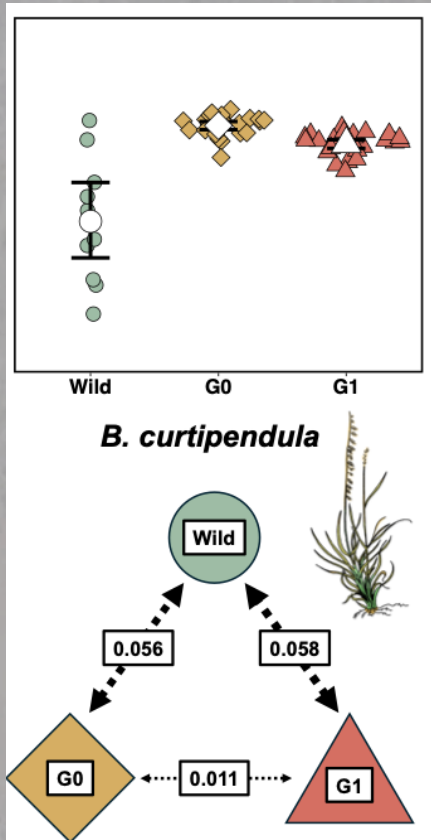
Results: *Bouteloua curtipendula*



- Higher genetic diversity among in G0 and G1
- Wild genetically differentiated from G0/G1

Faske et al., *In review*

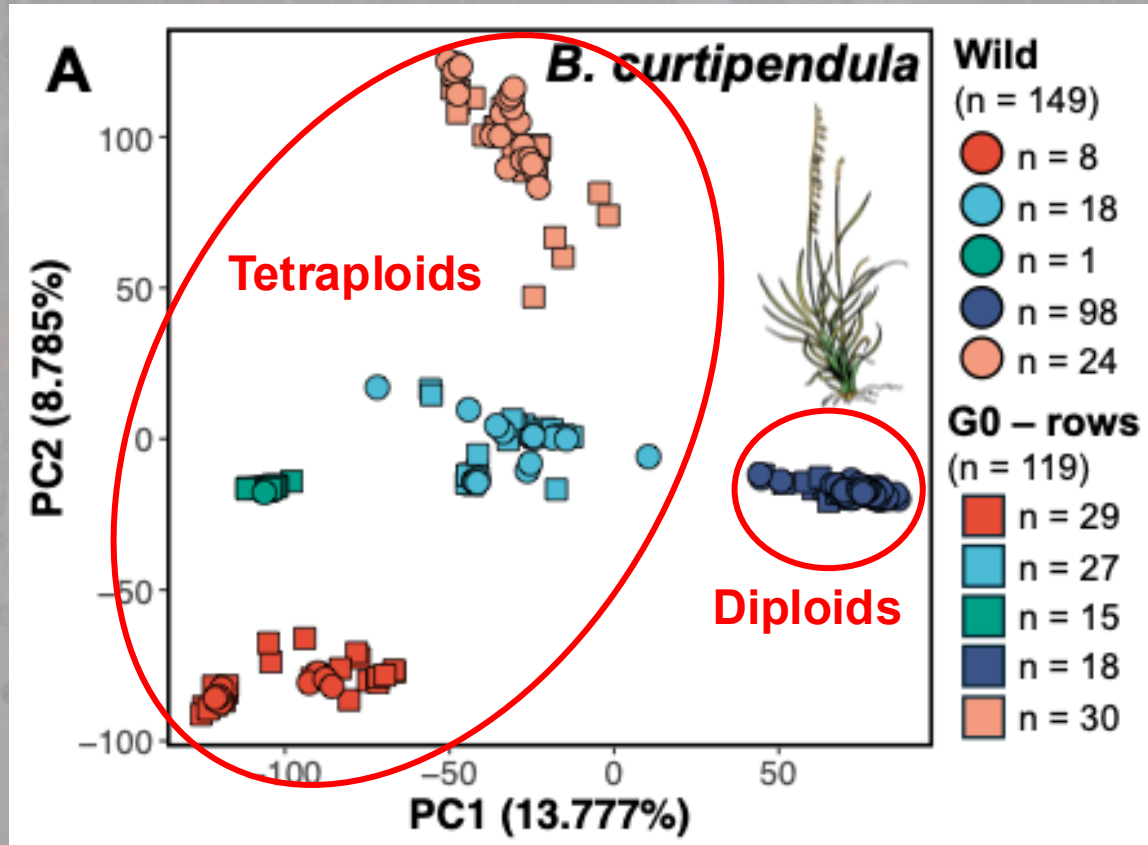
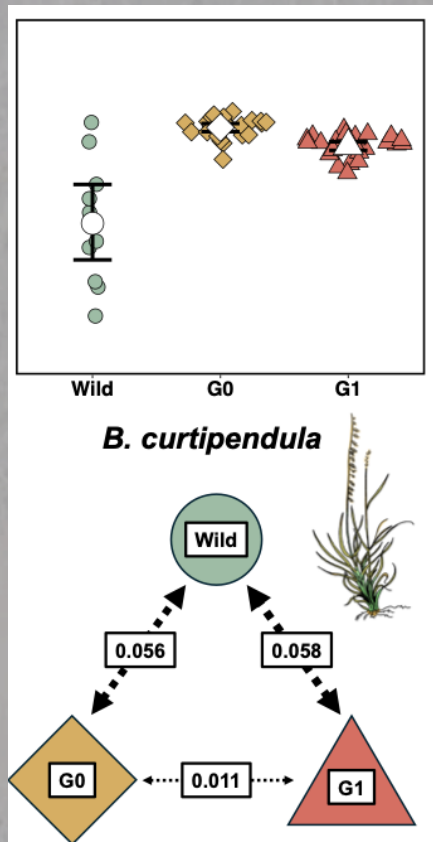
Results: *Bouteloua curtipendula*



- Individuals from Wild and G0 cluster into 5 genetic groups

Colors represent assigned genetic cluster

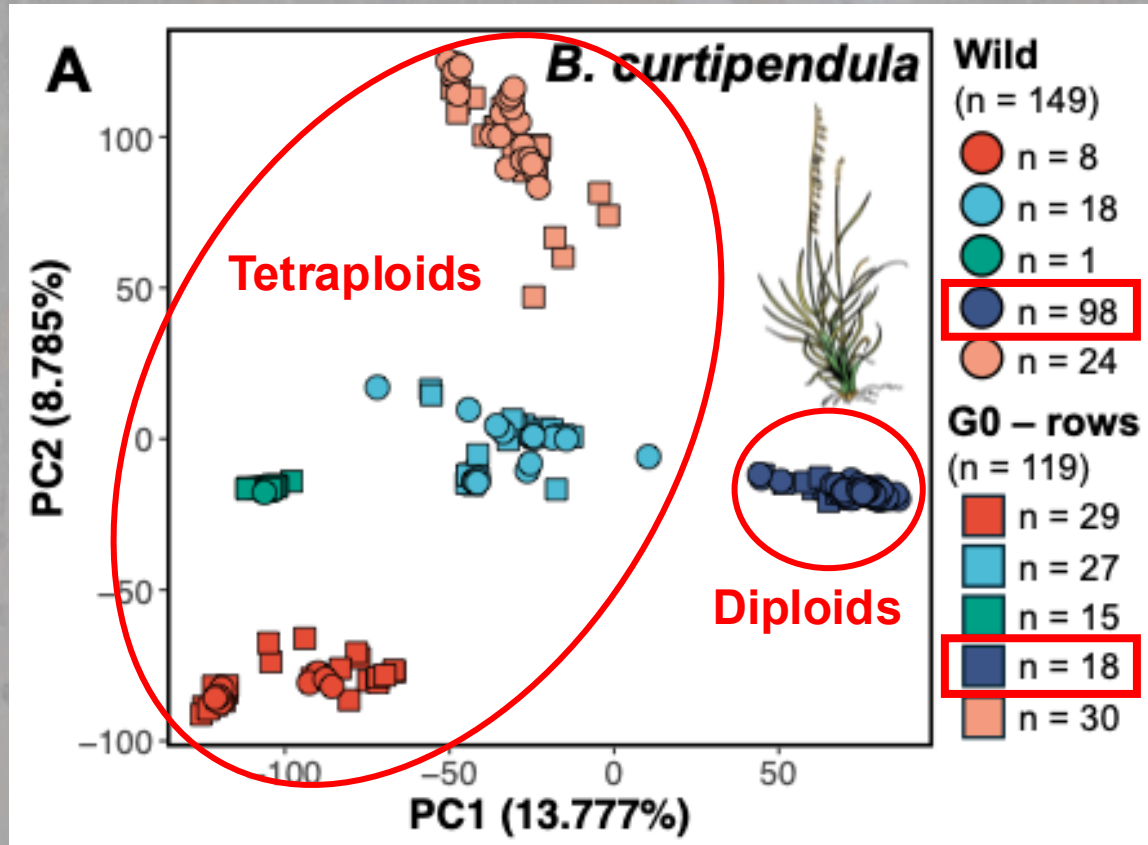
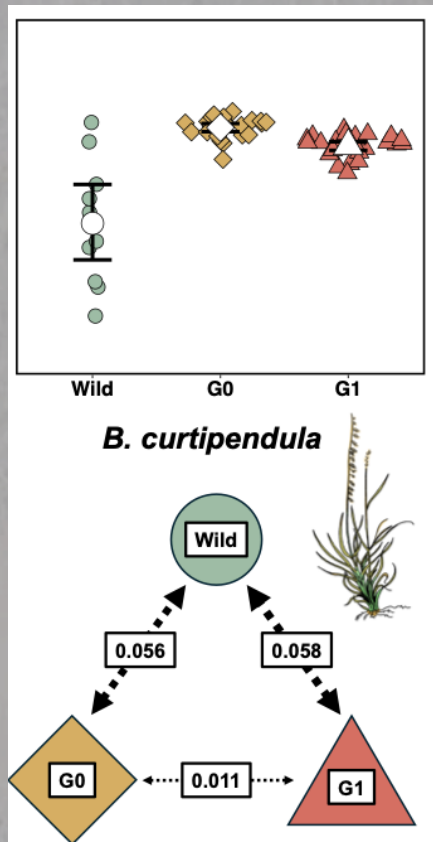
Results: *Bouteloua curtipendula*



Colors represent assigned genetic cluster

- Individuals from Wild and G0 cluster into 5 genetic groups
- Groups represent different ploidy levels

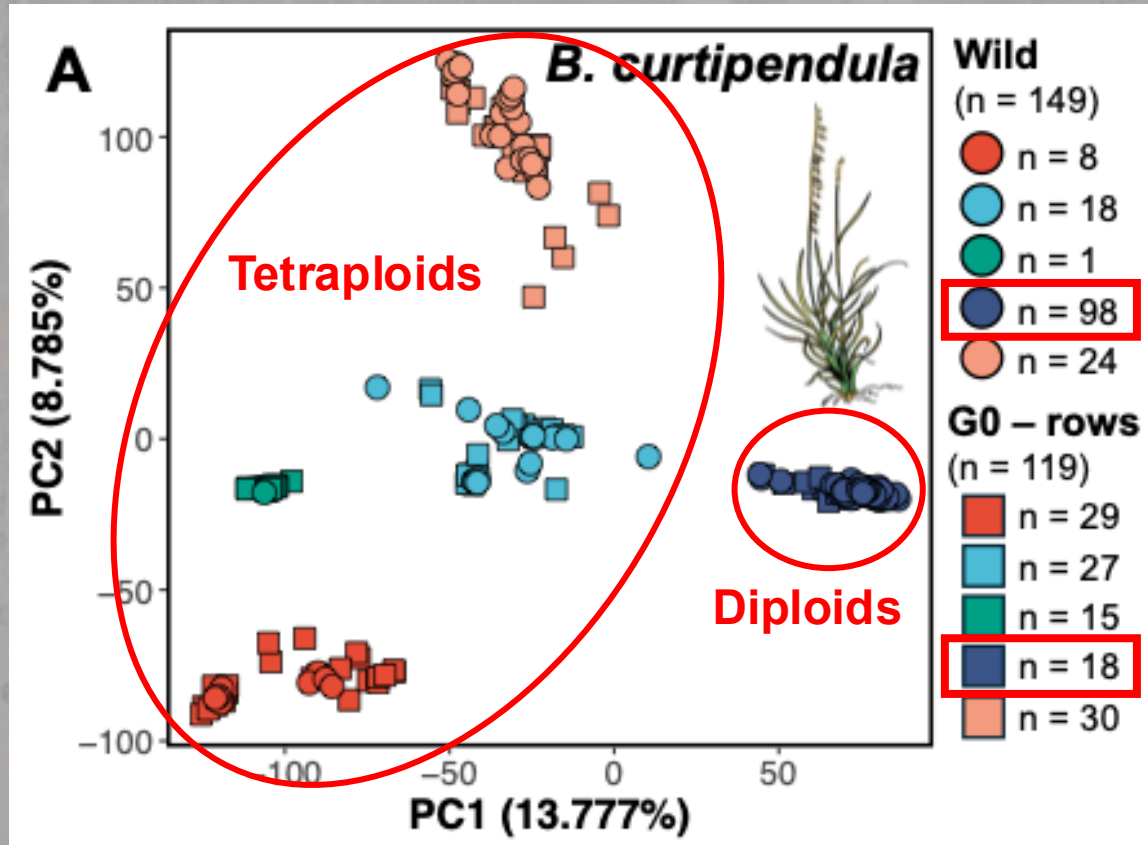
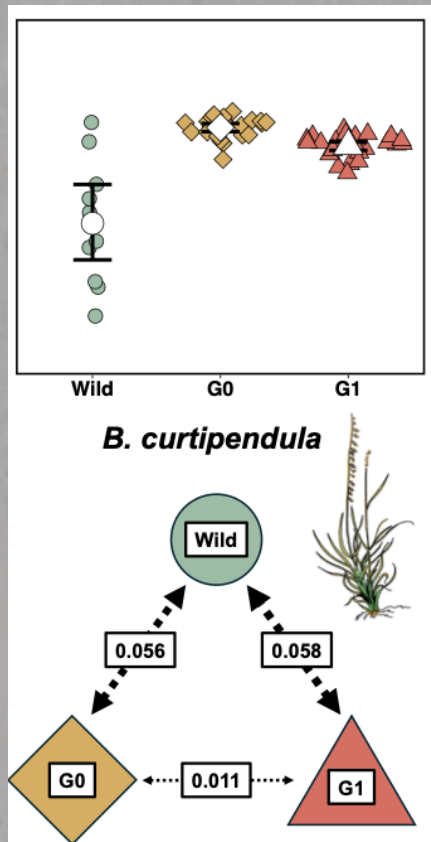
Results: *Bouteloua curtipendula*



Colors represent assigned genetic cluster

- Individuals from Wild and G0 cluster into 5 genetic groups
- Groups represent different ploidy levels
- Tetraploids much more common in G0 compared to Wild!!!

Results: *Bouteloua curtipendula*

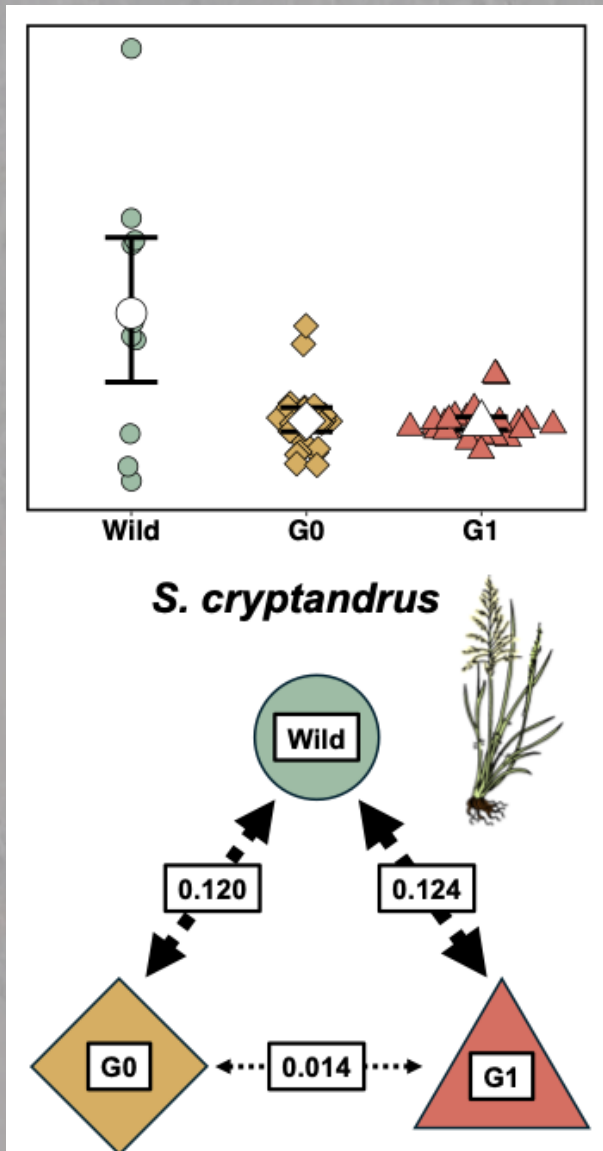


- Individuals from Wild and G0 cluster into 5 genetic groups
- Groups represent different ploidy levels
- Tetraploids much more common in G0 compared to Wild!!!

Tetraploids are overrepresented in agricultural fields, hence:

- higher genetic diversity
- higher genetic differentiation

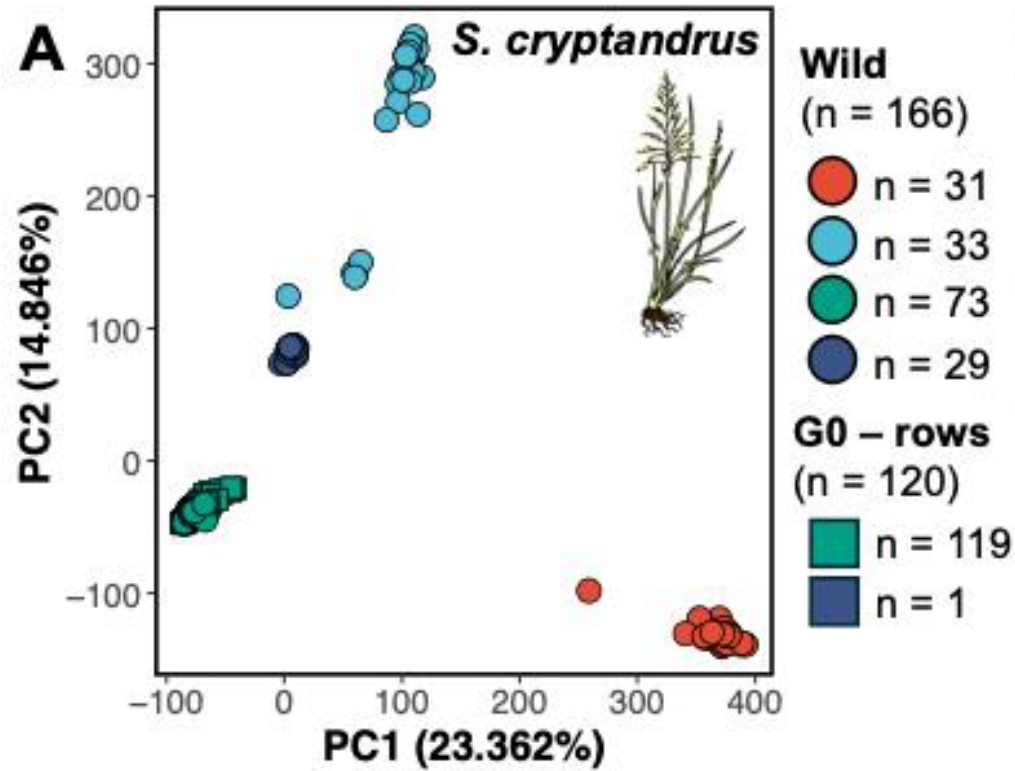
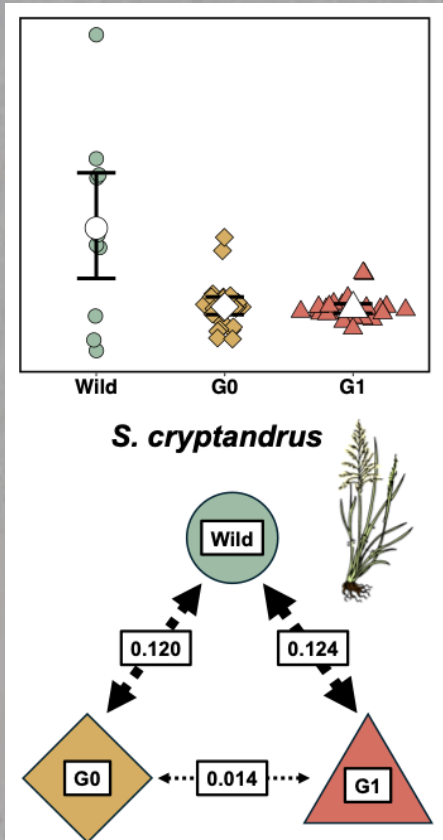
Results: *Sporobolus cryptandrus*



- Lower genetic diversity among in G0 and G1
- Wild genetically differentiated from G0/G1

Faske et al., *In review*

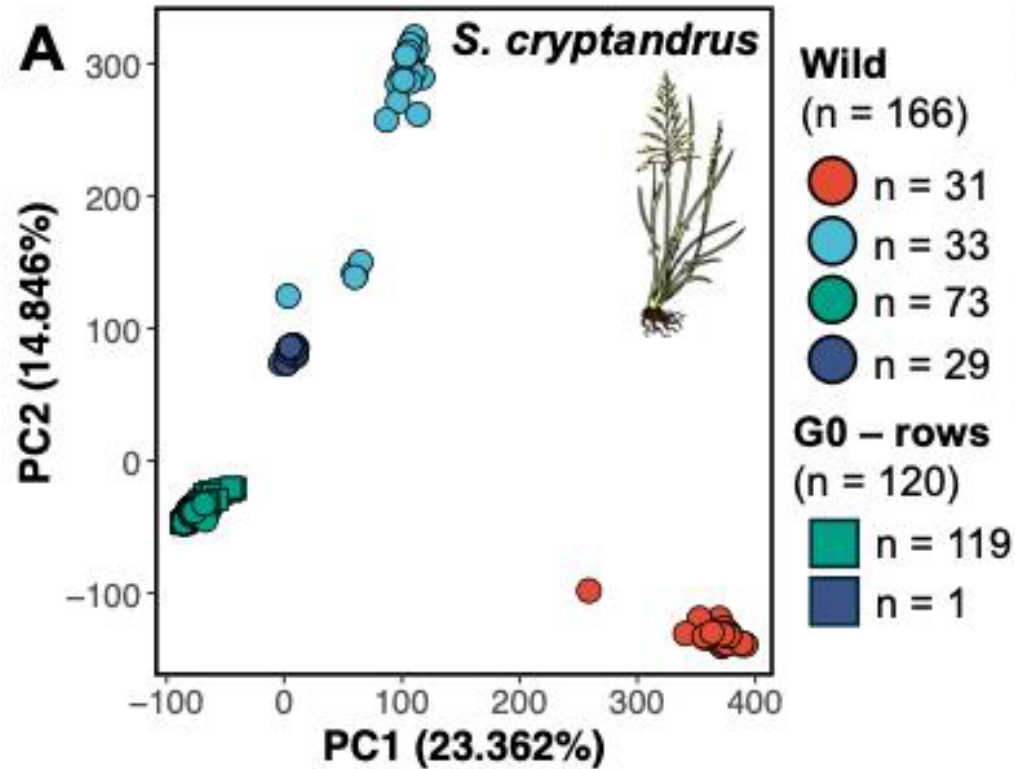
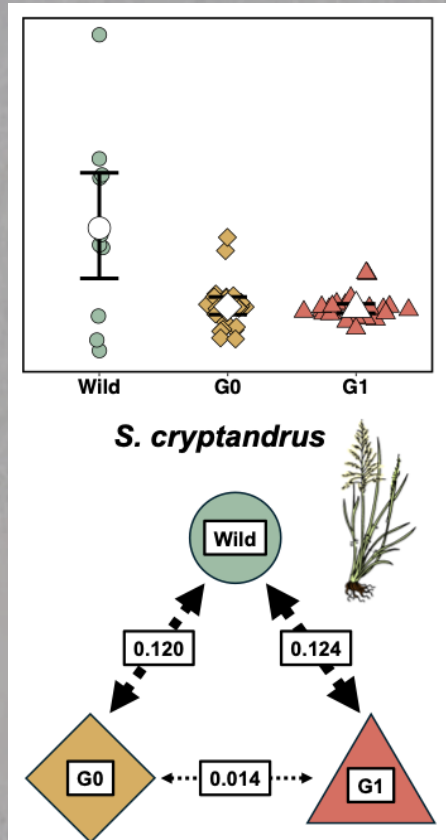
Results: *Sporobolus cryptandrus*



- 4 genetic lineages (Wild)

Colors represent assigned genetic cluster

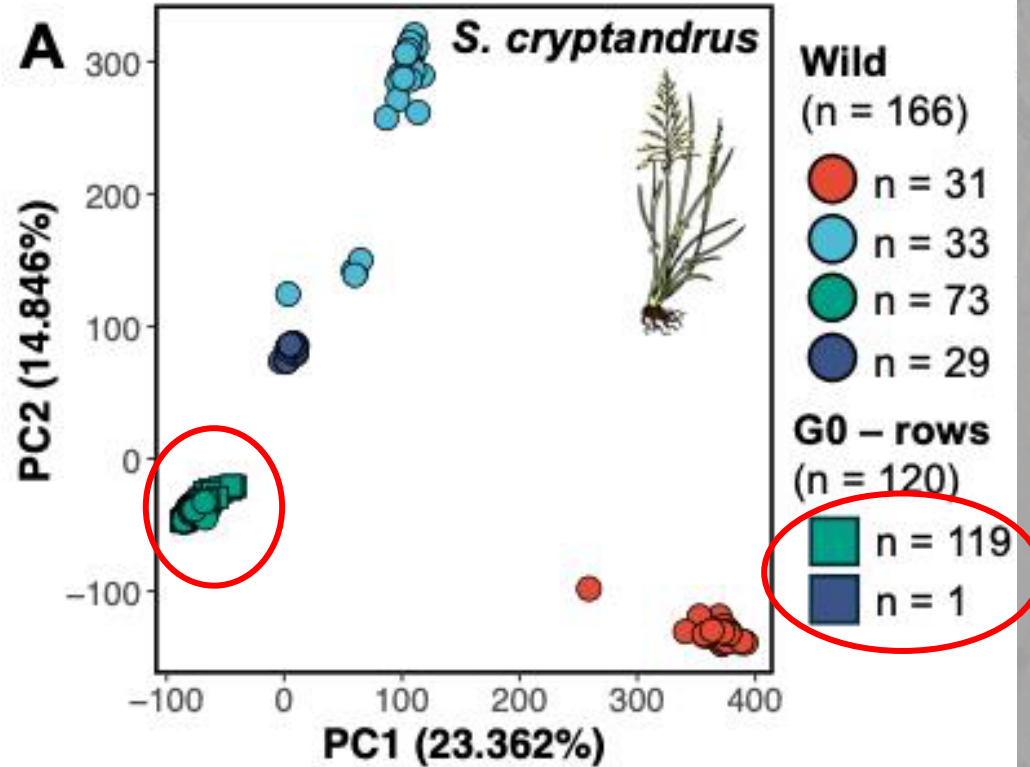
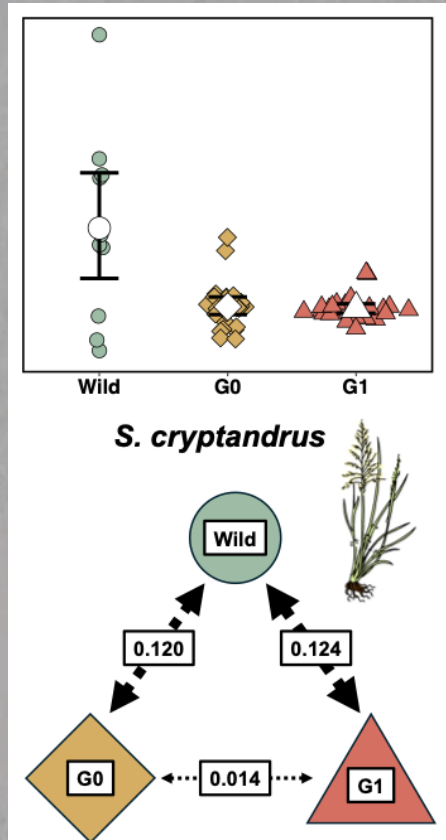
Results: *Sporobolus cryptandrus*



- 4 genetic lineages (Wild)
- All lineages can be present at each wild population

Colors represent assigned genetic cluster

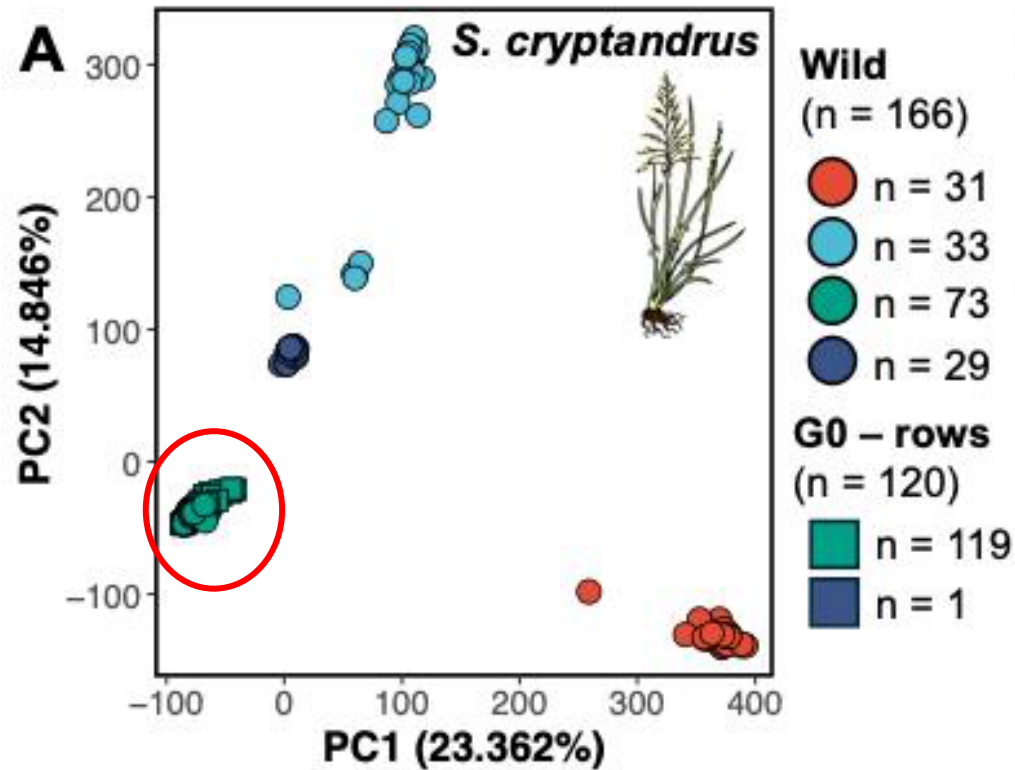
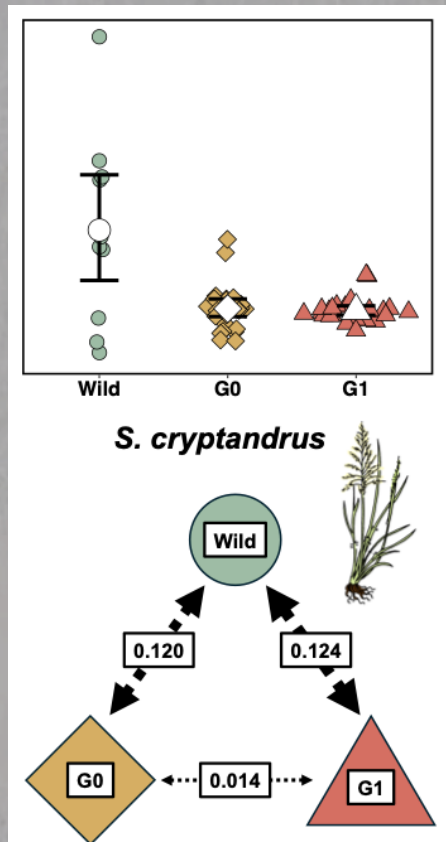
Results: *Sporobolus cryptandrus*



- 4 genetic lineages (Wild)
- All lineages can be present at each wild population
- Reduced to 2 lineages in production field (G0)

Colors represent assigned genetic cluster

Results: *Sporobolus cryptandrus*

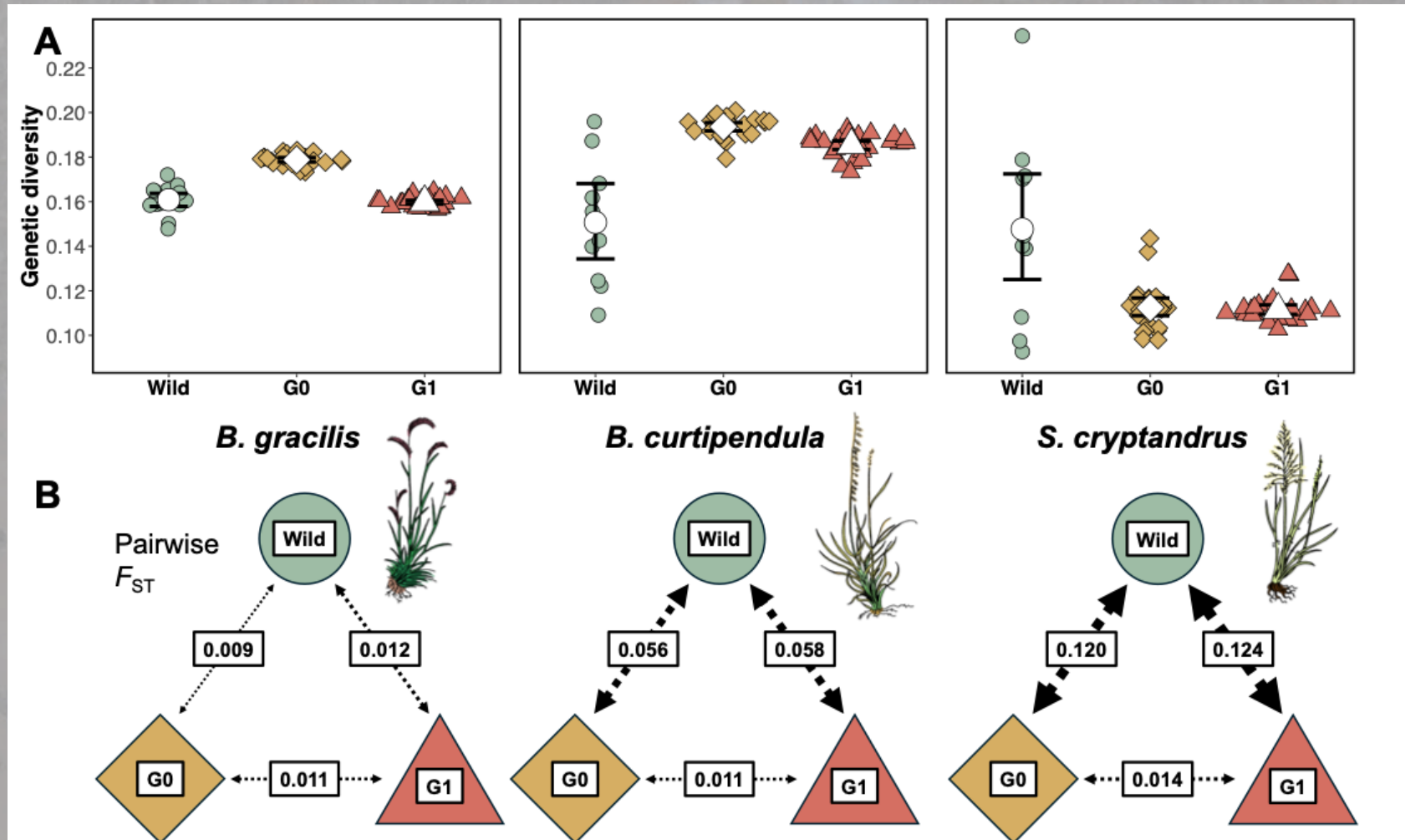


- 4 genetic lineages (Wild)
- All lineages can be present at each wild population
- Reduced to 2 lineages in production field (G0)

One genetic lineage dominates in agricultural fields, hence:

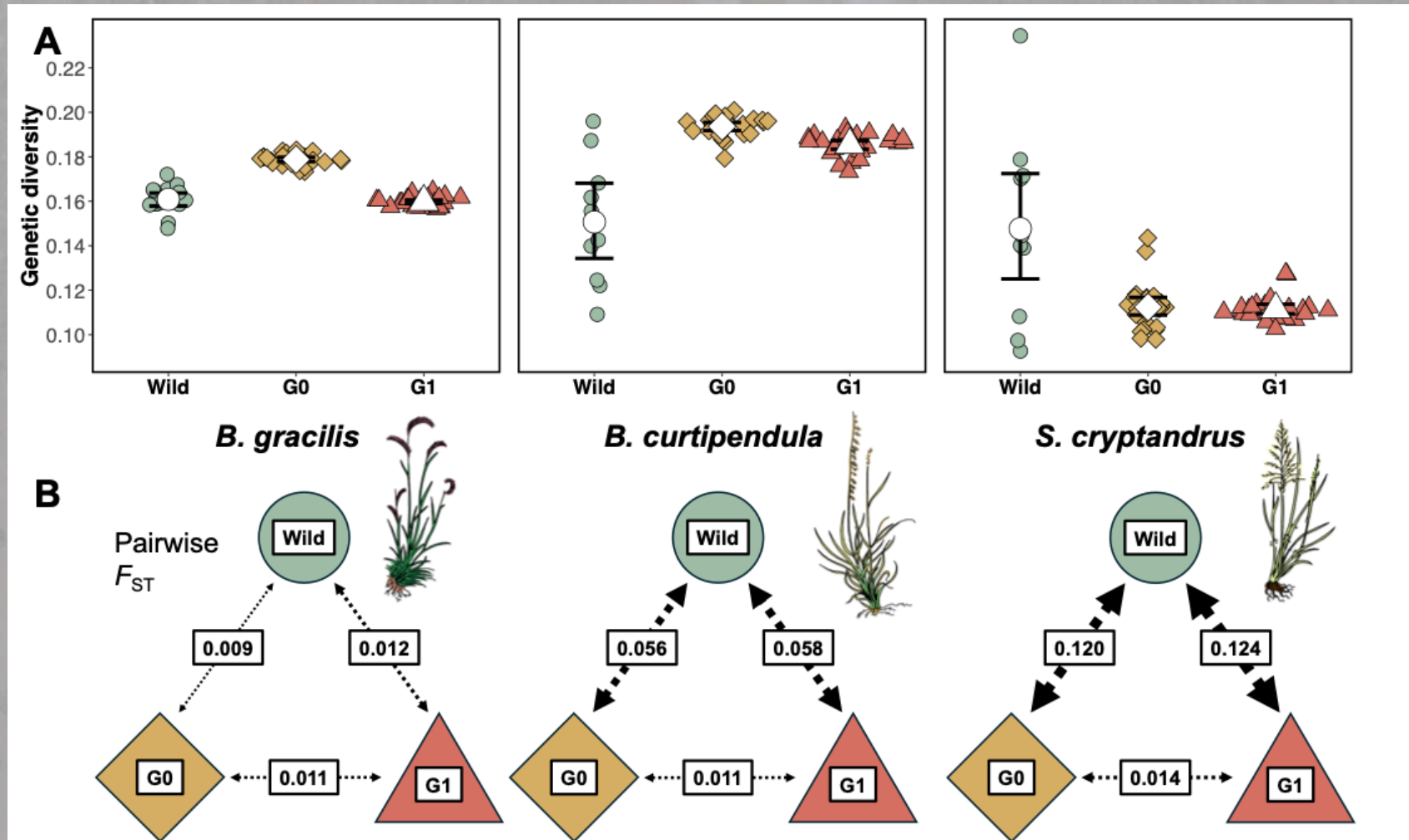
- lower genetic diversity
- higher genetic differentiation

Species-specific biology influences agricultural seed increase



Faske et al., *In review*

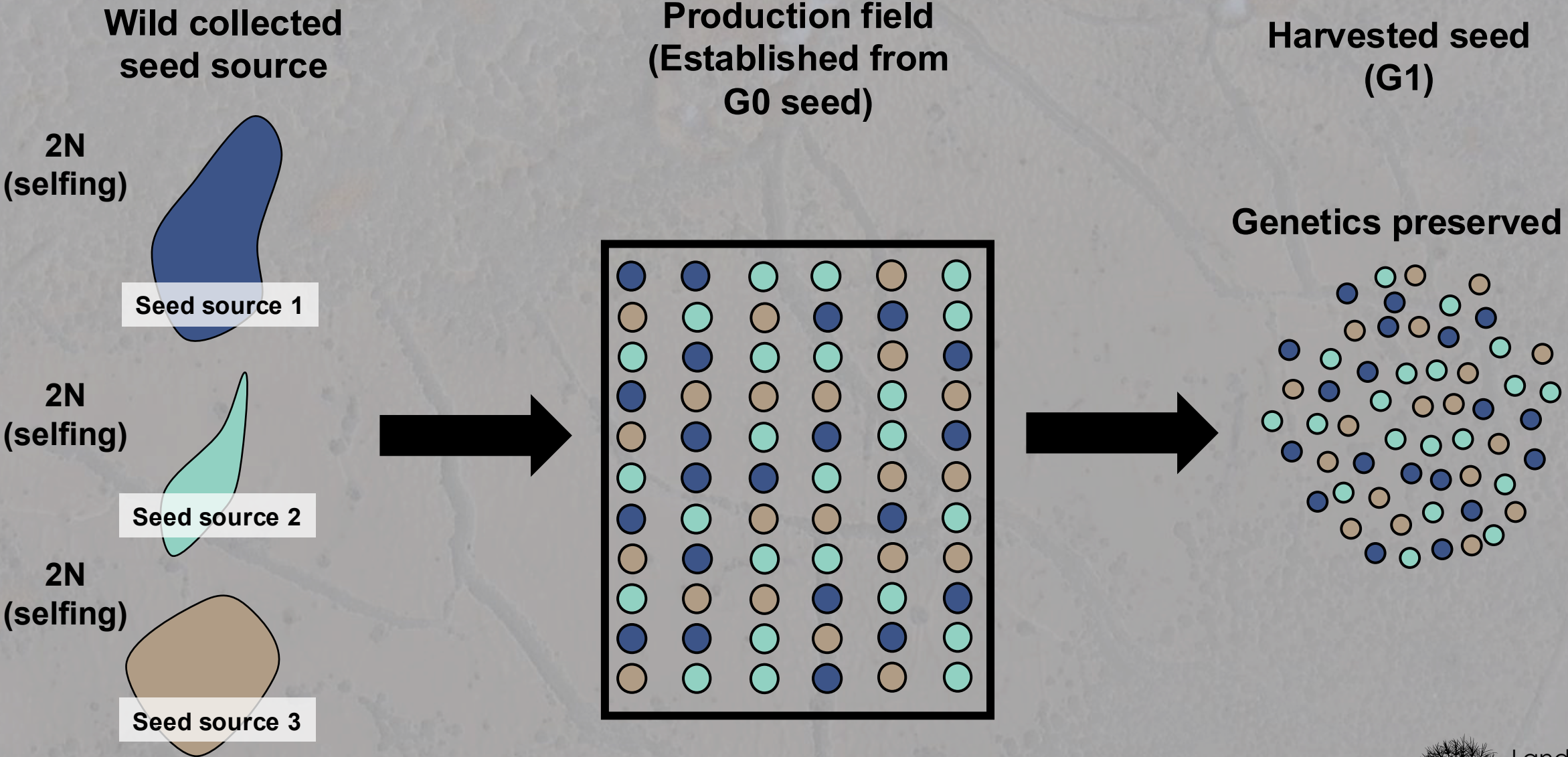
Species-specific biology influences agricultural seed increase



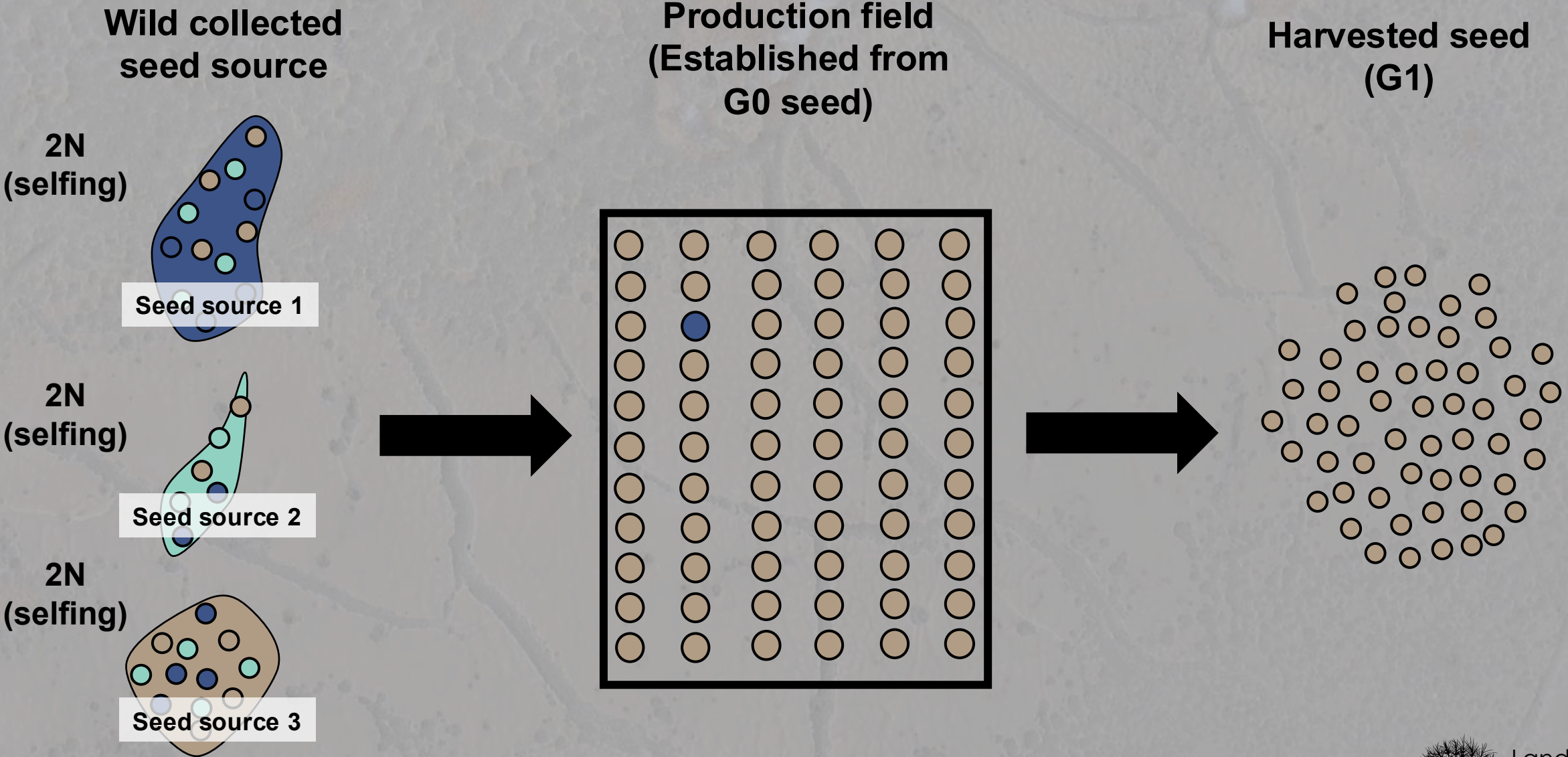
Faske et al., *In review*

Agricultural increase is not the culprit

Seed increase hypothesis for *S. cryptandrus*



Reality for *S. cryptandrus*



Where do the shifts occur (*Bouteloua curtipendula* and *Sporobolus cryptandrus*)?



Genetic shifts occur between seed collection and field establishment

- Factors to consider:
- timing of seed collection
 - single vs. multiple collection events
 - seed storage
 - greenhouse propagation of plugs
 - and more...

Take home messages

In our quest to increase the number of species and sources available to conduct ecologically and genetically appropriate restoration:

- Species biology matters
- We need more information about how seed collection and field establishment practices cause genetic shifts
- Does it matter?
 - Do we need all genetic lineages and ploidy levels present?



Take home messages

In our quest to increase the number of species and sources available to conduct ecologically and genetically appropriate restoration:

- Species biology matters
- We need more information about how seed collection and field establishment practices cause genetic shifts
- Does it matter?
 - Do we need all genetic lineages and ploidy levels present?

Using native seed in a project? Do you want to help understand these patterns and processes better? Contact us!



Many people supported this work:

John Bradford Melanie Gisler
Adrienne Pilmanis Gwen Wion
Zoe Davidson Laura Shriver
Maria Mullins
Carla Roybal
Shannon Lencioni

Funding support:
USDA-NIFA-AFRI grant #2022-67013-36116
New Mexico BLM L23PG00055

Reach out!

Rob Massatti: rob@landscollective.org
Trevor Faske: trevor@landscollective.org

Learn more & Contribute:
www.landscollective.org

