# Plant breeding through the lens of quantitative genetics

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#### **About me**



- 1. Malaysia
- 2. Indiana (BSc Biotech, Maths)
- 3. Wisconsin (PhD Genetics)
- 4. Freising, DE (Postdoc)
- 5. Edinburgh, UK (Postdoc)

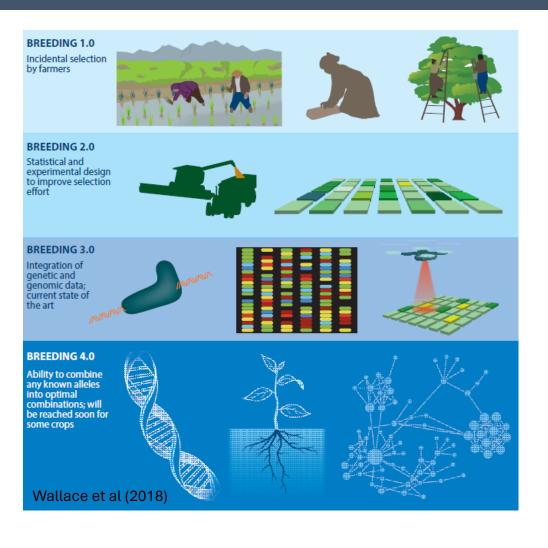
#### Talk outline

- Introduction
- Project 1: Crop domestication
- Project 2: Genetic diversity and pre-breeding
- Project 3: Novel crop breeding
- Future directions

#### Introduction

Plant breeding, complex traits and genetic gain

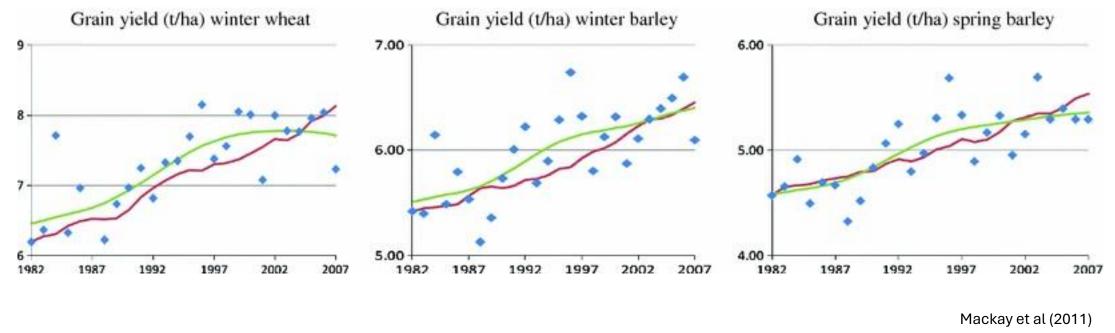
## **Plant Breeding**



- Domestication
- Improvement
- Experimental design
- Marker assisted selection (MAS)
- Genomic/Phenomic selection (GS/PS)
- Biology-Breeding
- Gene editing (GE)
- Functional variants
- Machine learning (ML/AI)

## **Breeding in major crops**

Yield increase is largely driven by genetic improvement.

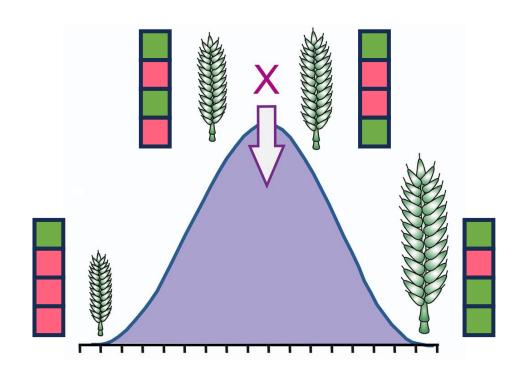


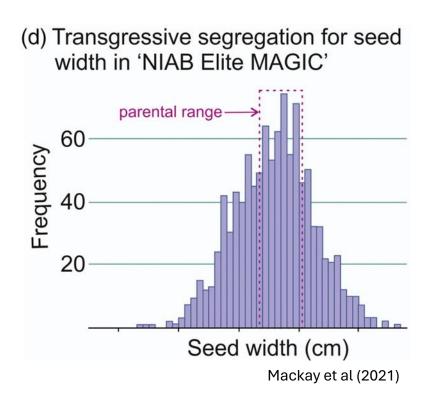
Green/blue: national yield

Red: variety effect

# Success in line breeding

Transgressive segregation: recombination and shuffling of causative genetic loci.



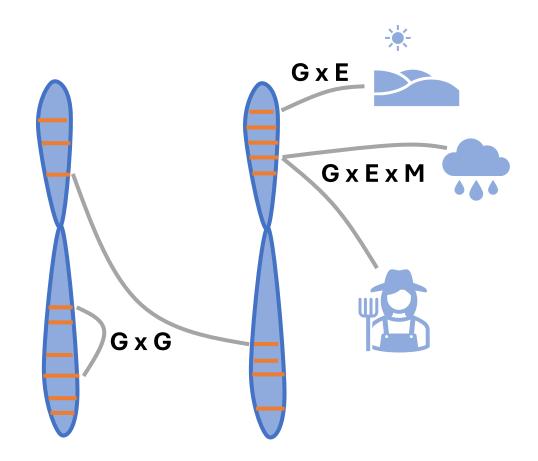


## **Complex traits**

Breeding targets: yield, flowering time, plant architecture, resource use efficiencies.

Quantitative/polygenic traits

Working model:



#### **Genetic Gain**

Breeder's equation (Lush 1937)

$$R = h^2 S$$

$$R = \frac{\sigma_g^2}{\sigma_p^2} \sigma_p i$$

$$R = \frac{\sigma_g}{\sigma_p} \, \sigma_g \, i$$

$$R = h\sigma_g i$$

Rate of genetic gain

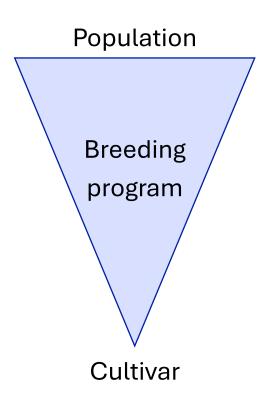
$$\Delta R = \frac{h\sigma_g i}{t}$$

$$\Delta R = \frac{sel.accuracy \times genetic\ variation \times sel.intensity}{time}$$

Framework for quantitative genetics

# Improving $\Delta R$

$$\Delta R = \frac{sel. accuracy \times genetic \ variation \times sel. intensity}{time}$$



- Multiparental population, mutation, pre-breeding.
- MAS, GS, GE, phenomics.
- Larger/efficient trials.
- Rapid cycling, speed breeding (SB/RGA).

# Plant breeding: a journey through time

Project 1: crop domestication

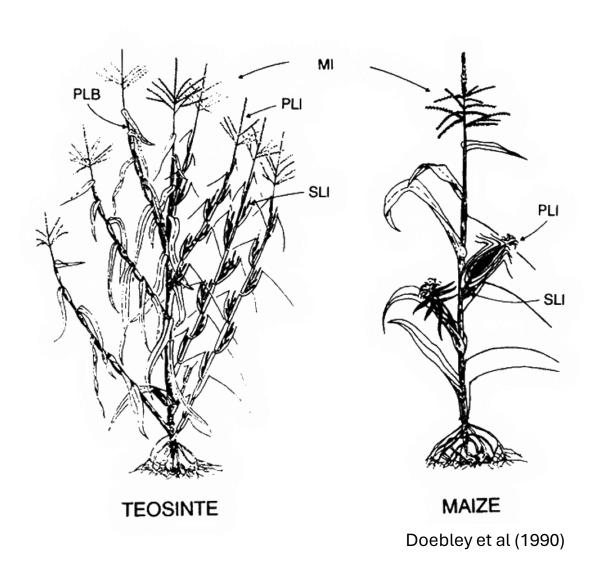
Project 2: genetic diversity and pre-breeding

Project 3: novel crop breeding

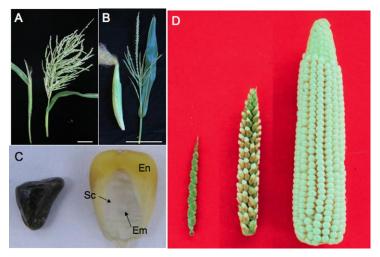
#### **Domestication**

This earliest form of <u>plant breeding</u> is known as <u>domestication</u>, where plants were selected to be more productive, easier to harvest, or more aesthetically or gastronomically pleasing (Flint-Garcia 2013).

#### Maize-teosinte model



- Tillers
- Lateral branches
- Terminal inflorescence
- Ear length/diameter/number
- Kernel number/size/glume



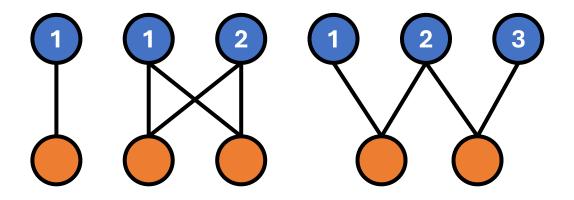
Hake and Ross-Ibarra (2015)

# Study population

#### 50 parents from Palmar Chico – Rio Balsas



#### **Crossing schemes**



#### Field trials

- Homestead, FL, 2013 2017
- 3,000 plants/year, randomized
- 18 domestication traits
- GBS, WGS





#### Field trials

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# Variance/covariance partitioning

Fitted standard mixed linear model  $y = X\beta + a + d + ay + e$ 

Trait = Fixed + Additive + Dominance + Additive \* Year + Residual

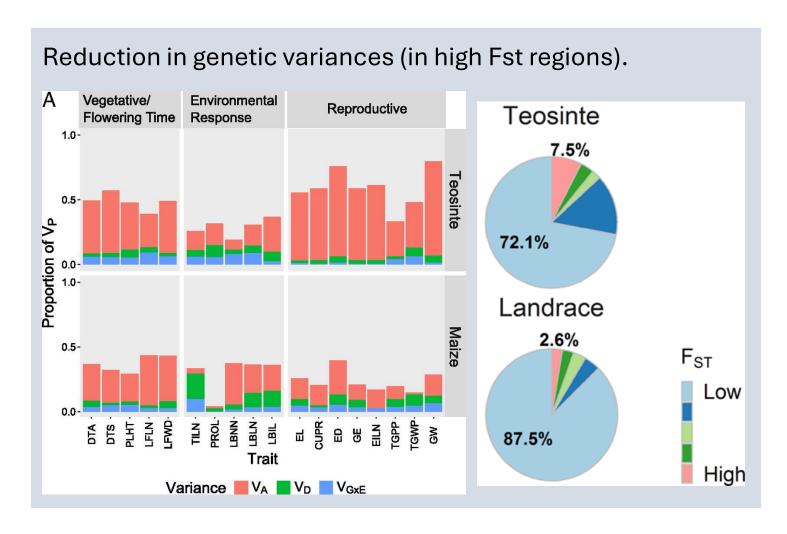
$$a \sim N(0, K_A \sigma_A^2)$$

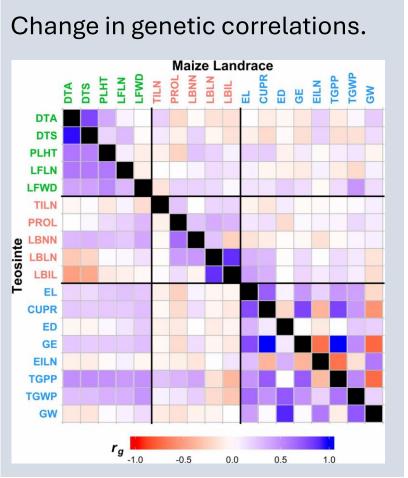
$$d \sim N(0, K_D \sigma_D^2)$$

$$ay \sim N(0, K_{AY}\sigma_{AY}^2)$$

Only the additive term was fitted in the bivariate model.

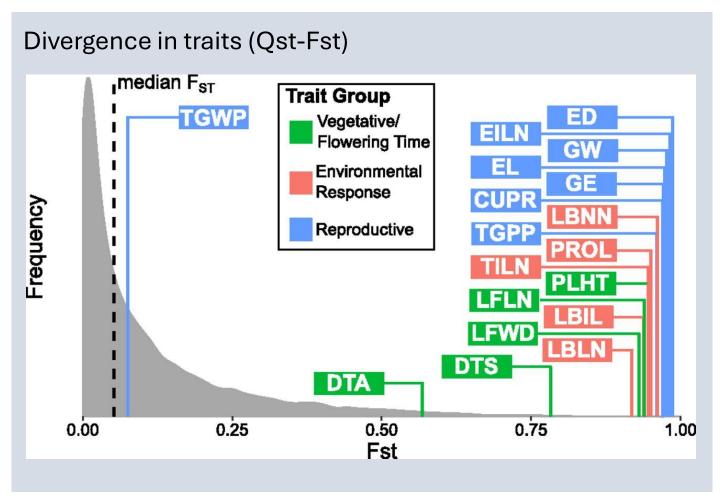
## QG modelling of domestication

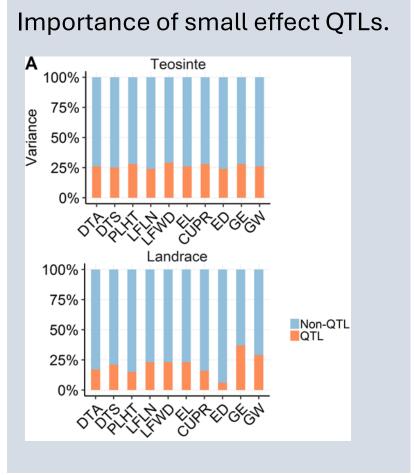




Yang et al (2019), Chen et al (2020, 2021)

## QG modelling of domestication





Yang et al (2019), Chen et al (2020, 2021)

# Plant breeding: a journey through time

Project 1: crop domestication

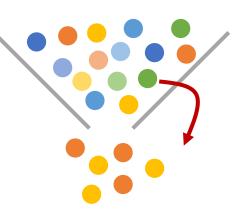
Project 2: genetic diversity and pre-breeding

Project 3: novel crop breeding

# Shuffling genetic diversity

#### Domestication/Improvement

- Bottleneck
- Selection



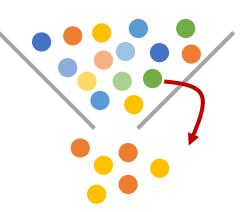
Marker assisted selection (MAS)

- Great for oligogenic traits.
- E.g. disease resistance.
- Inefficient for polygenic traits.

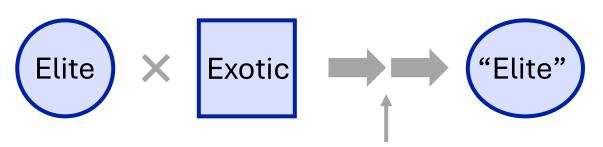
# Shuffling genetic diversity

#### **Domestication/Improvement**

- Bottleneck
- Selection



Polygenic introgression is challenging



Phenotypic/genomic selection

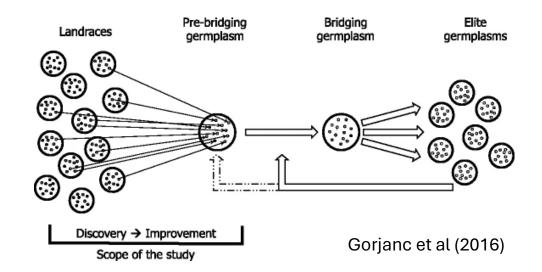
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#### **Pre-breeding**

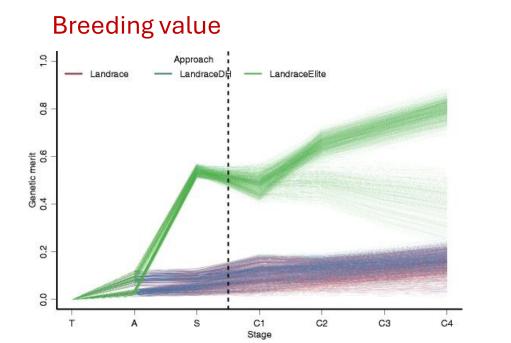


Seeds of Discovery (SeeD) studies and characterizes maize and wheat genetic diversity for use in breeding programs, which develop wheat varieties and maize hybrids improved through conventional technologies. These hybrids are better adapted to climate change, more resistant to pests and diseases and have higher yield potential.

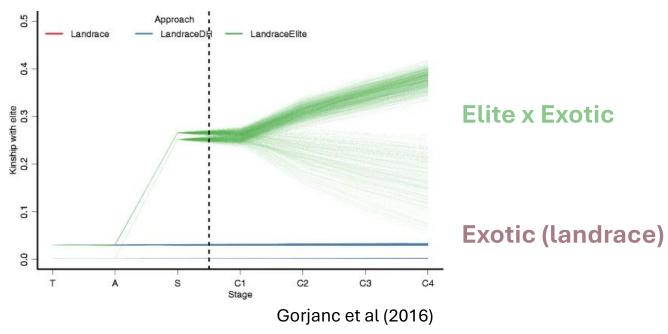


- Bridging population with elite-exotic crosses.
- Improve in elite-exotic first, then introduce into elite population.

# Selection bias in pre-breeding (simulation)



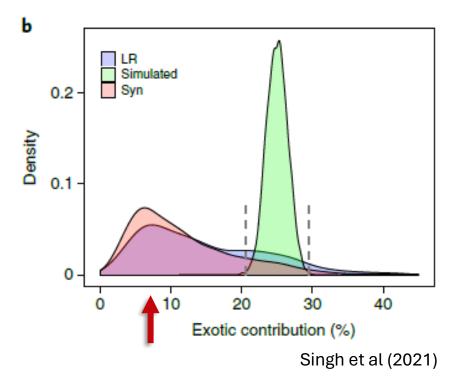
#### Kinship with elite parents



- Selection within exotic is slower than elite-exotic.
- Selection within elite-exotic reconstitutes the elite parent genome.

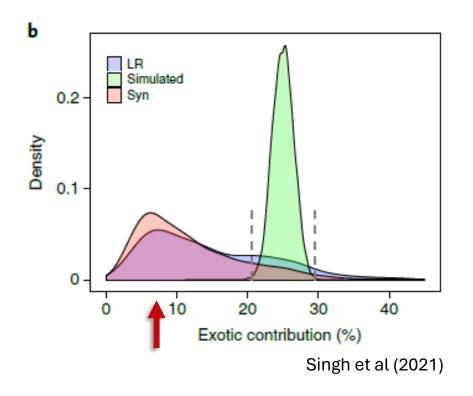
# Selection bias in pre-breeding (observed)

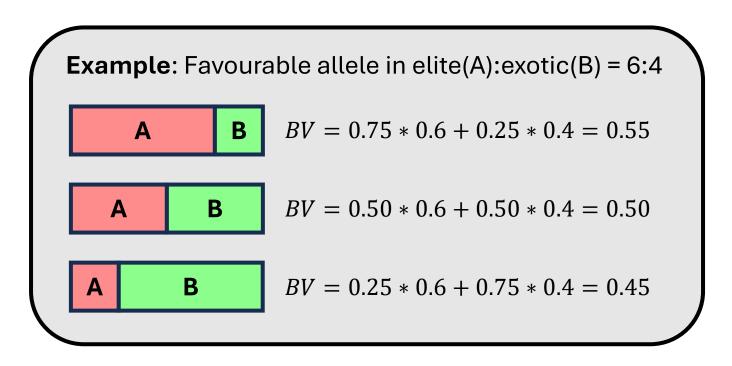
- Elite 2 x (Elite 1 x Exotic)
- Distribution of exotic parent genome after applying phenotypic selection.
- Validation of simulation outcomes in experimental observation.



# Selection bias in pre-breeding (observed)

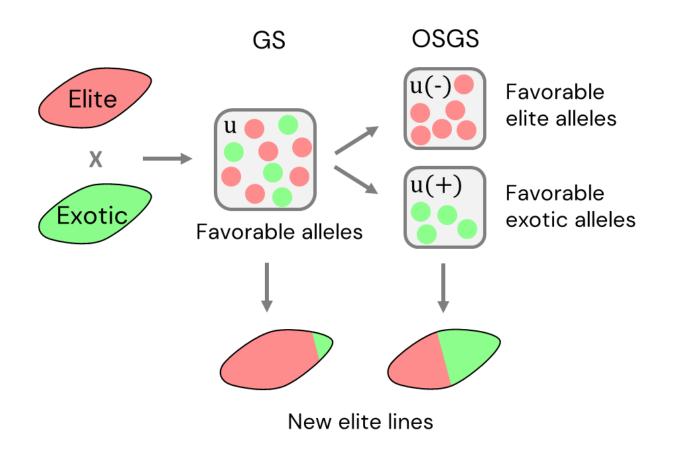
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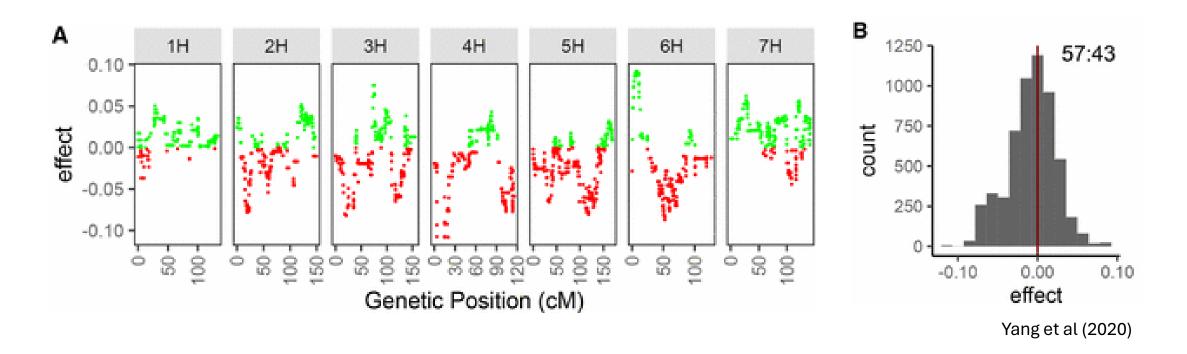
#### Origin specific genomic selection (OSGS)

OSGS: isolate and select on favorable parental contribution.

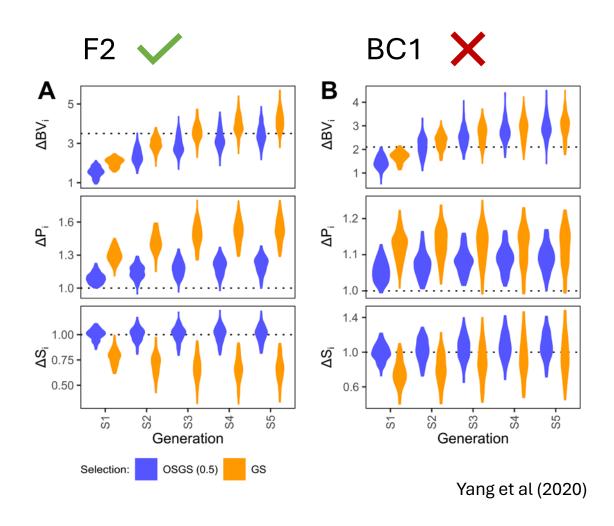


# OSGS in barley NAM (yield)

Partitioning favorable parental contributions.



#### OSGS in simulated data



- OSGS vs GS.
- Weighted selection in OSGS.
- Comparable breeding values.
- OSGS maintains elite-exotic balance.
- F2 > BC1.

# Plant breeding: a journey through time

Project 1: crop domestication

Project 2: genetic diversity and pre-breeding

Project 3: novel crop breeding

# Novel crop – purslane (Portulaca oleracea)



## Developing a breeding program for purslane

- Env: Vertical farm
- Trait: Omega-3 level

TABLE 2: Plant sources of omega-3 fatty acids (g/100 g).

Category	Fruits/vegetables	Amount (g)
Low	Avocados, California raw	0.1
	Broccoli	0.1
	Strawberries	0.1
	Cauliflower, raw	0.1
	Kale, raw	0.2
	Spinach, raw	0.1
	Peas, garden dry	0.2
	Cowpeas, dry	0.3
	Beans, navy, sprouted, cooked	0.3
	Corn, germ	0.3
Medium	Bean, common dry	0.6
	Leeks, freeze-dried, raw	0.7
	Wheat, germ	0.7
	Spirulina, dried	0.8
	Purslane	0.9
	Oat, germ	1.4
	Beachnuts	1.7
	Soybeans kernels, roasted	1.5
	Soybeans, green	3.2

Uddin et al (2014)

Identify breeding targets.

Survey variation in phenotypes, GxExM.

Engage with stakeholders.

Register varieties.

Create populations and select.

Trial in vertical farms.

#### Short vs long day



Fluorescent vs LED





# Developing a breeding program for purslane

#### Royal Highland Show





Identify breeding targets.

Survey variation in phenotypes, GxExM.

Engage with stakeholders.

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Trial in vertical farms.

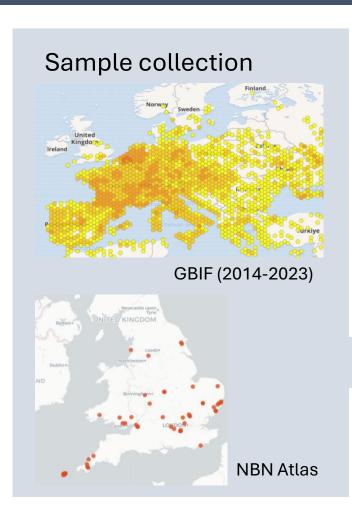
#### Green vs golden purslane







## Developing a breeding program for purslane



Identify breeding targets.

Survey variation in phenotypes, GxExM.

Engage with stakeholders.

Register varieties.

Create populations and select.

Trial in vertical farms.

#### Growth chambers



Hydroponic trials

VF – under construction



What's next?

RUE: resource use efficiency



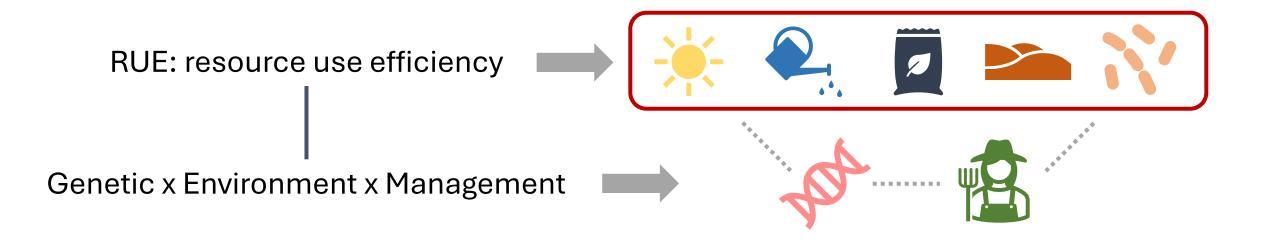


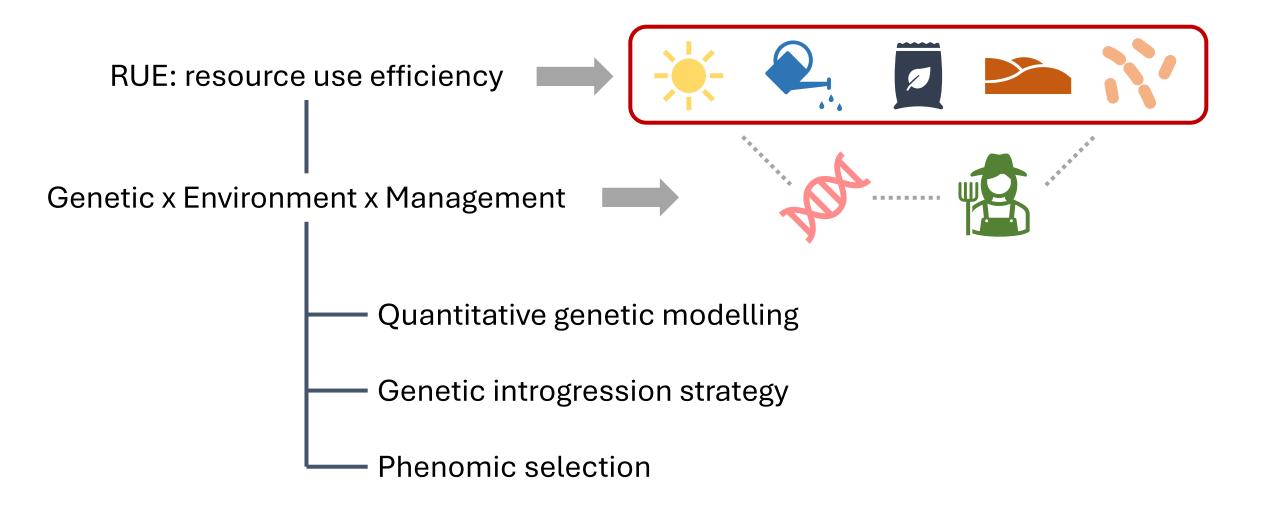












#### Summary

- Plant breeding, complex traits and genetic gain.
- Plant breeding: a journey through time:
  - QG-perspective on maize domestication
  - Genetic introgression in pre-breeding
  - Breeding for novel crop
- Interest in RUE and GEM.



- Breeding efficiency
- Sustainability
- Climate resilience

#### Acknowledgement

#### Many thanks for the opportunity to present today!

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Wayne Powell

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David Marshall

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