

Introduction

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Breeder Equation

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Accuracy

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Blueberry

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Final Consideration

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(HOS 6932)– Survey of Breeding Tools and Methods

Theoretical and Practical Aspects for Genomic Selection Implementation

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Introduction

In the last class ...

- Importance of training and testing data set
- Metrics to measure the predictive ability
- RR-BLUP vs. GBLUP

In this class ...

- Why and how to switch to genomic selection?
- Example in blueberry
- Final discussion

Introduction

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Prediction of Total Genetic Value Using Genome-Wide Dense Marker Maps

T. H. E. Meuwissen,* B. J. Hayes[†] and M. E. Goddard^{‡,§}

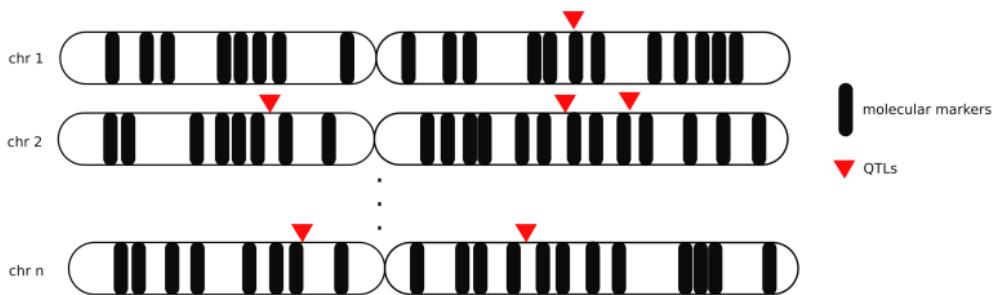
*Research Institute of Animal Science and Health, 8200 AB Lelystad, The Netherlands, [†]Victorian Institute of Animal Science, Attwood 3049, Victoria, Australia and [‡]Institute of Land and Food Resources, University of Melbourne, Parkville 3052, Victoria, Australia

Manuscript received August 17, 2000
Accepted for publication January 17, 2001



Introduction

General Idea



$$\boxed{\text{GEBV}} = \boxed{\text{DNA}} \boxed{a} + \boxed{e}$$

↓ ↓

Genomic Estimated
Breeding Values

$$\boxed{\text{GEBV}} = \boxed{\text{Plant}} \boxed{u} + \boxed{e}$$

↓ ↓

$\boxed{\text{DNA}} \rightarrow \boxed{\text{Pedigree}}$

Relationship Matrix

Introduction

Background

- After this pioneer study genomic prediction has popularized in plant and animal breeding
- Organization of breeding programs can be very different between crops:
 - ▷ Clonally propagated species
 - ▷ Perennial crop
 - ▷ Genomic Selection in Hybrid Breeding
 - ▷ Self-pollinated crops
 - ▷ Family structures
- It is a dynamic process: implementation can change over time according to the evolution of technologies, knowledge, and breeding goals.

Breeder's Equation

Central Question

Why and how to switch to genomic selection?

Simple answer: GS can impact all terms in the Breeder's equation:

$$\Delta G = \frac{i \times r \times \sigma_g^2}{L}$$

- ΔG = genetic gain per cycle
- i = intensity of selection
- r = accuracy of selection
- L = breeding cycle time

Discuss element-by-element in the Breeder's equation

Breeder's Equation

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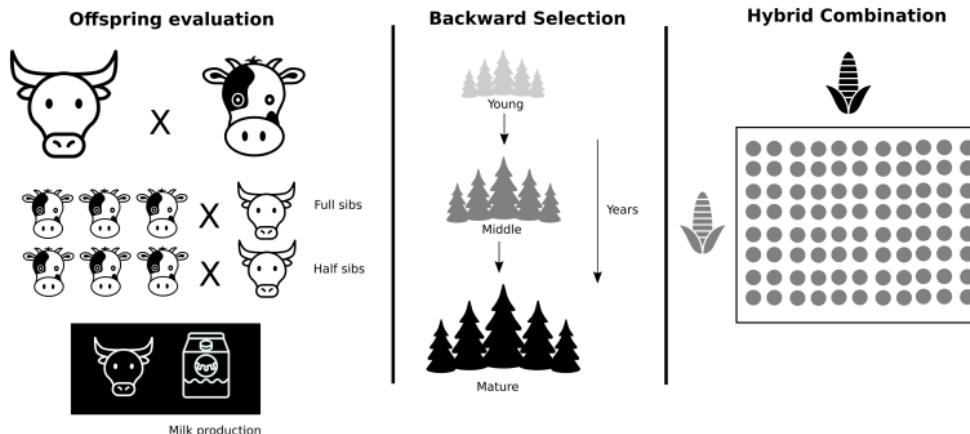
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Breeder's Equation

Breeding cycle time

- Breeding cycle is constrained by the way in which phenotype is collected
- ↑ Genetic Gain by ↓ breeding cycle

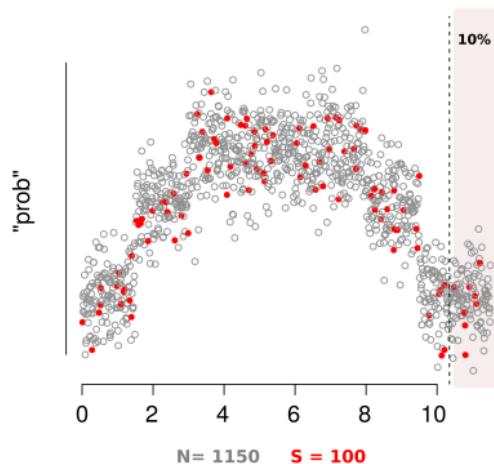


Breeder's Equation

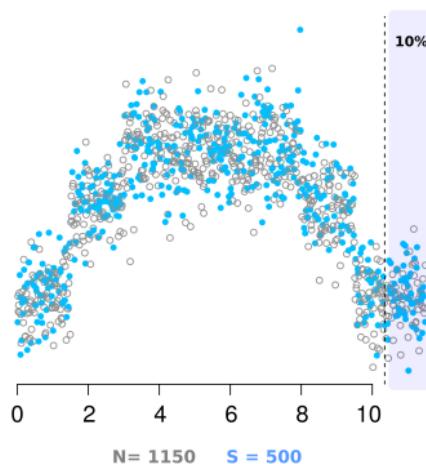
Intensity

- Lower costs with the same or higher testing capacity
- ↑ Genetic Gain by ↑ intensity

Phenotypic Selection



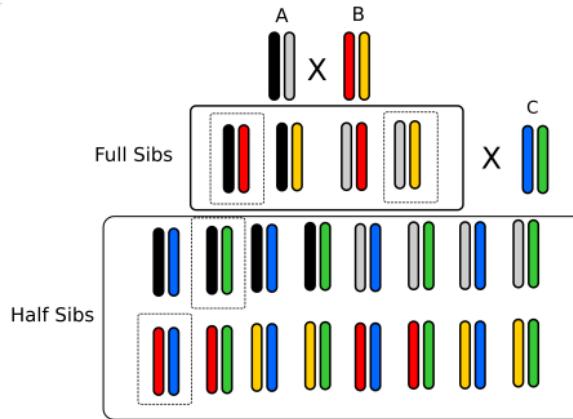
Genomic Selection



Breeder's Equation

Accuracy

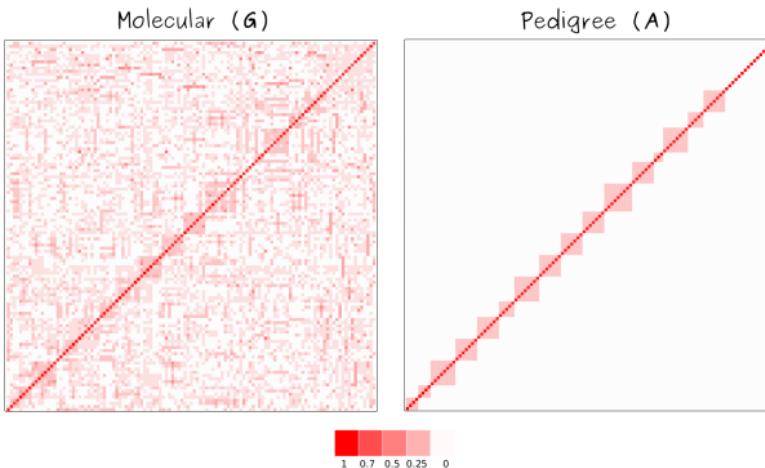
- Without genomic, full sibs have the same estimated breeding values
- Some genotypes can more related to the other, due to the inheritance of random samples of alleles from both parents.
- Mendelian Sampling variance
- ↑ Genetic Gain by ↑ accuracy



Breeder's Equation

Genetic Diversity

- Best cross: combination between highest BV and diversity
- Genomic relationship are more precise than realized kinship
- ↑ Genetic Gain by ↑ diversity



Dealing with accuracy

Background

- Predictive accuracy is a key metric for successful of genomic selection
- There are many deterministic formulas
- Check the hands-on

Expected accuracy

$$r = \sqrt{N_p h^2 [N_p h^2 + M_e]^{-1}}$$

- Daetwyler et al. (2013)
- N_p is the number of individuals in the TRN pop
- h^2 is the heritability of the trait
- M_e is the number of independent chromosome segments

Dealing with accuracy

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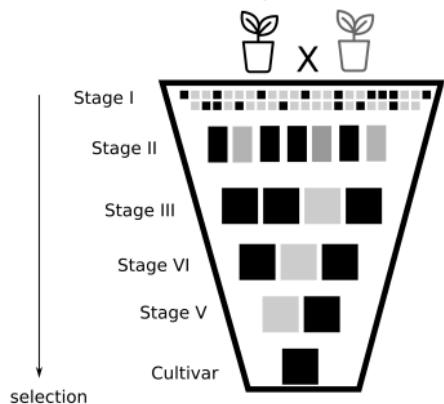
Expected accuracy

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Dealing with accuracy

Simulation Example:



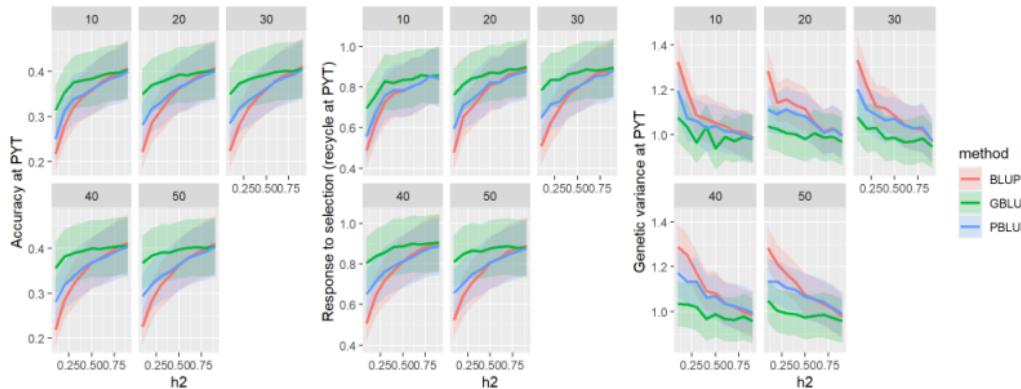
- Simulated 1000 materials in a clonal species
- Excellence in Breeding Platform
- Traits with different h^2
- Different family structures
- Selection based in GBLUP, PBLUP and BLUP

CGIAR – YouTube Channel – Basics of Genomic Selection

Dealing with accuracy

Simulation

Heritability vs. Population Structure vs. Accuracy

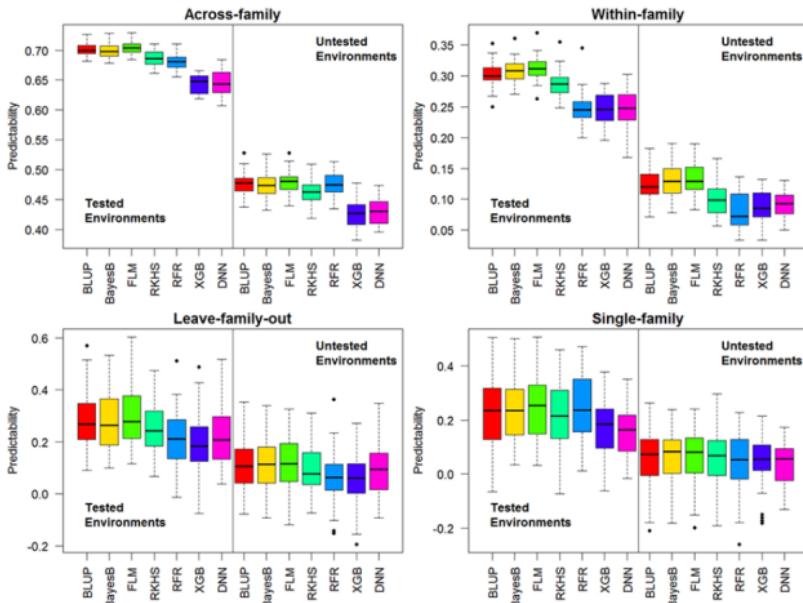


- GBLUP is the best for low-to-middle h^2
- With $\uparrow h^2$, we have \downarrow differences between ABLUP, BLUP or GBLUP.
- Lower gains in small families (large M_e)
- GBLUP tends to reduce faster the genetic variability

Dealing with accuracy

Soybean

Population Structure vs. Accuracy

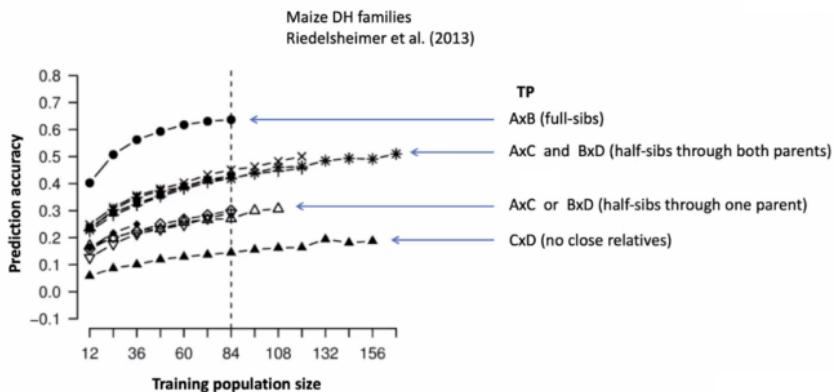


Xavier, 2021. <http://dx.doi.org/10.1590/1984-70332021v21Sa15>

Dealing with accuracy

Maize

Predicting progeny of A x B with different training population set

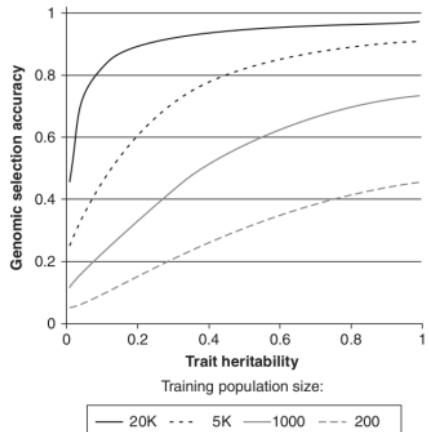


Riedelsheimer, 2013. <https://doi.org/10.1534/genetics.113.150227>

Dealing with accuracy

Barley

Heritability vs. Population Size vs. Accuracy



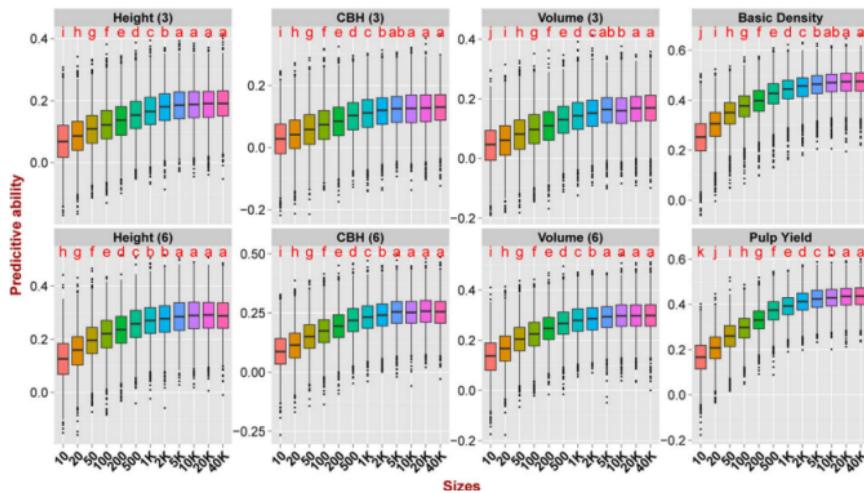
Predicted accuracy of genomic selection assuming the characteristic of elite barley:: effective population size of 50 and genetic map length of 1100 cM.

Hayes et al., 2009. DOI: 10.1017/S0016672308009981

Dealing with accuracy

Eucalyptus

SNP density vs. Accuracy

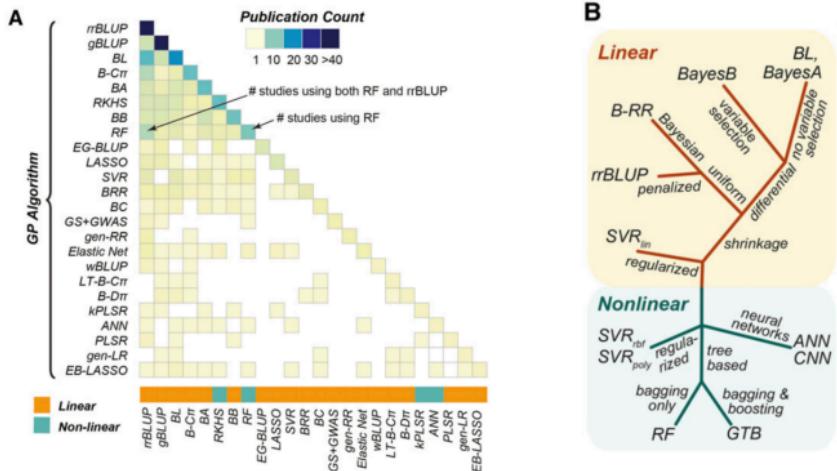


- Depends of LD and h^2 of the trait
- Most of the case, we can see a plateau

Tan et al., 2017. DOI 10.1186/s12870-017-1059-6

Predictive Models

Genomic Selection Models



- Different genetics and biological assumptions
- rrBLUP and GBLUP are the most popular

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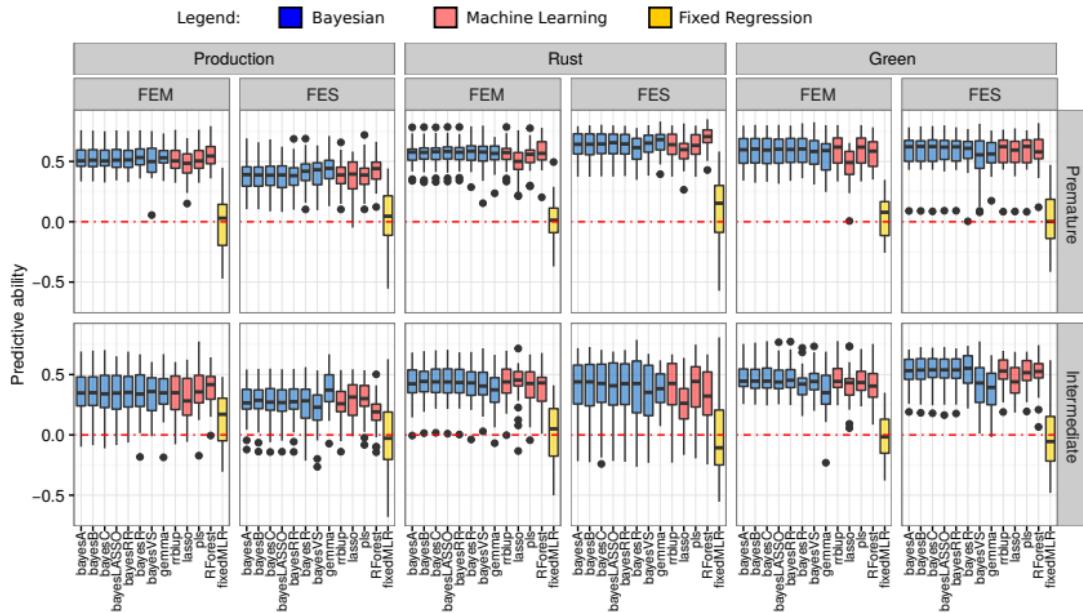
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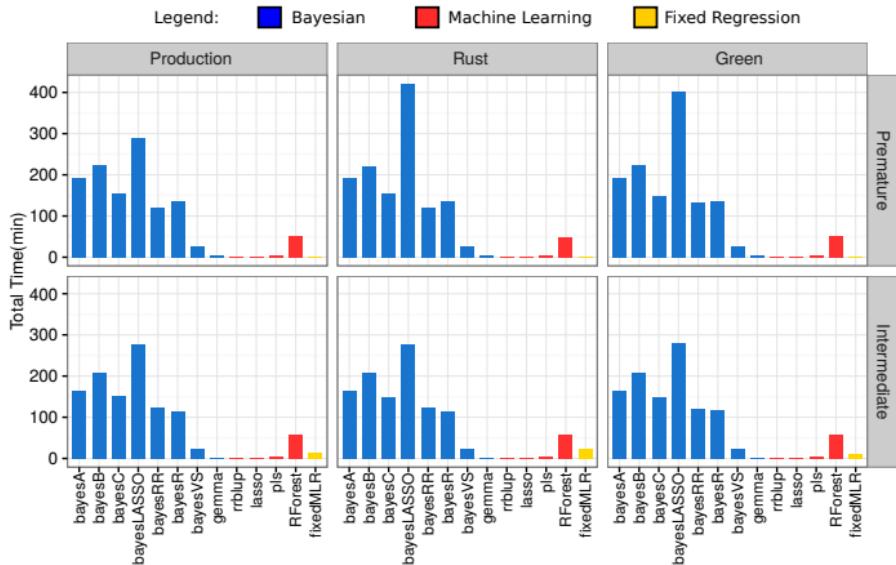
Predictive Models

Coffee example



Predictive Models

Coffee example



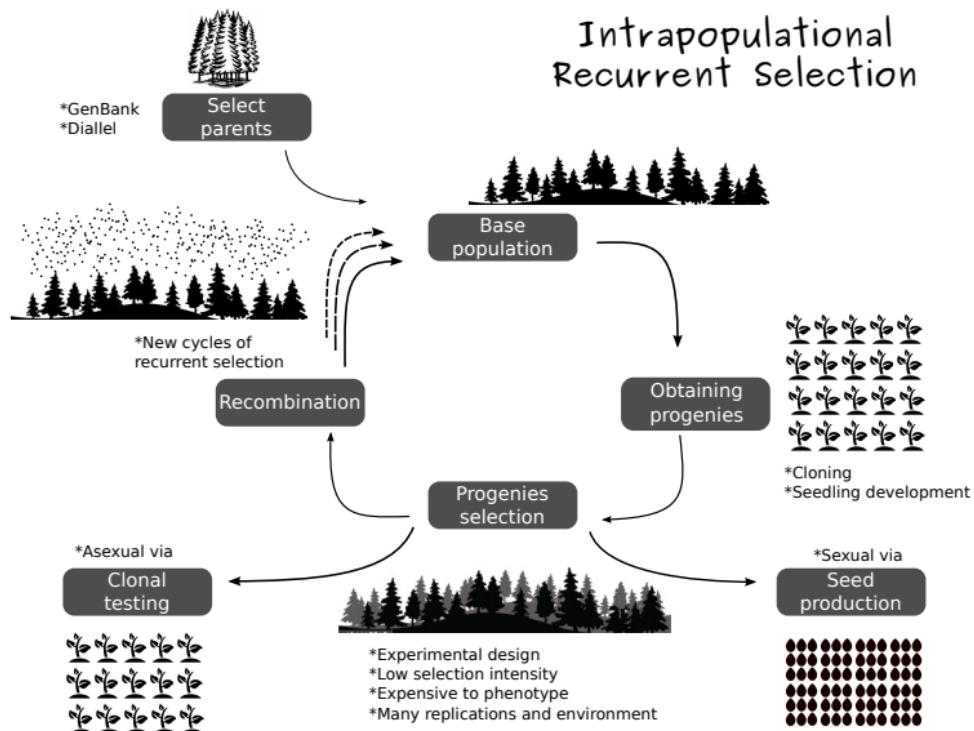
Genomic Selection in Blueberry

Background

- Blueberry is an autotetraploid species ($2n = 4x = 48$) species
- Perennial with a long juvenile phase
- Multi-year evaluations and large experimental areas
- High sensibility to inbreeding depression
- Breeding program most based in phenotypic recurrent selection



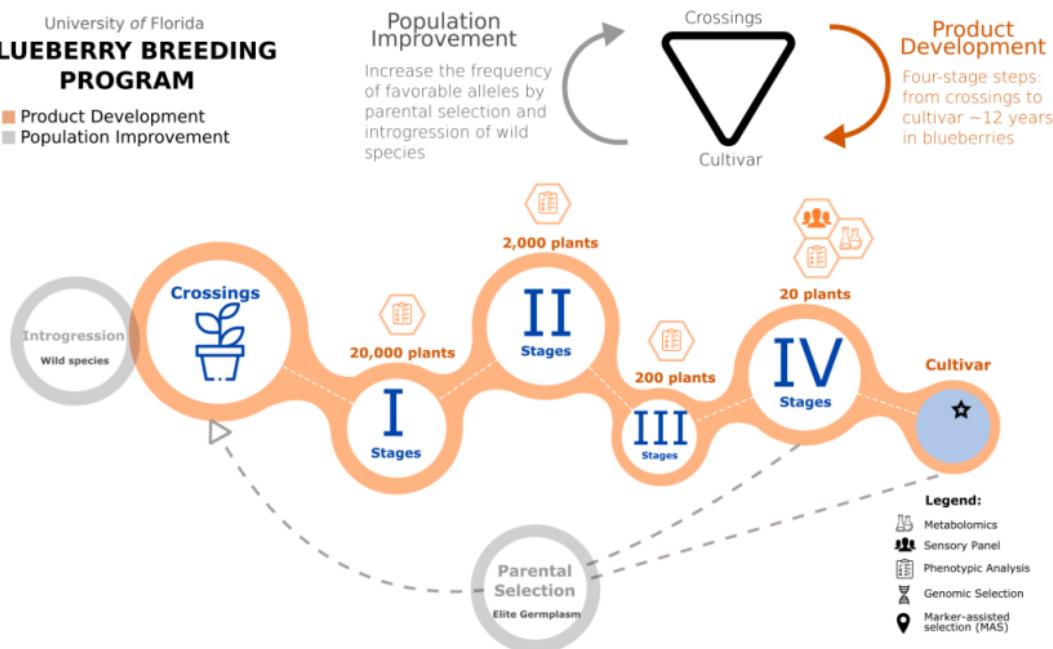
Genomic Selection in Blueberry



Genomic Selection in Blueberry

University of Florida
BLUEBERRY BREEDING PROGRAM

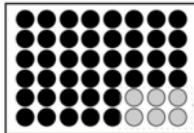
■ Product Development
 ■ Population Improvement



Genomic Selection in Blueberry

Calibration set

- Genotypes in earlier stages
- Genotypes in earlier and advanced stages



- * Diverse population
- * 1,837 genotypes
- * 2014 and 2015 seasons

① Cross-validation | Ratio (6x,60x)
Dosage (6x,60x)

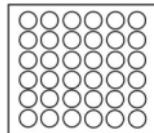
② Across-stages Scenario



- * Advanced stage
- * 114 genotypes
- * 2019 season

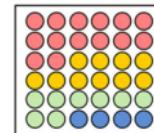
Testing set

③ General Scenario

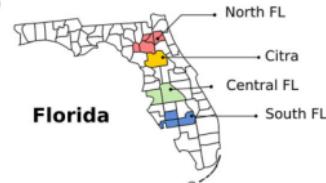


- * Advanced stage
- * 280 genotypes
- * Phenotypes collected from 2014 to 2020
- * Phenotypes corrected for year, location and age effects

④ Stratified Scenario



- * Advanced stage
- * 280 genotypes
- * Phenotypes collected from 2014 to 2020
- * Different FL regions
- * Phenotypes corrected for year and age effects



Florida

Phenotypes evaluated over four regions in the Florida State

Addressing different questions using different CV schemes:

- Scenario 1: cross-validation using the same population
- Scenario 2: same phenotypes evaluated in early and advanced stages
- Scenario 3: prediction the general performance in commercial conditions
- Scenario 4: prediction across environments

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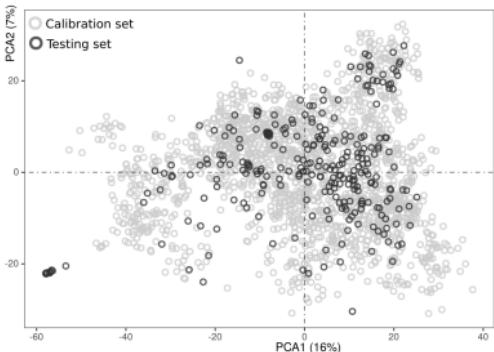
Blueberry
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Final Consideration
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(A)

Principal Component Analysis

Calibration vs. Testing Set



- * Testing set: advanced stages
- * Calibration: early stages

Introduction
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Breeder Equation
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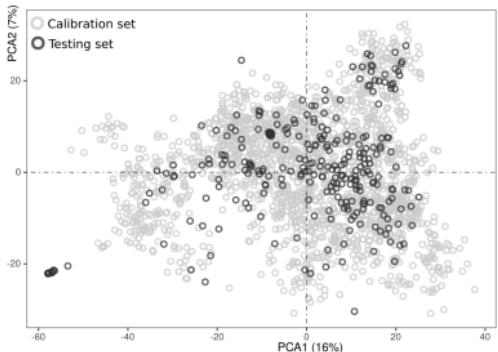
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Final Consideration
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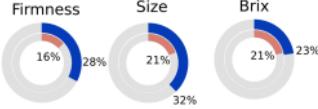
(A)

Principal Component Analysis Calibration vs. Testing Set



(B)

Across-stages Scenario



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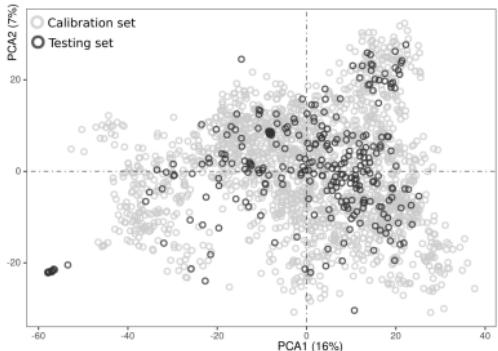
Breeder Equation
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Accuracy
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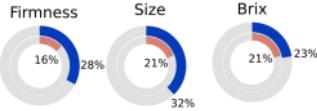
Blueberry
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Final Consideration
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(A) Principal Component Analysis
Calibration vs. Testing Set



(B) Across-stages Scenario



(C) General Scenario



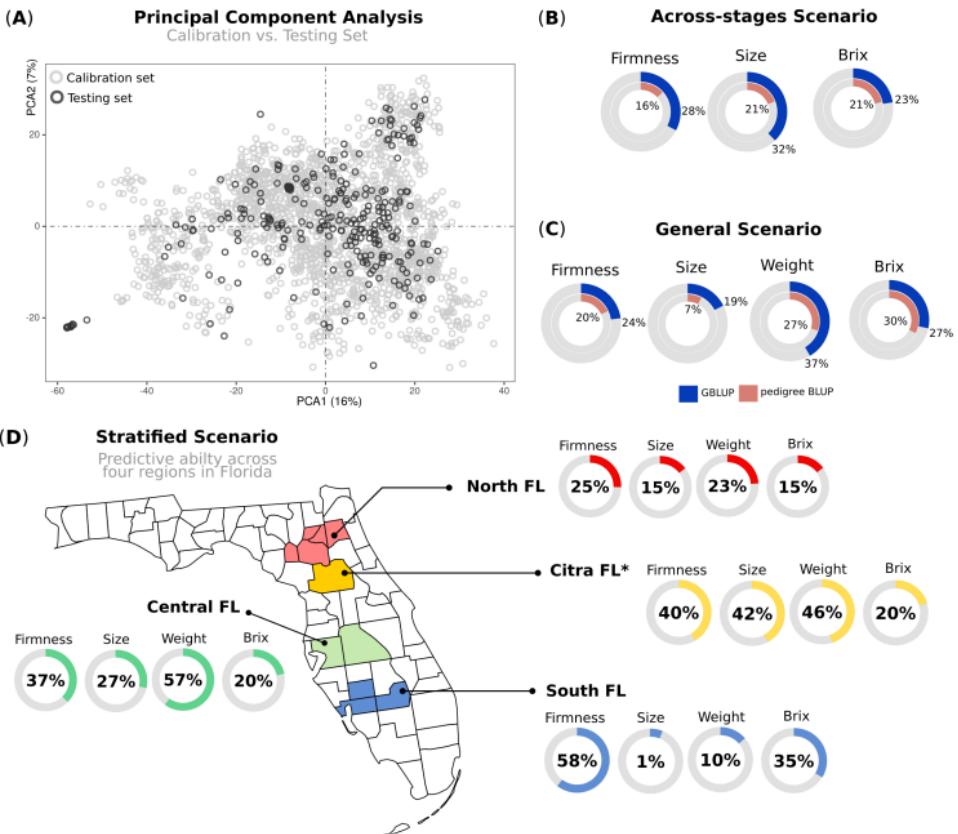
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Blueberry
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Final Consideration
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Final Consideration

My personal opinion for practical implementation:

- Genomic selection is a tool to assist breeding and not the other way around
- Have a very good understanding about your crop !!
- Implementation require a solid breeding program
- Genotyping might be not so easy and cheap
- Collect good phenotype is imperative
- Test new methods is valid. But don't forget to include an additive GBLUP as a benchmark
- It is multidisciplinary topic.

References

Other references

-  R2D2 Consortium et al., 2021. Why and How to Switch to Genomic Selection: Lessons From Plant and Animal Breeding Experience. <https://doi.org/10.3389/fgene.2021.629737>
-  Excellence in Breeding Platform. <https://excellenceinbreeding.org/>
-  Ferrão et al., 2021. Genomic Selection in an Outcrossing Autotetraploid Fruit Crop: Lessons from Blueberry Breeding. doi: 10.3389/fpls.2021.676326
-  Ferrão et al., 2017. Genomic Selection - State of Art. Book: Genetic Improvement of Tropical Species. In: Hugo Campos and Peter D. S Caligari (Ed.). Springer, 2017
-  Isik et al., 2017. Genetic Data Analysis for Plant and Animal Breeding. Book link

Introduction

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Breeder Equation

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Accuracy

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Blueberry

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Final Consideration

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Thank you !!!

Question ??

About Me:

SCAN ME

