

Master's Thesis

# Development and Validation of a Model for Genomic Selection for Late Blight in Potato

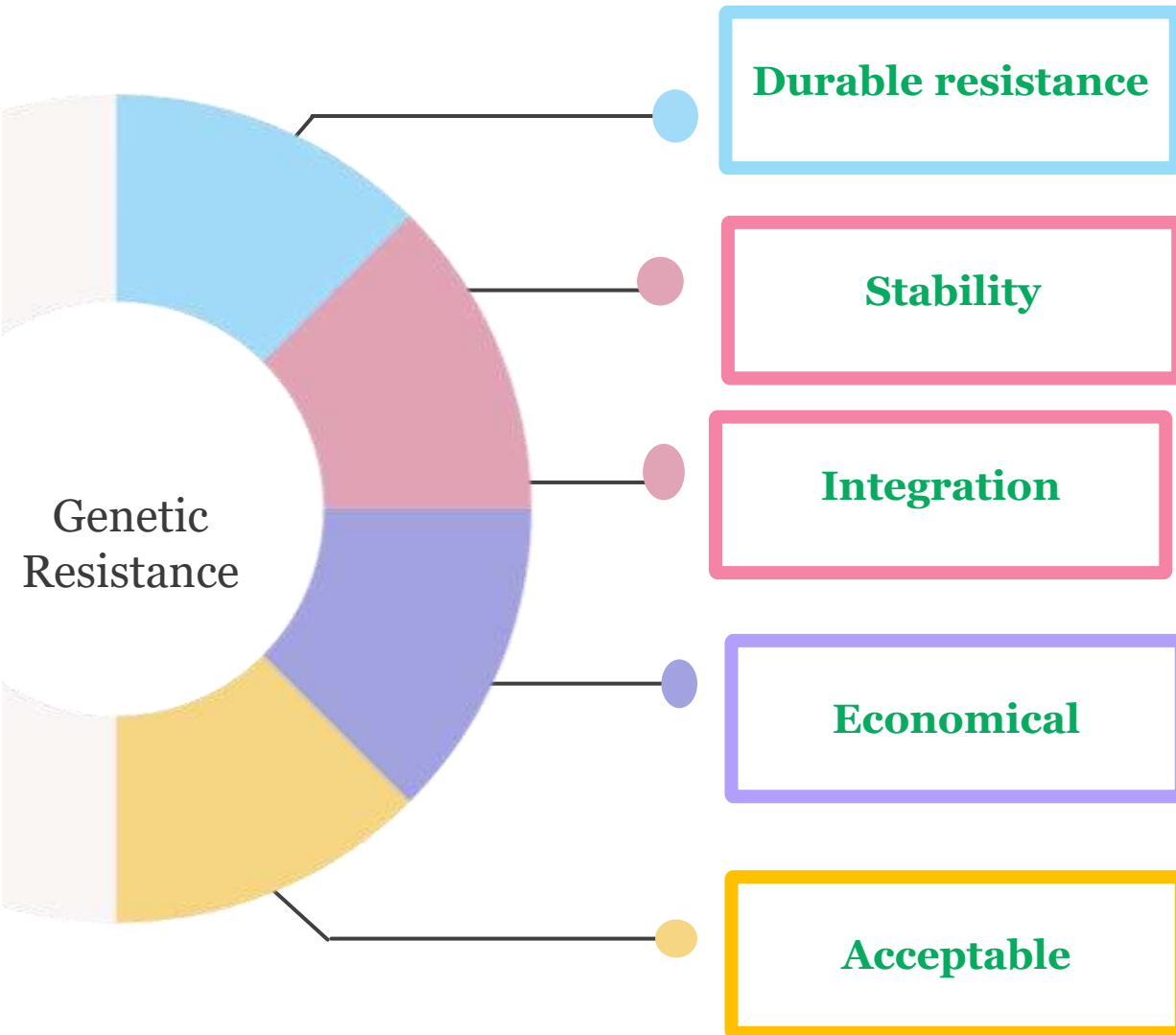
Sagar Kafle



Supervisors: Prof. Just Jensen, Prof. Hermann Bürstmayr

Science for [life]

# Why Plant Protection Needs Breeding



Long lasting solution

Stable resistance across environments

Integration with IPM

Economically viable

Organic production

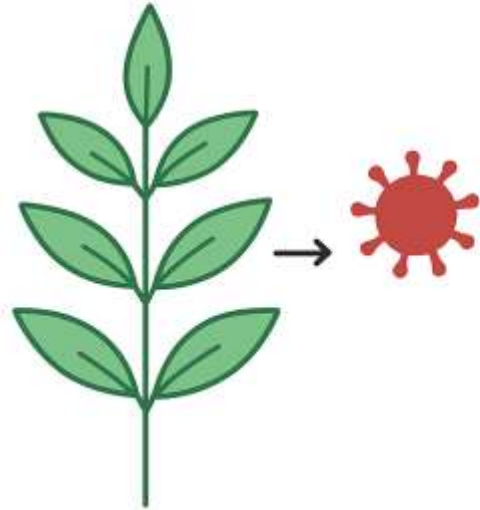
Monogenic

Non-durable

Race specific

Introgression

### Qualitative resistance



Complete protection



### Quantitative resistance



Partial protection



Polygenic (QTLs)

Durable

Race non-specific

MAS/GS

Highly evolving pathogen

Long breeding cycle

Complex Potato Genetics

## R-gene Resistance

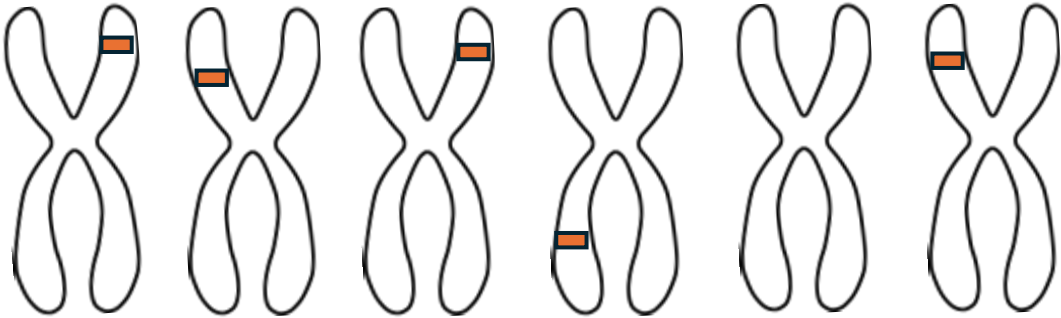


High selective pressure

Non-Durable

Identify & Introgression

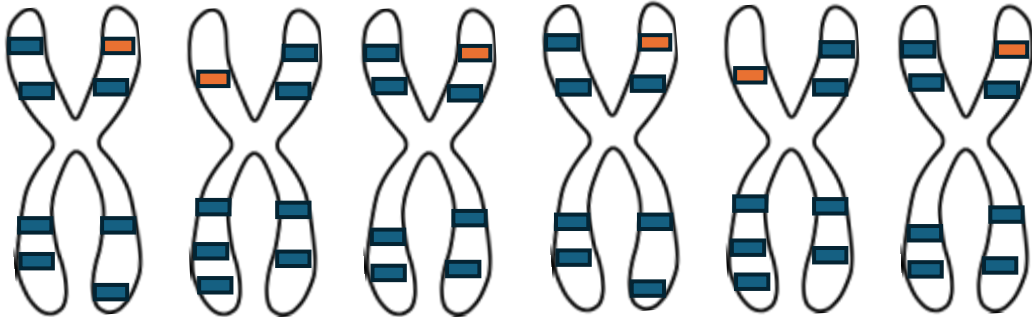
## Major QTLs (MAS)



Durable

Map and introgress

## Major and Minor QTLs (GS)



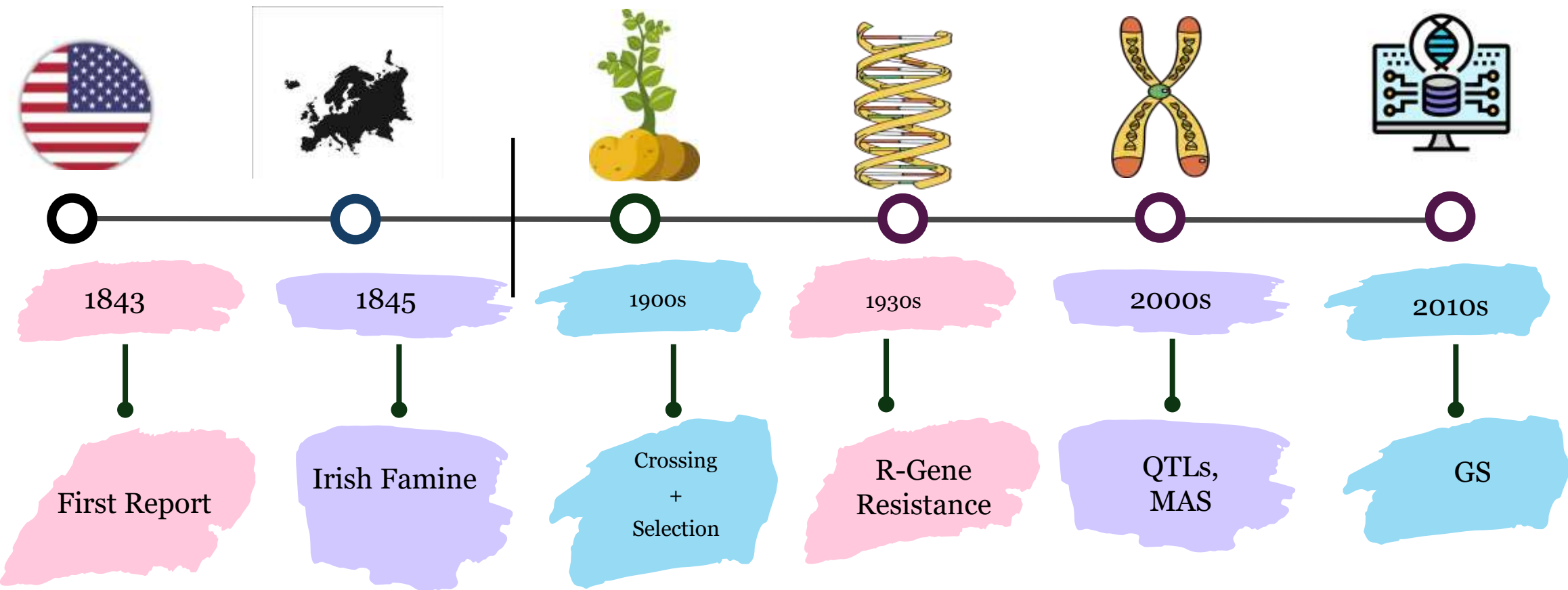
Durable  
Less selective pressure

Modelling

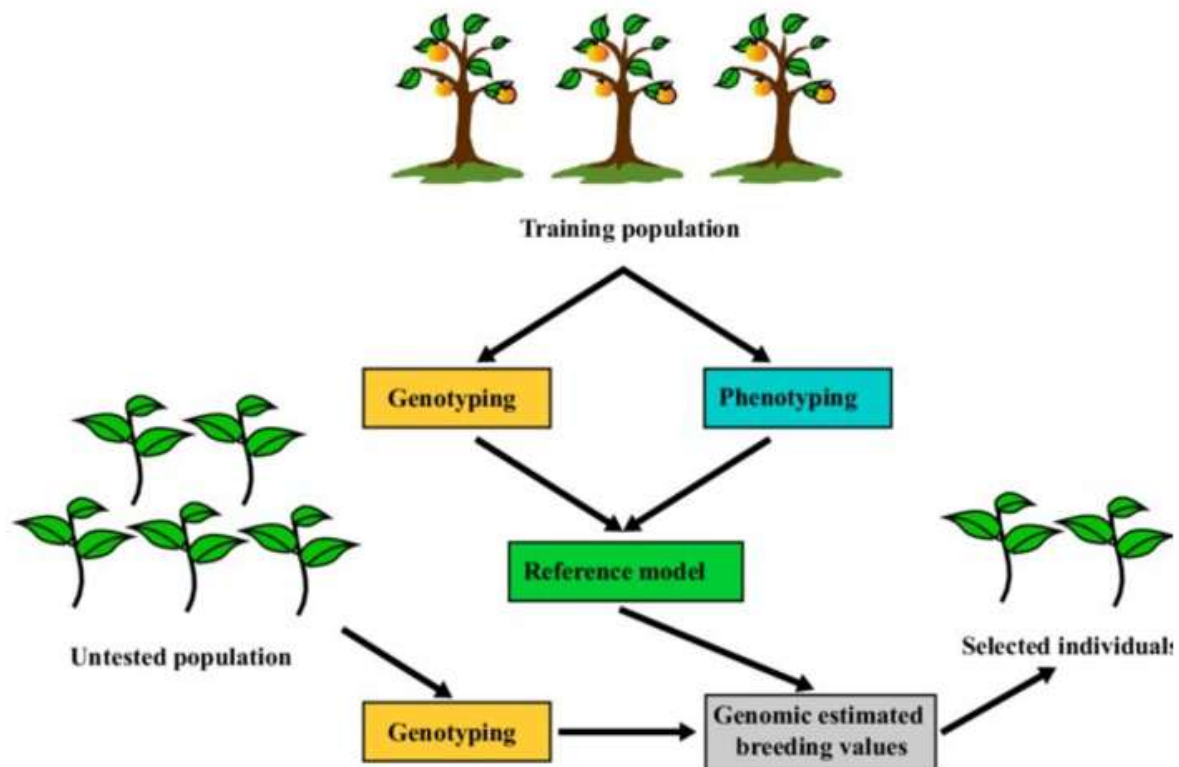
Poland et al., 2009

Genetic Resistance

# LATE BLIGHT: A Problem over century



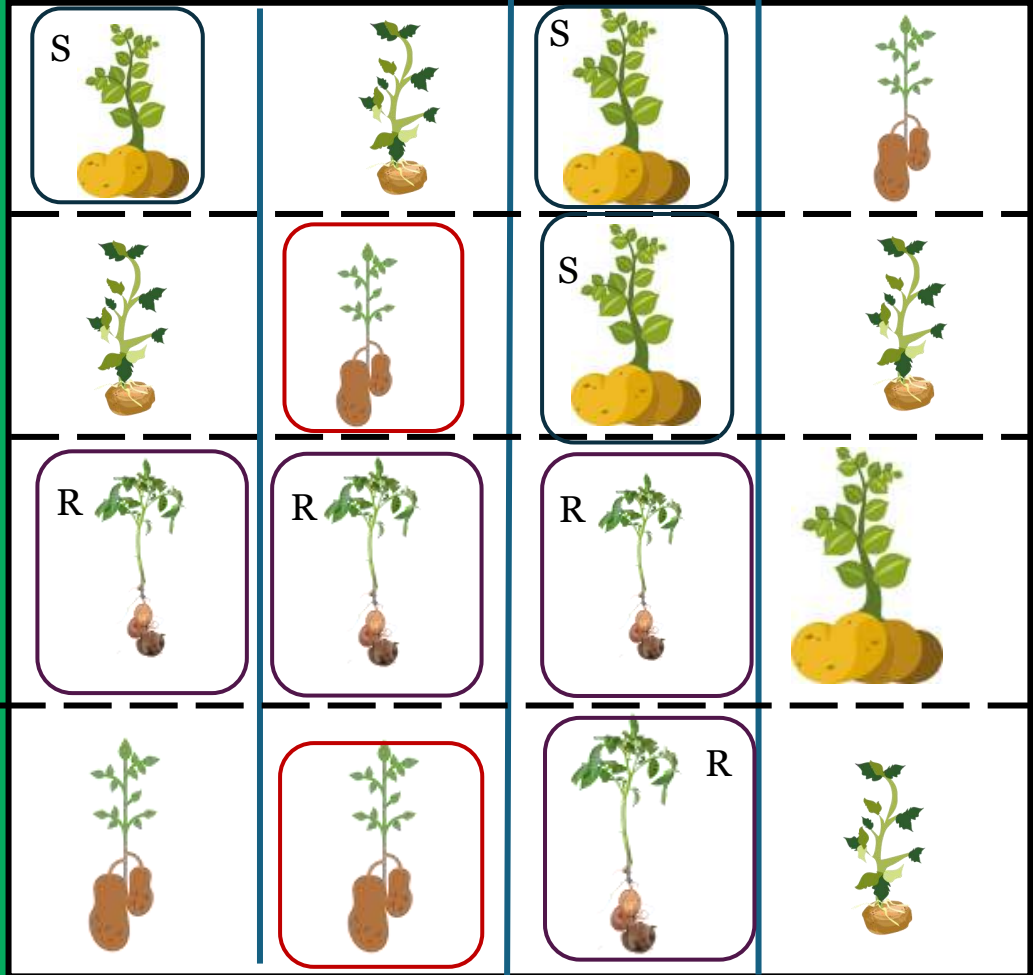
# Genomic Selection



- ✓ Out of 144 QTLs, 25 % -Resistance Gene Analogues (RGA), 50% -Durable Resistance Loci (DRLs) (Danan et al. 2011)

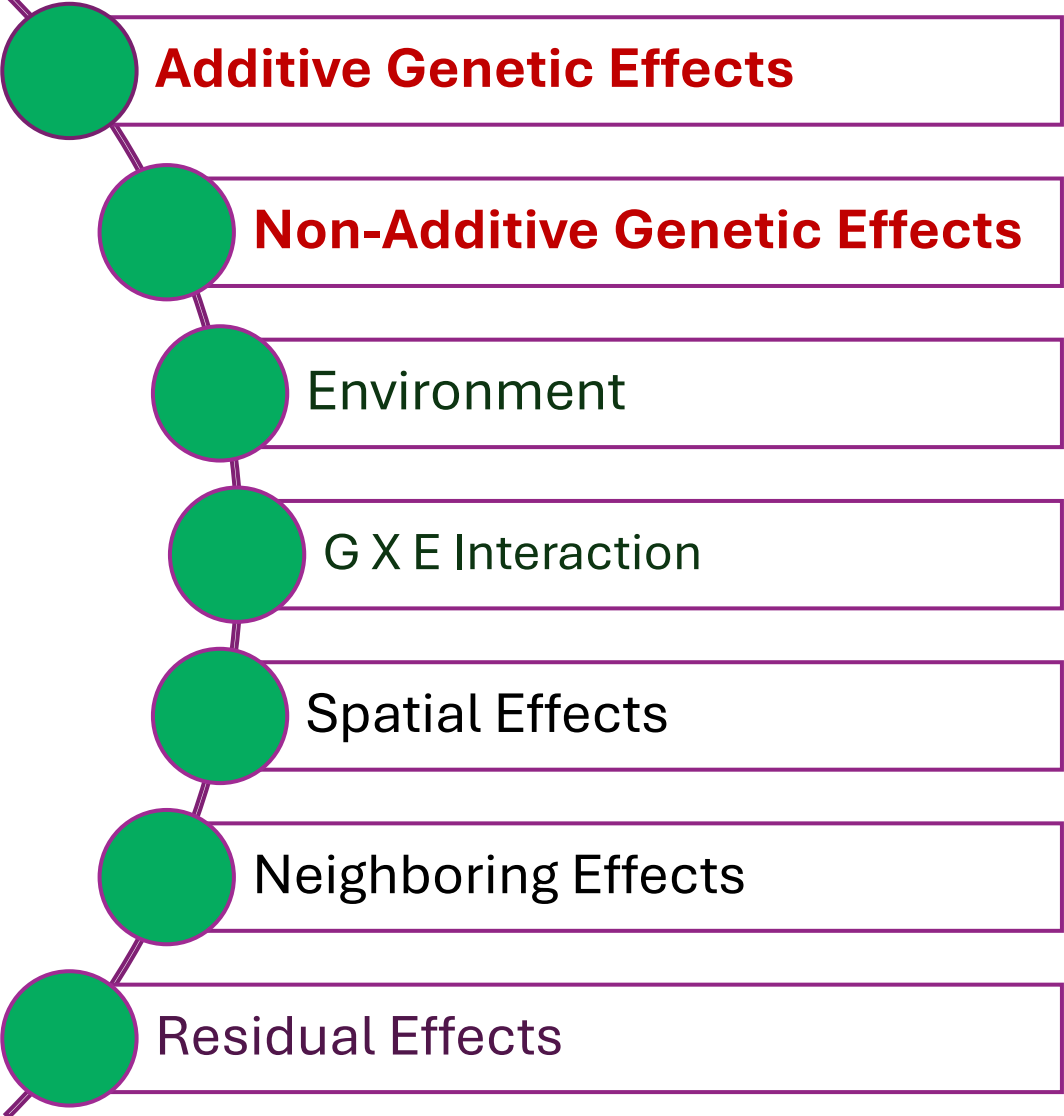
- ✓ Prediction of untested lines
- ✓ Breeding value of tested lines

# Sources of Variation



Low  
Fertility

High  
Fertility



# Materials and Methods

## Plant Materials



Table 1 Description of potato lines tested over different years.

	2024	2023	2022
<b>No. of plots</b>	4000	2970	3078
<b>Lines tested</b>	1212	975	1717
<b>Lanes (Blocks)</b>	3	2	2
<b>Rows</b>	18	18	18
<b>Columns</b>	84	110	86

Table 2 Description of potato lines, grouped by market segment.

Group	Notation	No. of records	No. of lines
1	Very Early	74	11
2	Early	109	3
3	Table North(North Europe)	1703	346
4	Table South without frying quality	1253	259
5	Red	16	1
6	Salad types	165	28
7	Table north with frying quality	1163	341
8	French fry	1502	415
9	Chips/Crisps	763	194
10	Starch	2985	1465
11	Starch	4	1
12	Starch	334	132
13	Unknown	4	1

# Field Design

Blocks: 2

Testing Rows: 18  
Spreader Rows: 10

Columns: 86

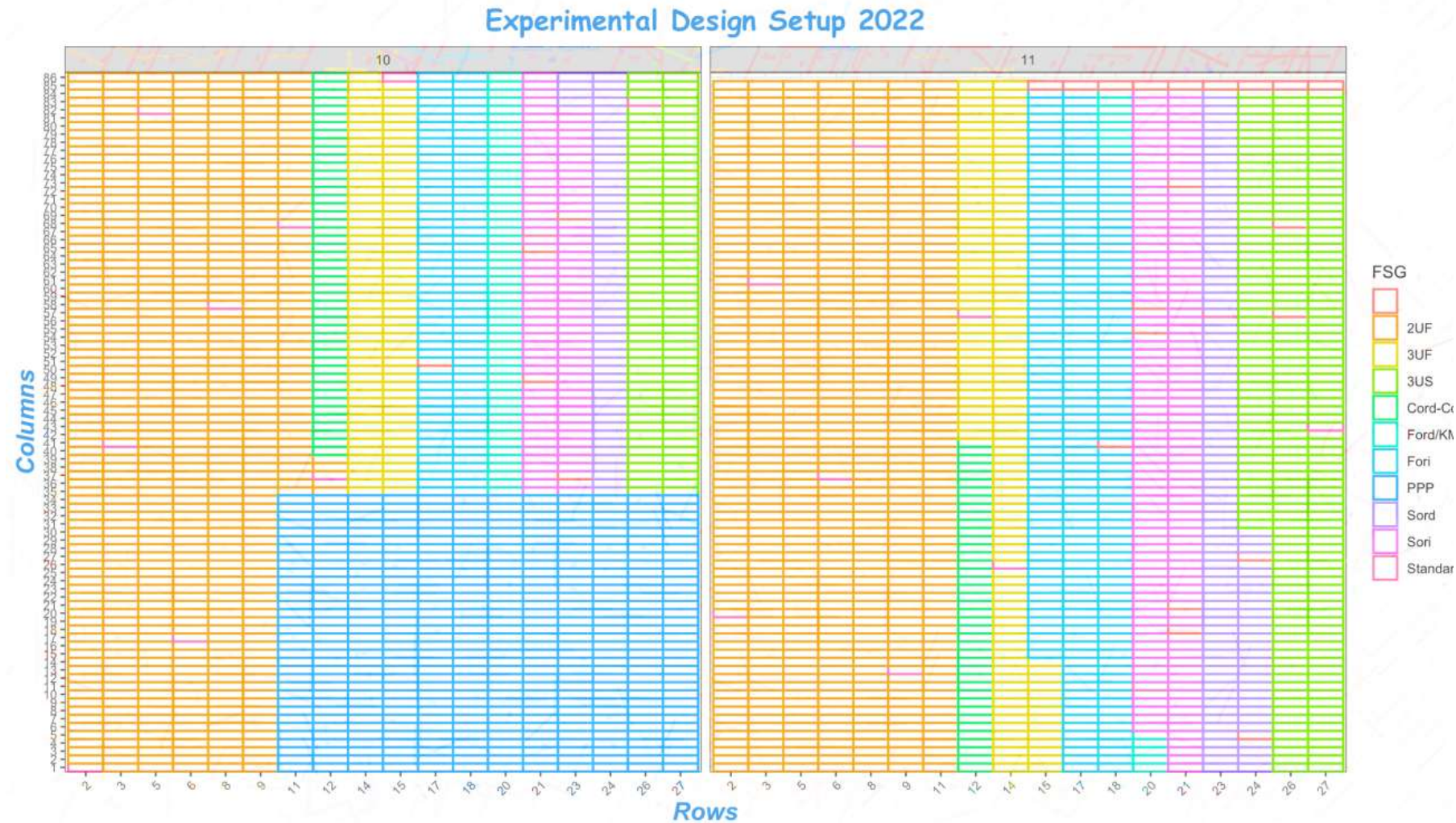
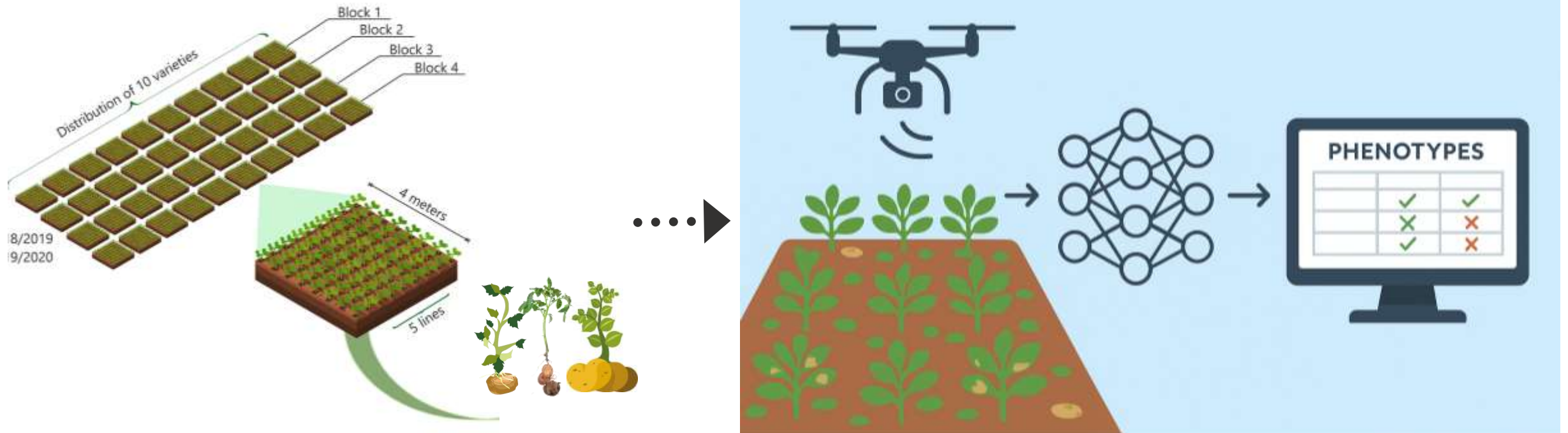


Fig 1: Experimental field design for year 2022

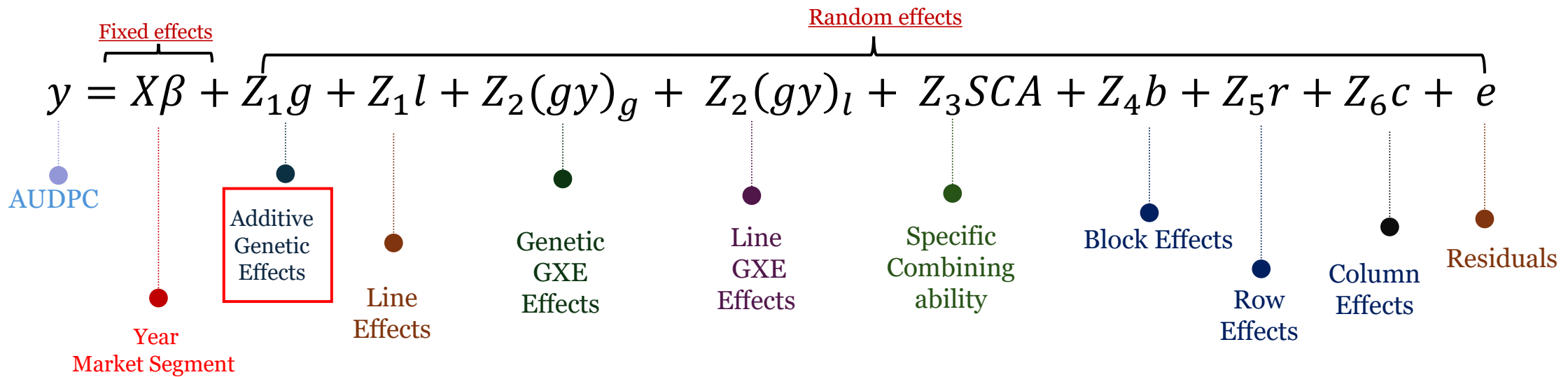
# High Throughput Phenotyping System



High Throughput Phenotyping



# MODEL 2: Row-Column Model



**ABLUP:**  $y = X\beta + Z_1g + Z_2(gy)_g + Z_3SCA + Z_4b + Z_5r + Z_6c + e$

# Results: Variance Components

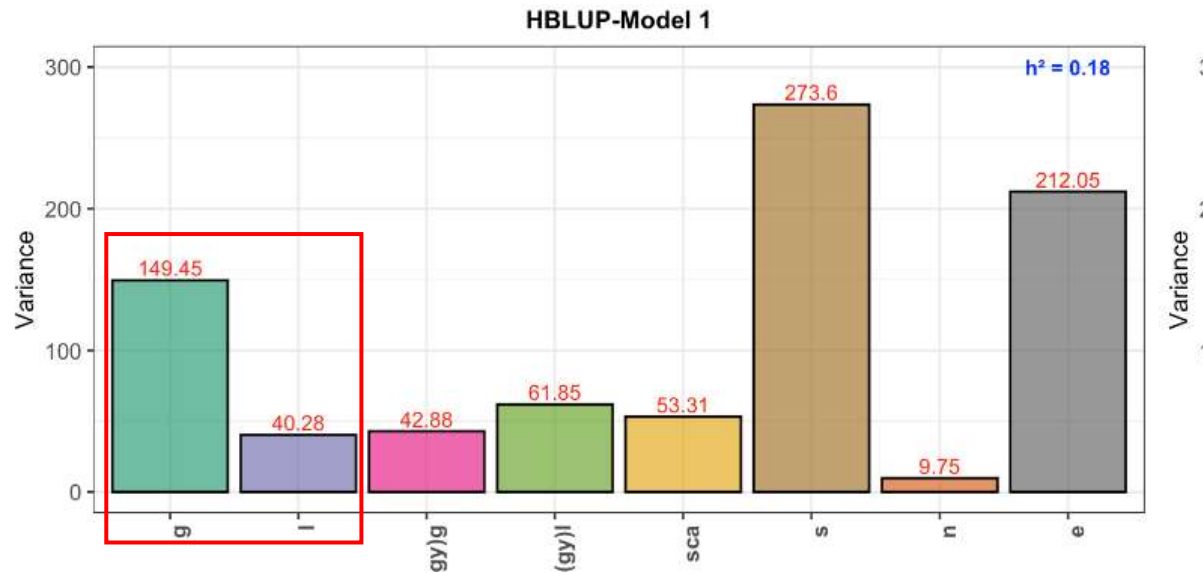


Fig 2. Estimated variance components for AUDPC of late blight of potato.

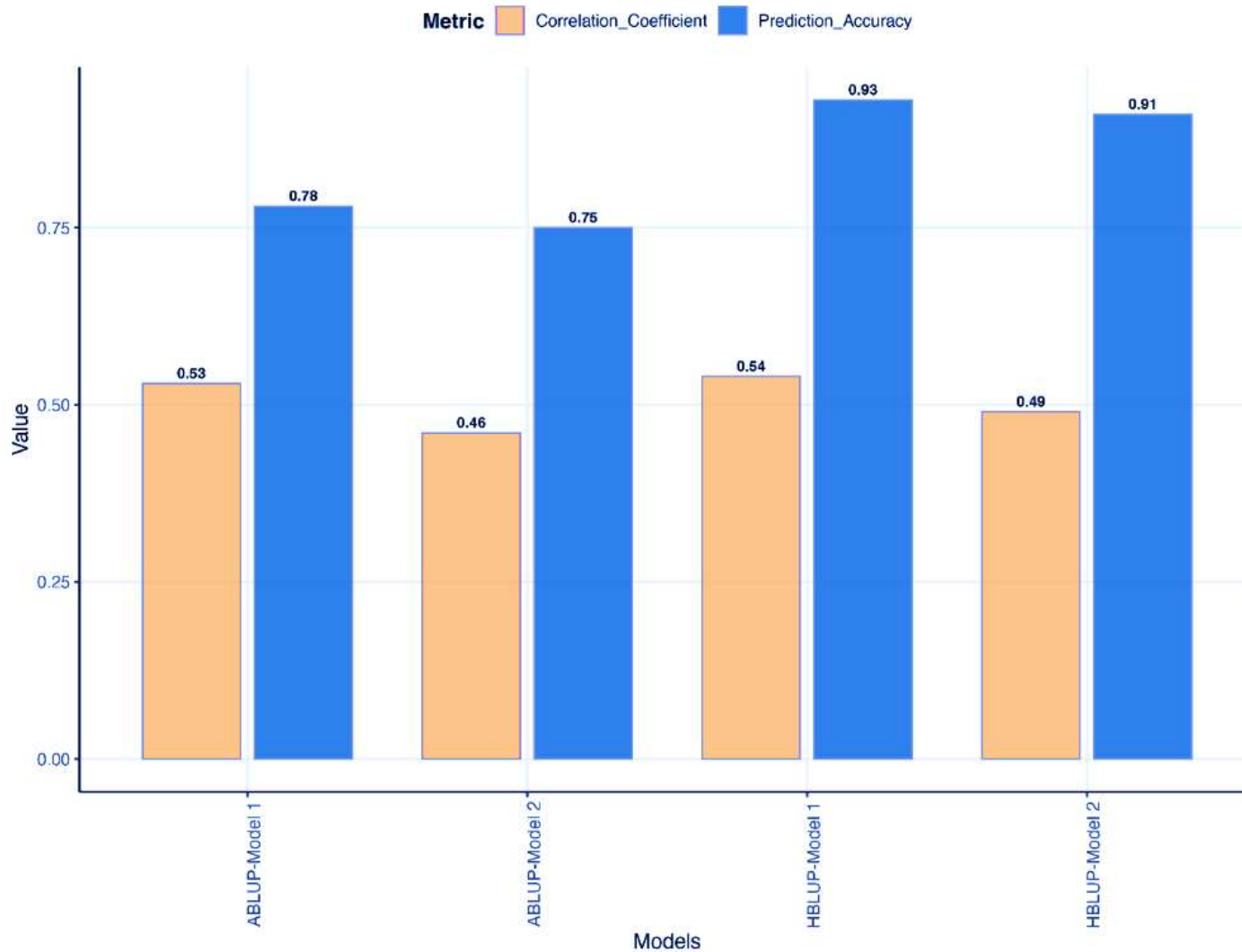
- ✓ Spatial Effects affects the disease score the most
- ✓ Genetic component also significantly influence late blight resistance

Table 3. Heritability of traits for different models.

Model	$h_{ns}^2$	$h_{yc}^2$
HBLUP Model 1	0.18	0.34
ABLUP Model 1	0.24	0.47
HBLUP Model 2	0.16	0.29
ABLUP Model 2	0.21	0.38

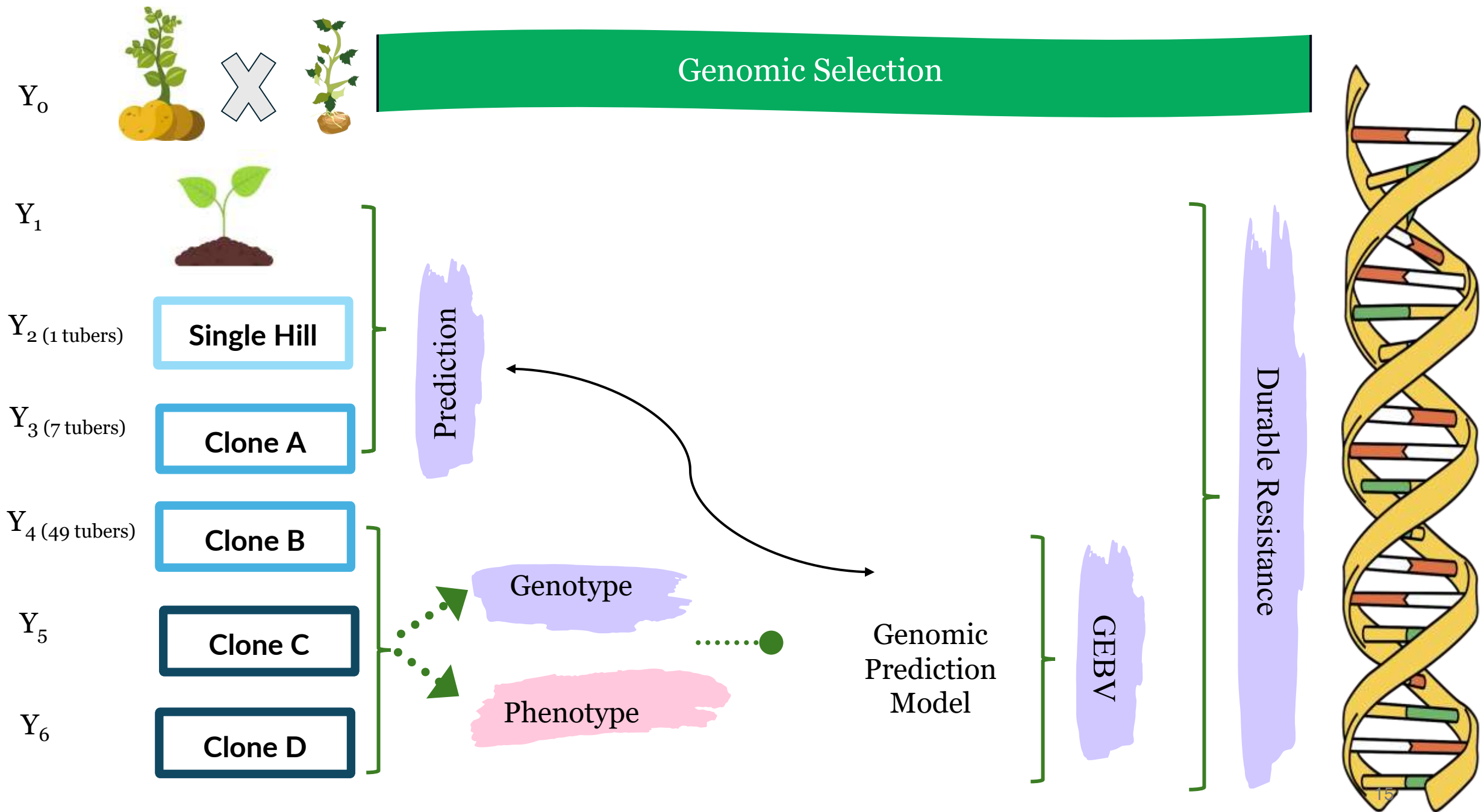
# Accuracy of Prediction

Prediction Accuracy and Correlation Coefficient for Different Models



- ✓ Accuracy of prediction for late blight resistance ranged from 0.75-0.93
- ✓ Relatively higher accuracy of prediction

Figure 3. Accuracy of prediction across different models.



## 5. Conclusions

- ✓ Late blight of potato: Globally the most destructive disease of potato
- ✓ Fungicide application remains common, but are increasingly unpopular and subjected to regulatory constraints
- ✓ Given the polygenic nature of the late blight, genomic selection could offer such a promising solution for developing resistance to late blight.
- ✓ Combining gene pyramiding with quantitative resistance may offer sustainable, high-level disease resistance.
- ✓ Even after 100+ years of breeding, truly durable resistance remains difficult to achieve.

## 7. References

Danan, Sarah, Jean-Baptiste Veyrieras, and Véronique Lefebvre. 2011. “Construction of a Potato Consensus Map and QTL Meta-Analysis Offer New Insights into the Genetic Architecture of Late Blight Resistance and Plant Maturity Traits.” *BMC Plant Biology* 11:16. doi:10.1186/1471-2229-11-16.

Poland, Jesse A., Peter J. Balint-Kurti, Randall J. Wisser, Richard C. Pratt, and Rebecca J. Nelson. 2009. “Shades of Gray: The World of Quantitative Disease Resistance.” *Trends in Plant Science* 14(1):21–29. doi:10.1016/j.tplants.2008.10.006.

Poland, Jesse, and Jessica Rutkoski. 2016. “Advances and Challenges in Genomic Selection for Disease Resistance.” *Annual Review of Phytopathology* 54:79–98. doi:10.1146/annurev-phyto-080615-100056.

Paluchowska, Paulina, Jadwiga Śliwka, and Zhimin Yin. 2022. “Late Blight Resistance Genes in Potato Breeding.” *Planta* 255(6):127. doi:10.1007/s00425-022-03910-6.

**Supervisor  
Committee**

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*Thank You...*