

# Genomics Update

4<sup>th</sup> International Kernza Conference

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**KANSAS STATE**  
**UNIVERSITY**



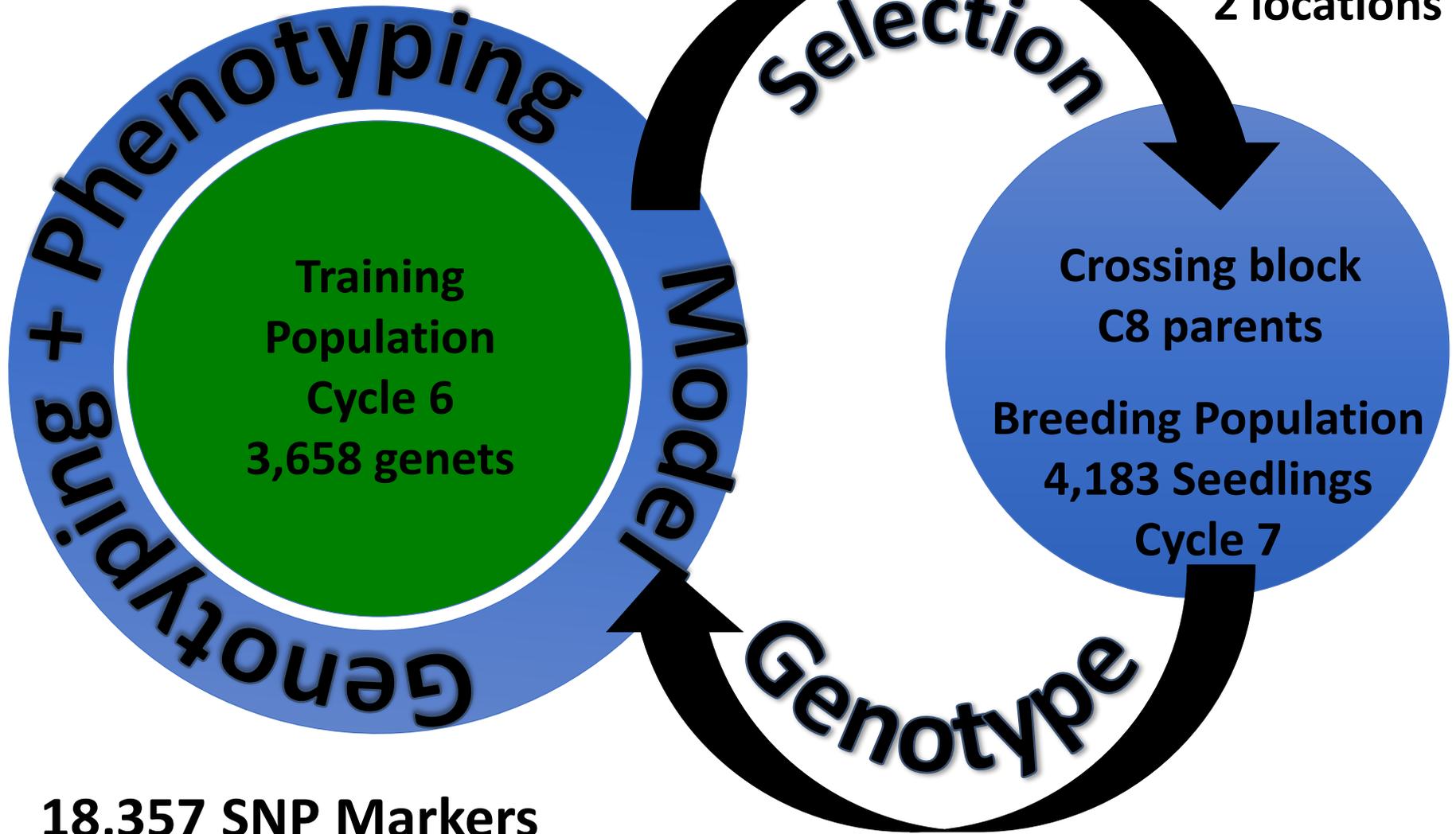
Malone Family Land  
Preservation Foundation

# Genomics Update

- Cycle 7 GS Results and Future Direction
- Current Status of Genome Assembly
- Paternity Analysis

**2 years, 46 traits**

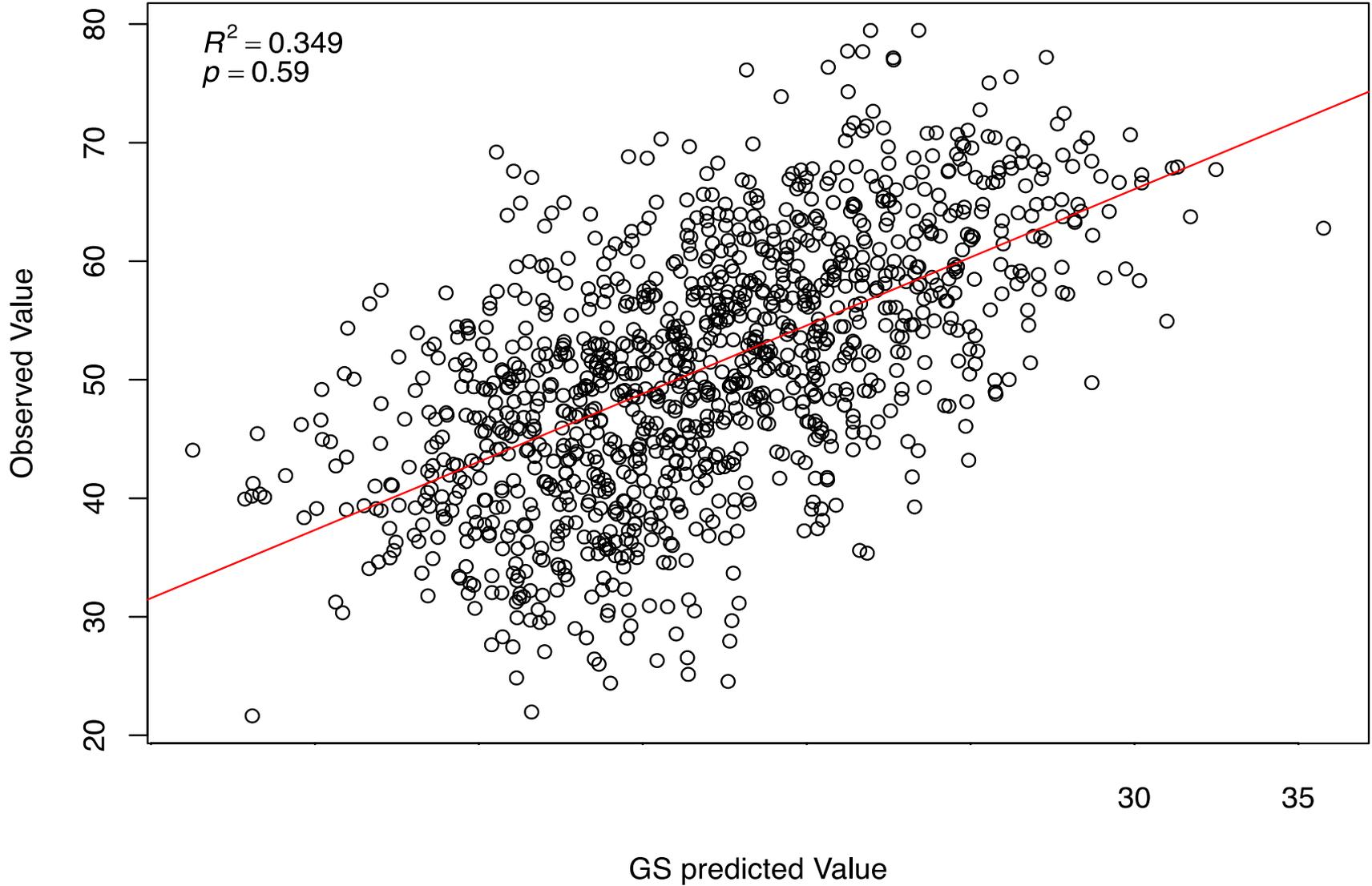
**Training population:  
1200 genets  
2 locations**



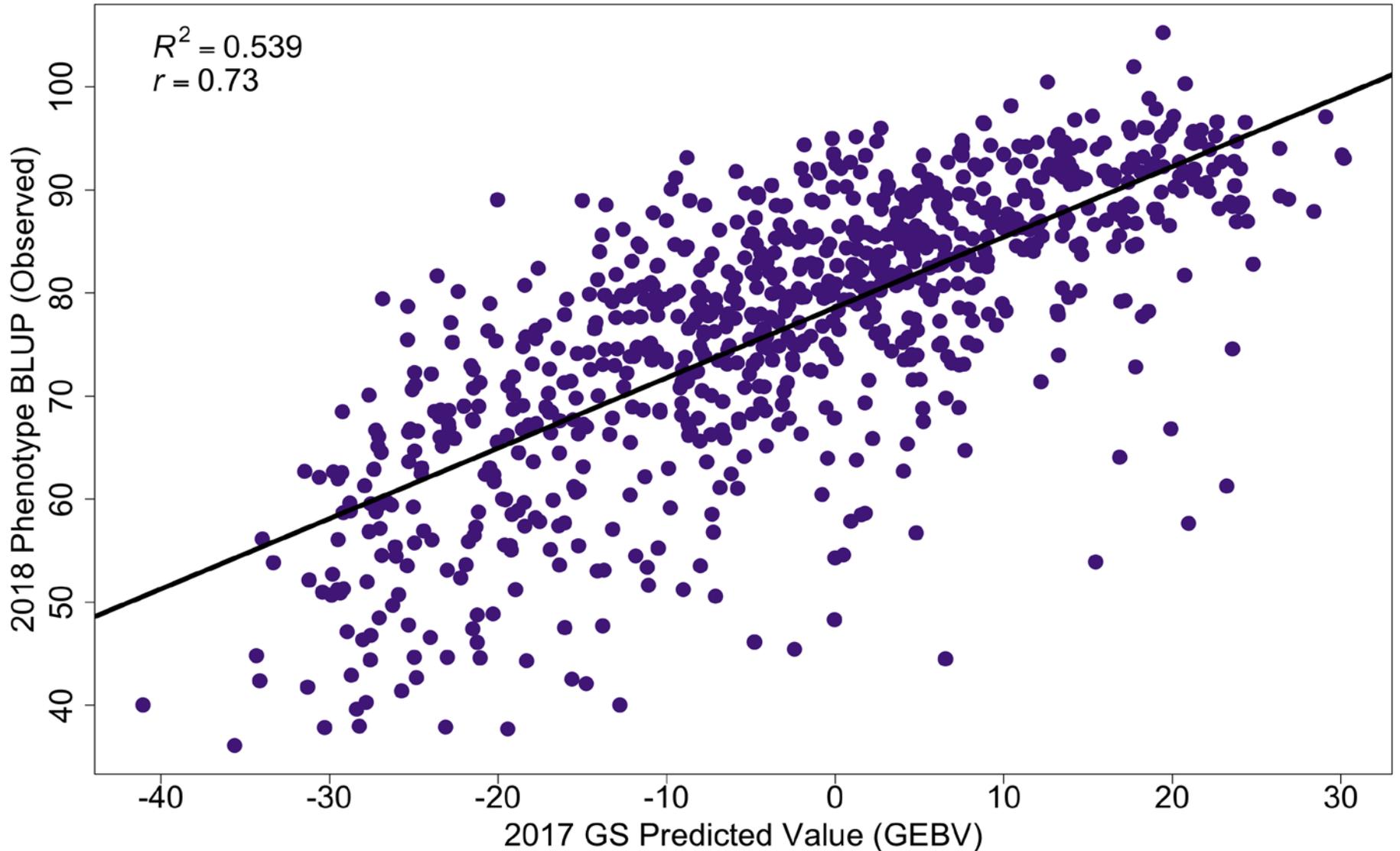
**18,357 SNP Markers**

**22,701 SNP Markers**

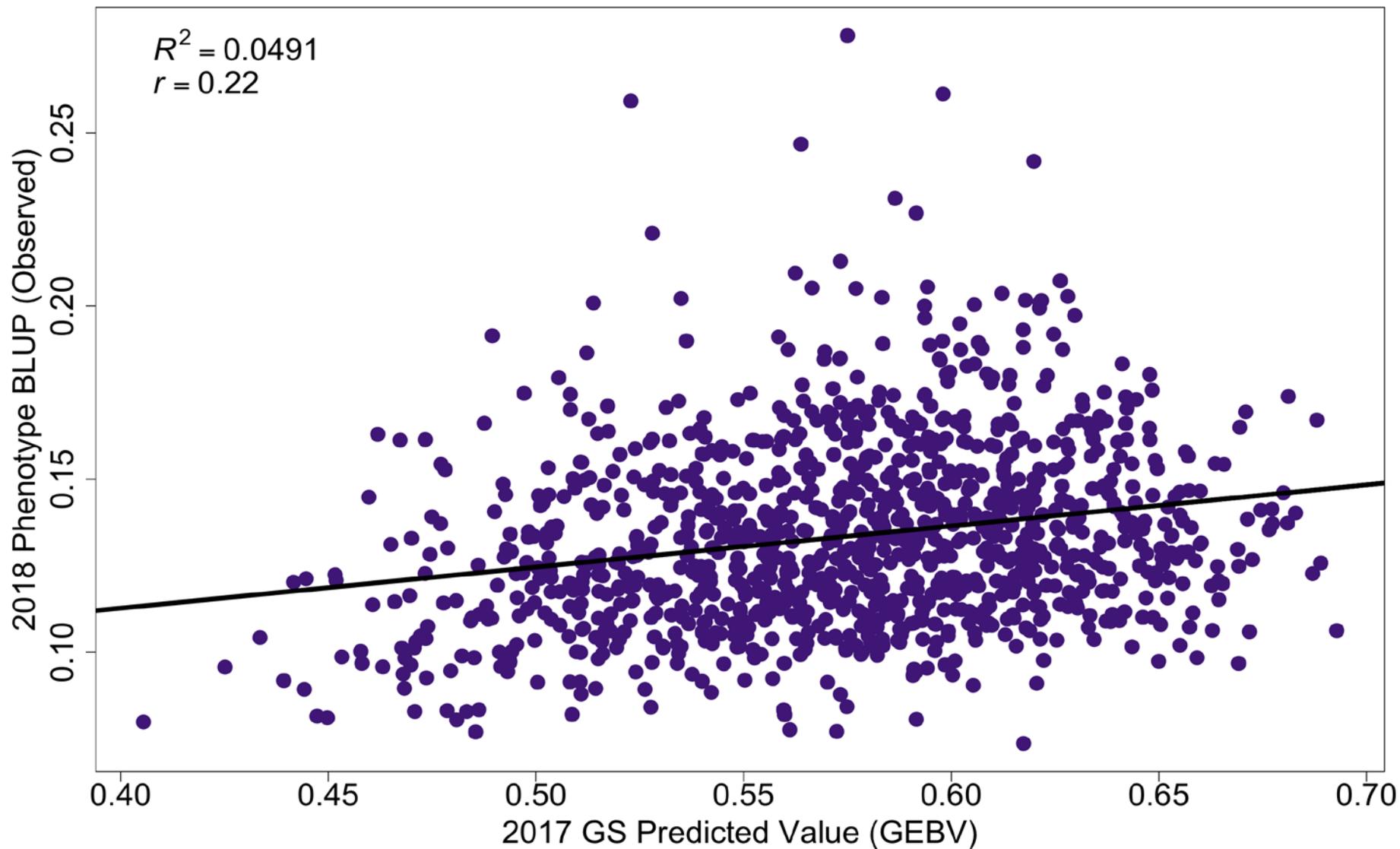
# Stem Angle



# Free Threshing 2018

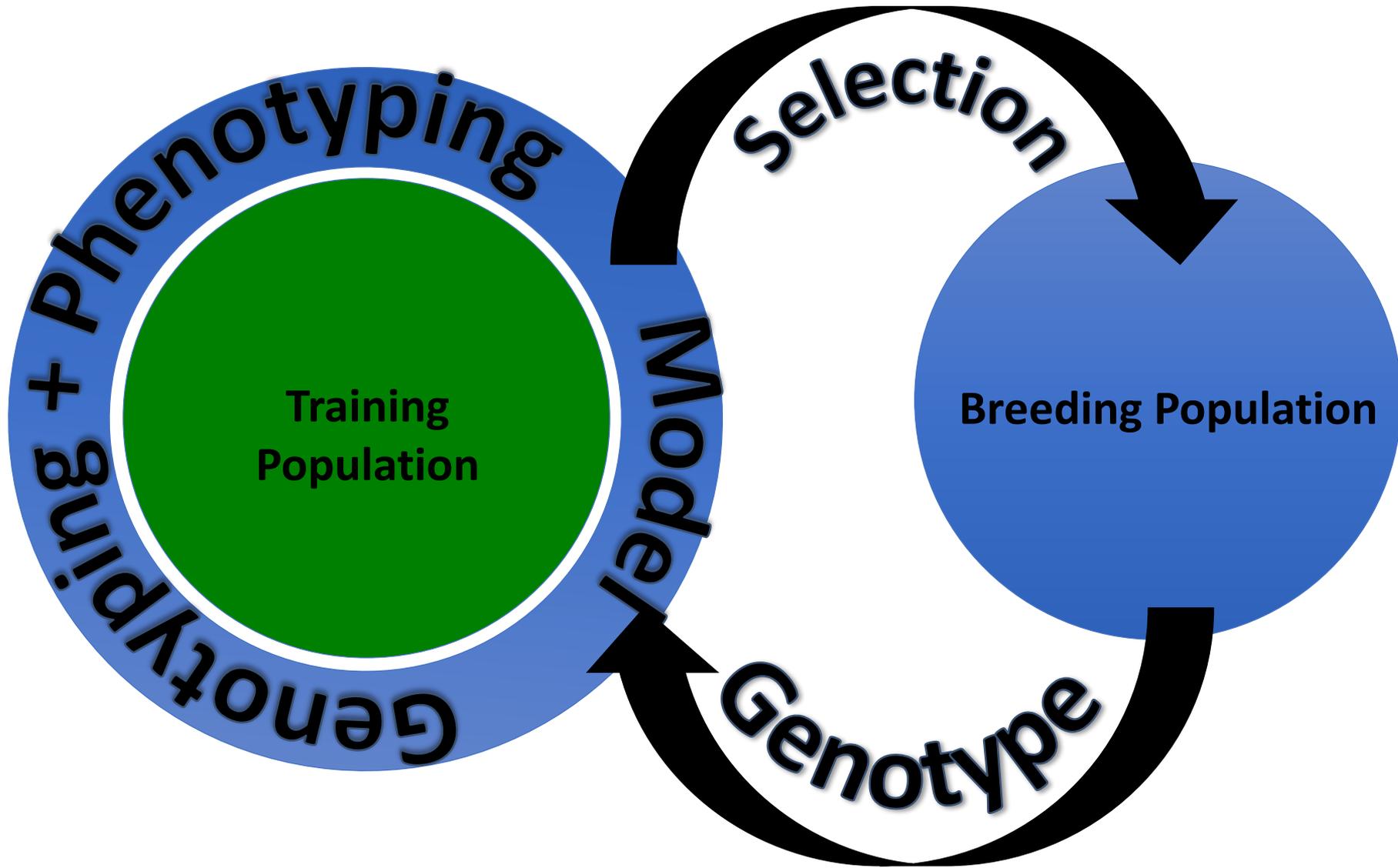


# Spike Yield 2018



# Traits 2018

Trait	$h^2$	r
Free Threshing	0.36	0.73
Shattering	0.65	0.71
Seed Length	0.72	0.68
Plant Height	0.48	0.59
Stem Angle	0.58	0.59
Spikelets per Head	0.21	0.57
Milligrams per Seed	0.53	0.55
Maturity	0.63	0.54
Percent Fertility	0.48	0.50
Seeds per Head	0.44	0.45
Seed Density	0.66	0.31
Spike Yield in grams	0.40	0.22
Florets per Spike	0.24	0.14



**Genotyping + Phenotyping**

**Training Population**

**Model**

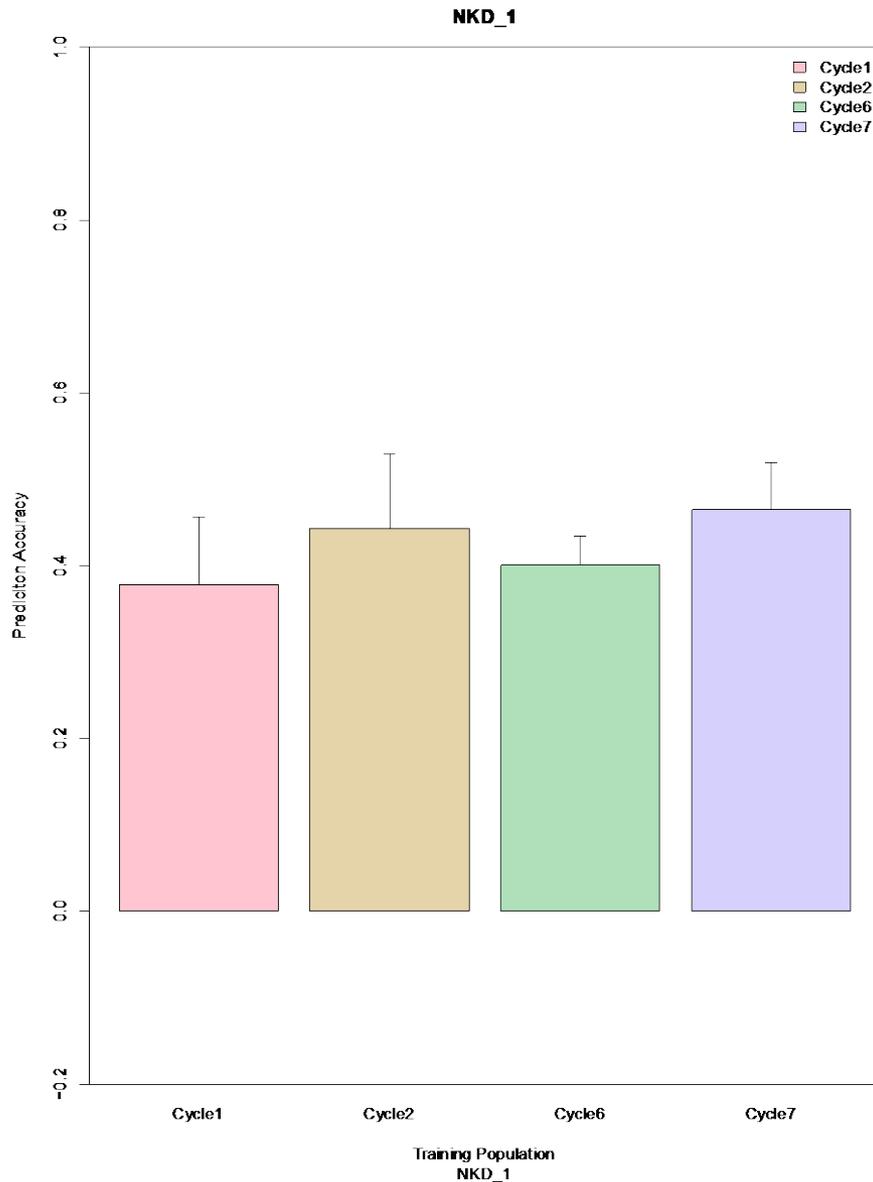
**Selection**

**Breeding Population**

**Genotype**

# IWG Breeding Programs

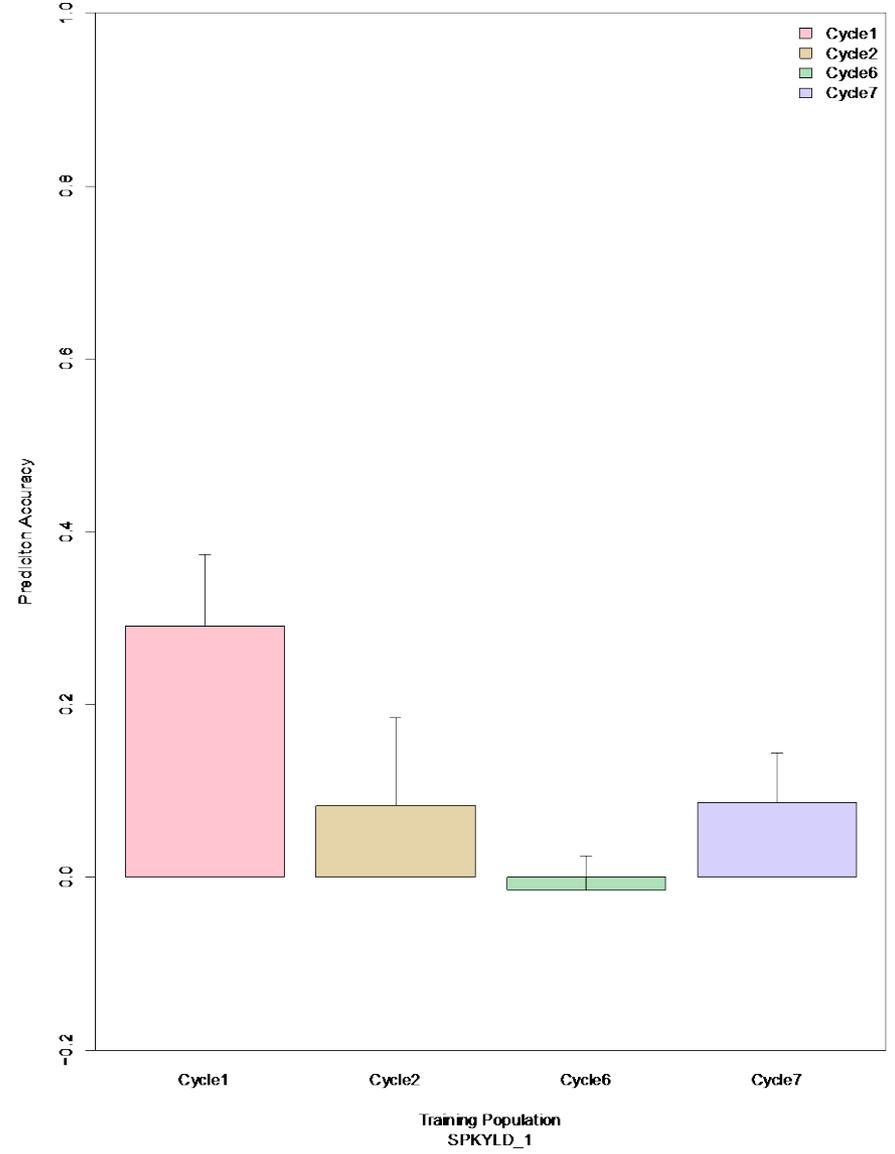


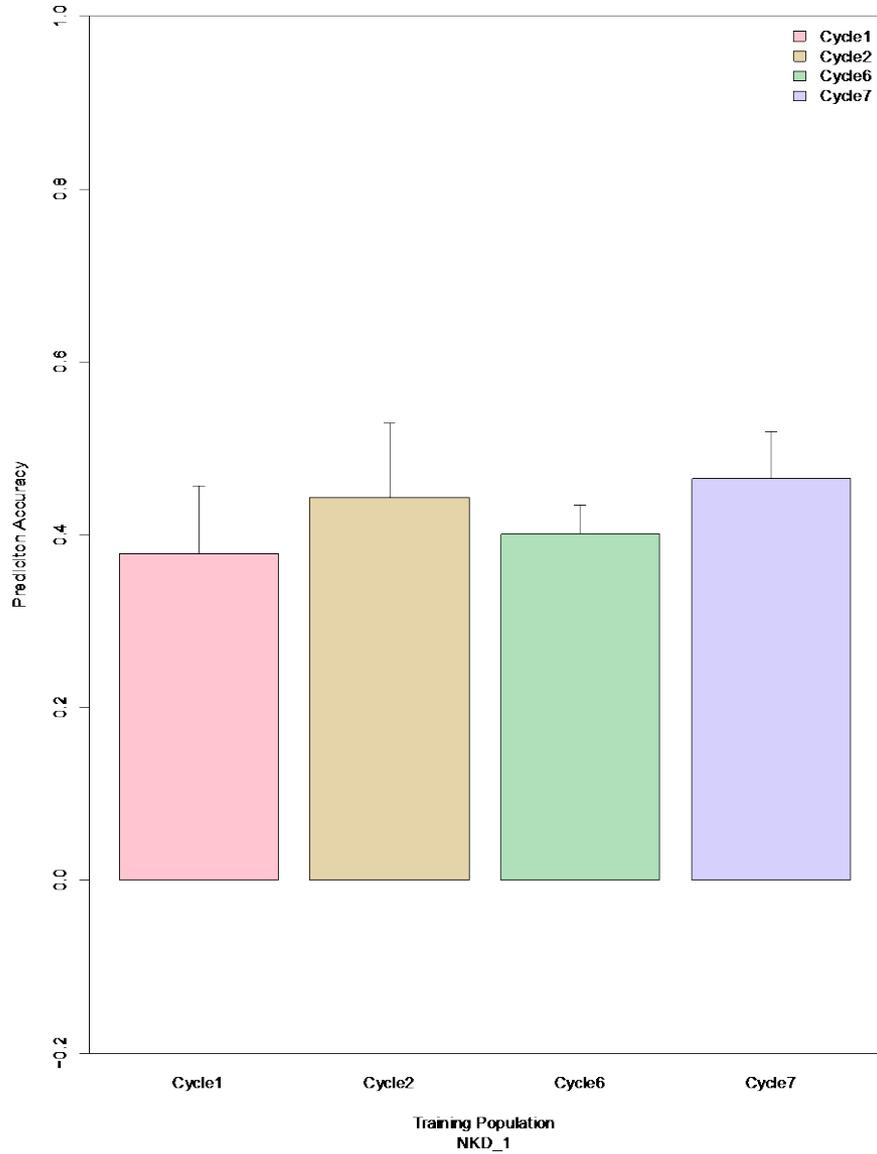
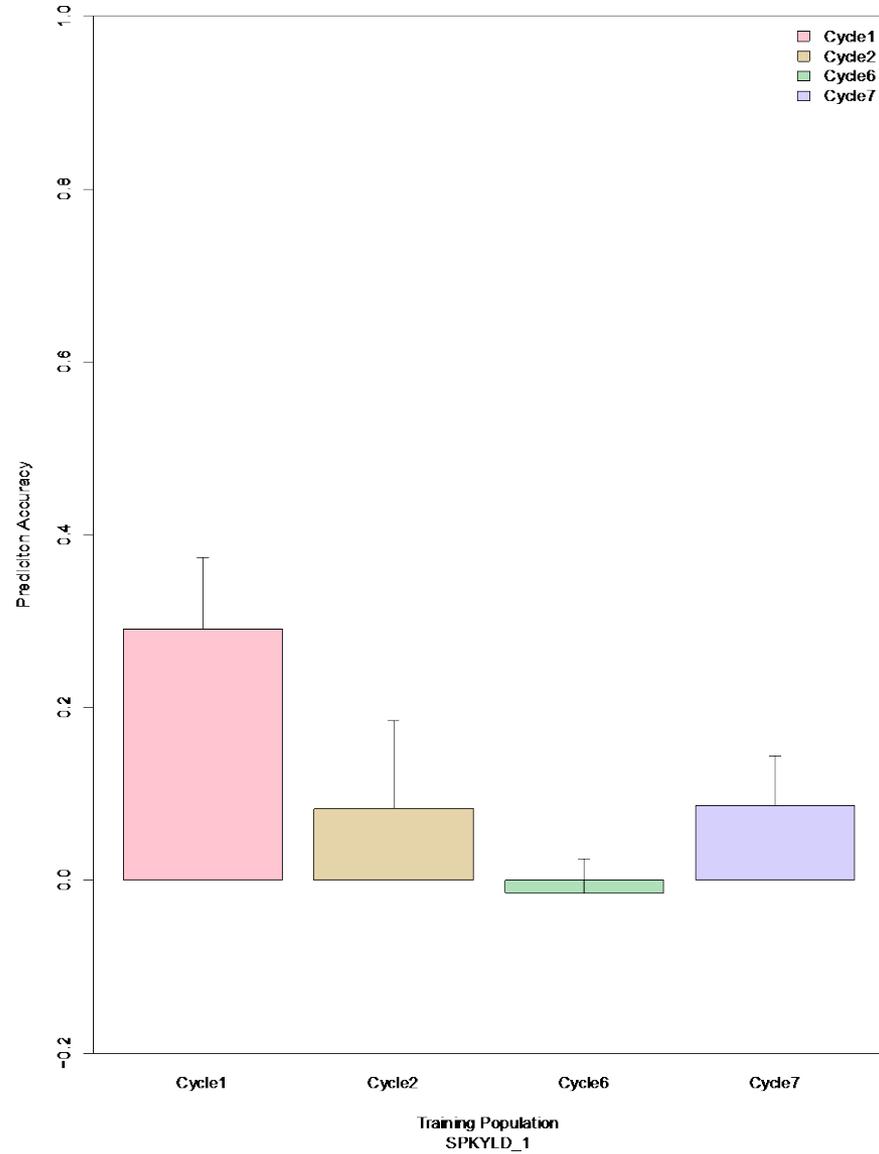


Cycle 1 & 2 UMN  
Cycle 6 & 7 TLI

Using all other cycles as training population to predict each cycle—Cycle 2, 6, & 7 form the training population to predict Cycle 1.

# SPKYLD\_1



**NKD\_1****SPKYLD\_1**

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# Genomic effects of neo-domestication



**Rodale Research Center  
(Pennsylvania)**

300 wild USDA PIs /  
forage varieties

Best 14

2 breeding cycles

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## Rodale Research Center (Pennsylvania)

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Best 14

2 breeding cycles

Best 12  
'Rodale Selections'

## The Land Institute (Kansas)

Starting material for first  
breeding cycle

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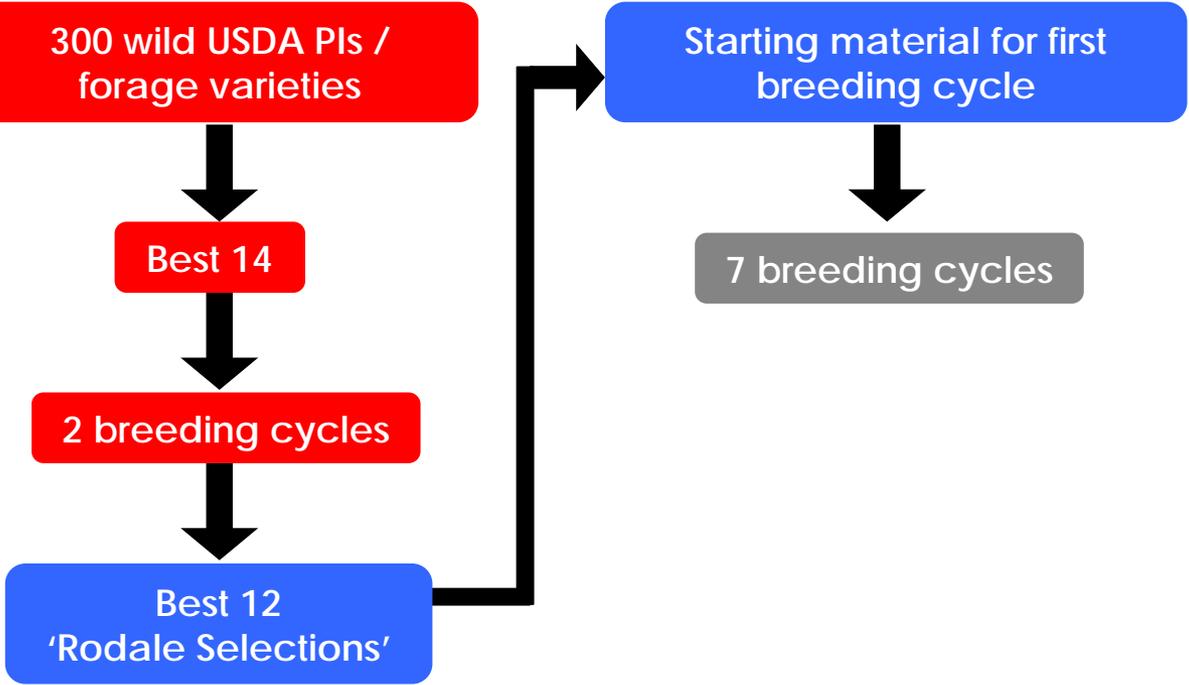
2 breeding cycles

Best 12  
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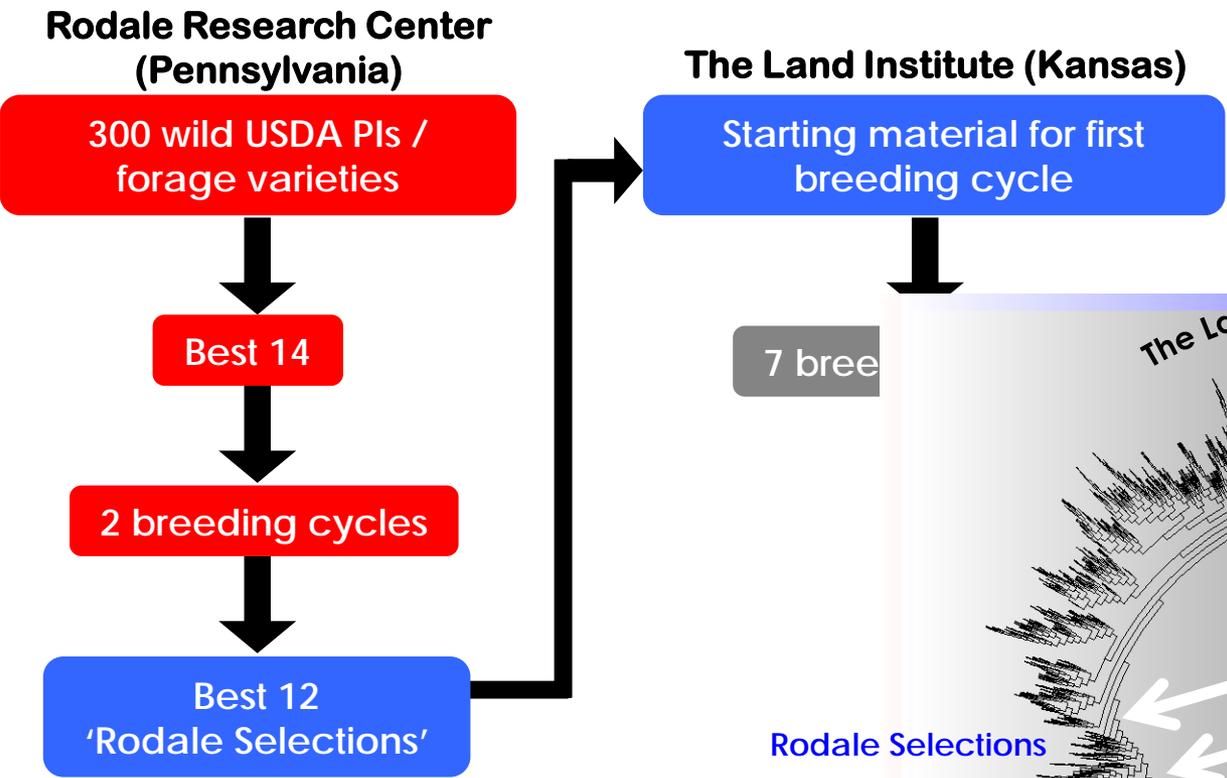
## The Land Institute (Kansas)

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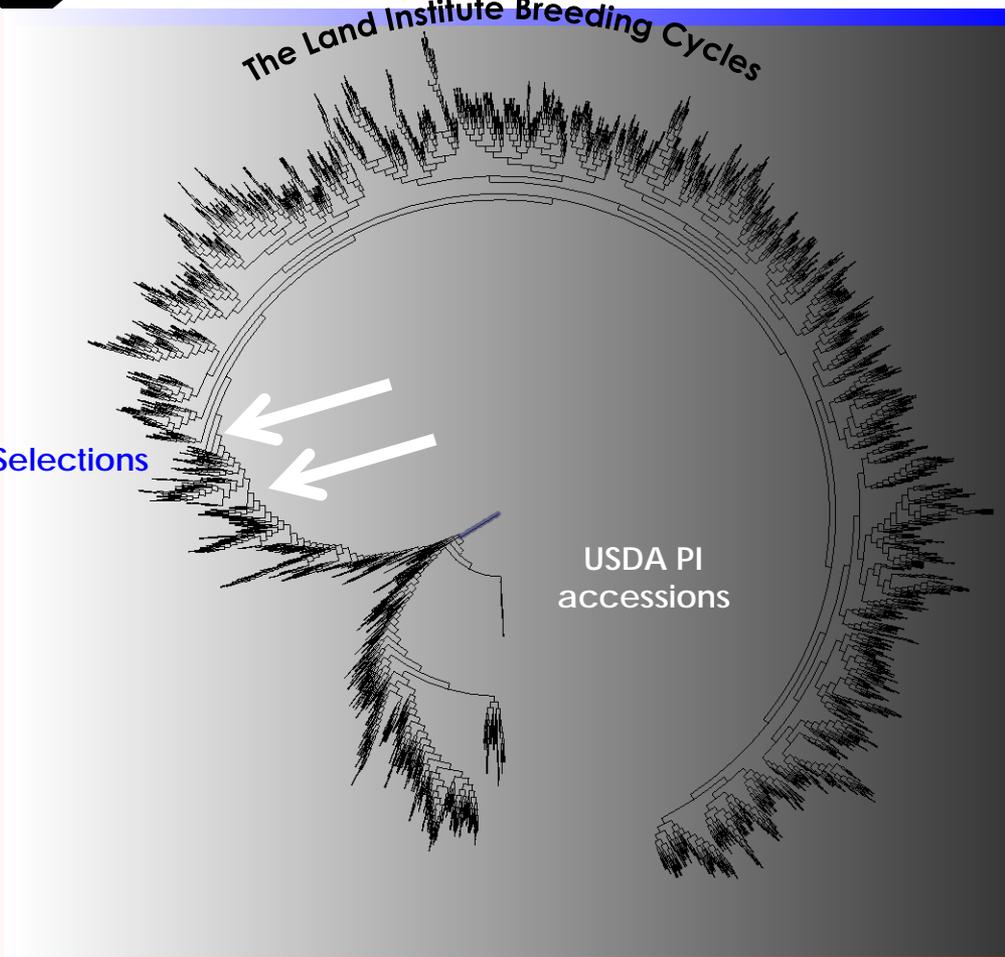
7 breeding cycles



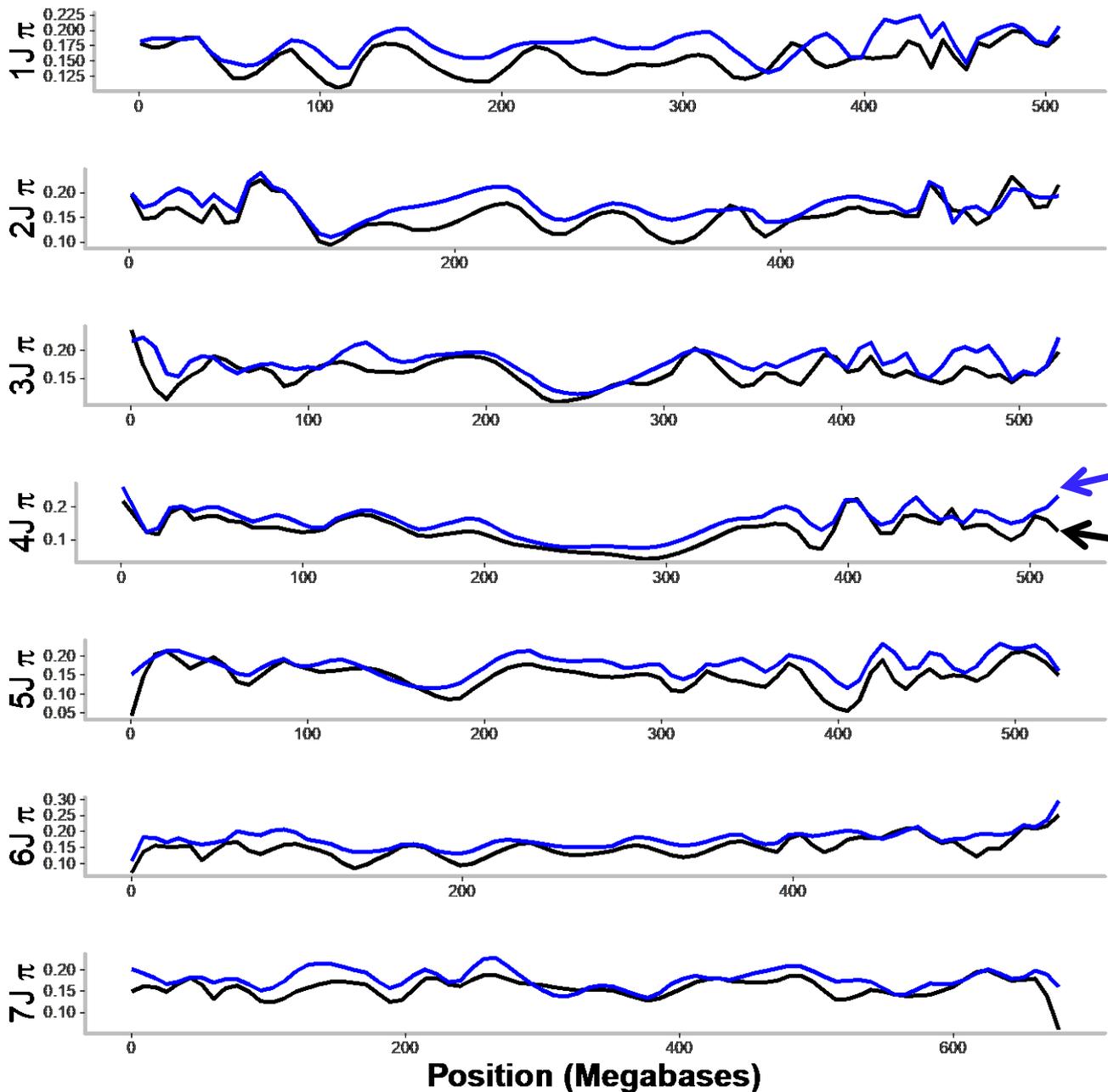
# Genomic effects of neo-domestication



**Genotyping (via GBS) of  
TLI Cycles 5-7 +  
Rodale Selections +  
USDA PI accessions**



# Genomic effects of neo-domestication - $\pi$



Rodale selections

TLI Cycle 7

**Genome-wide  
reductions in  
nucleotide  
diversity across  
the genome**

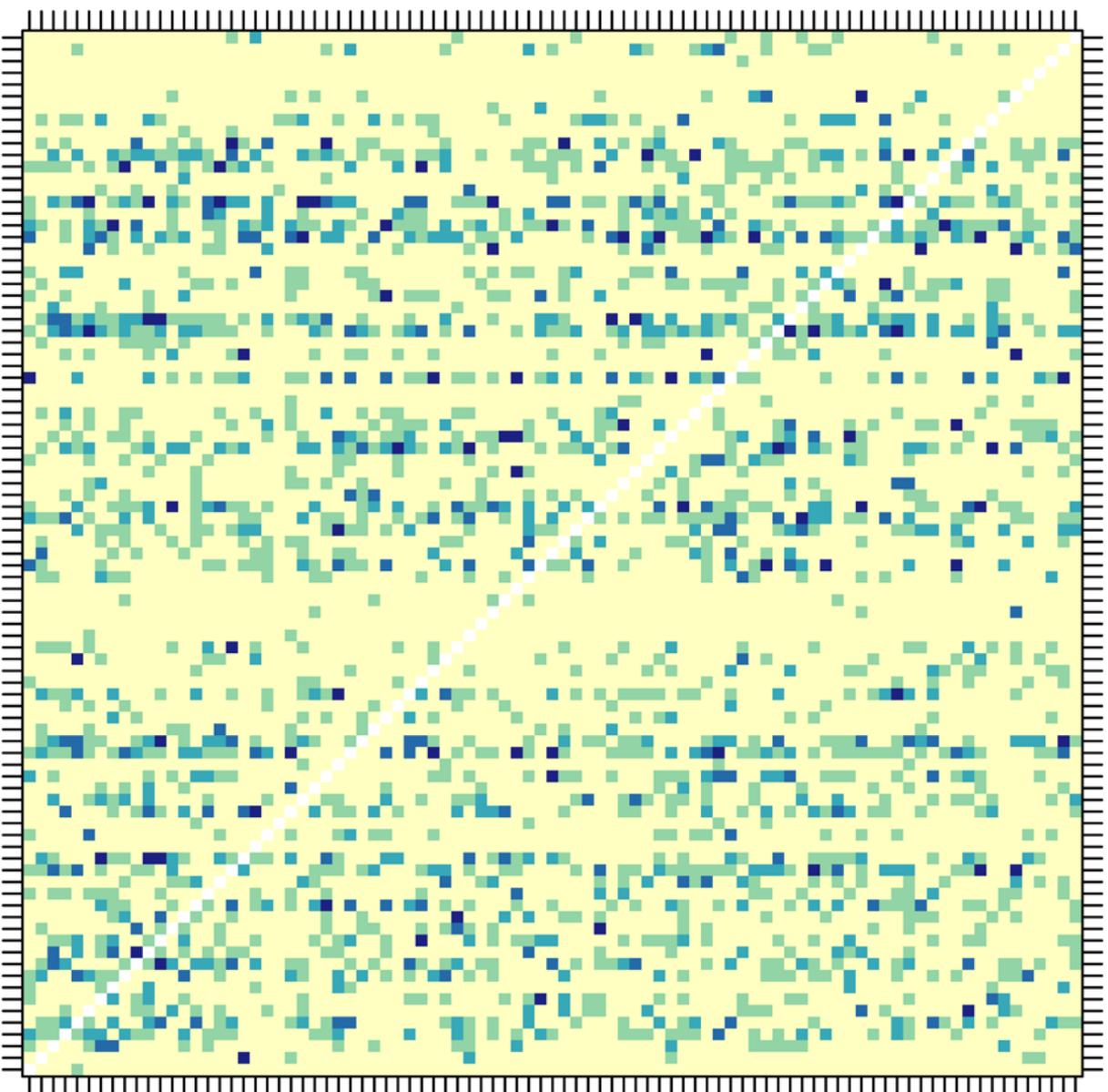
Slide by Kevin Dorn

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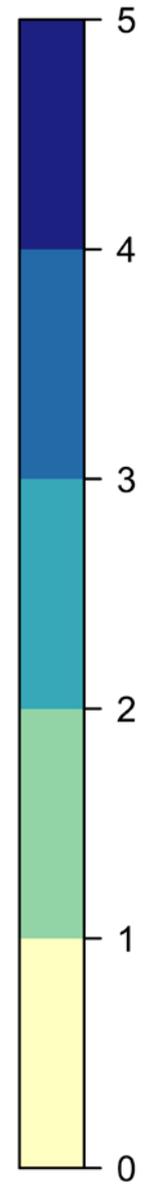
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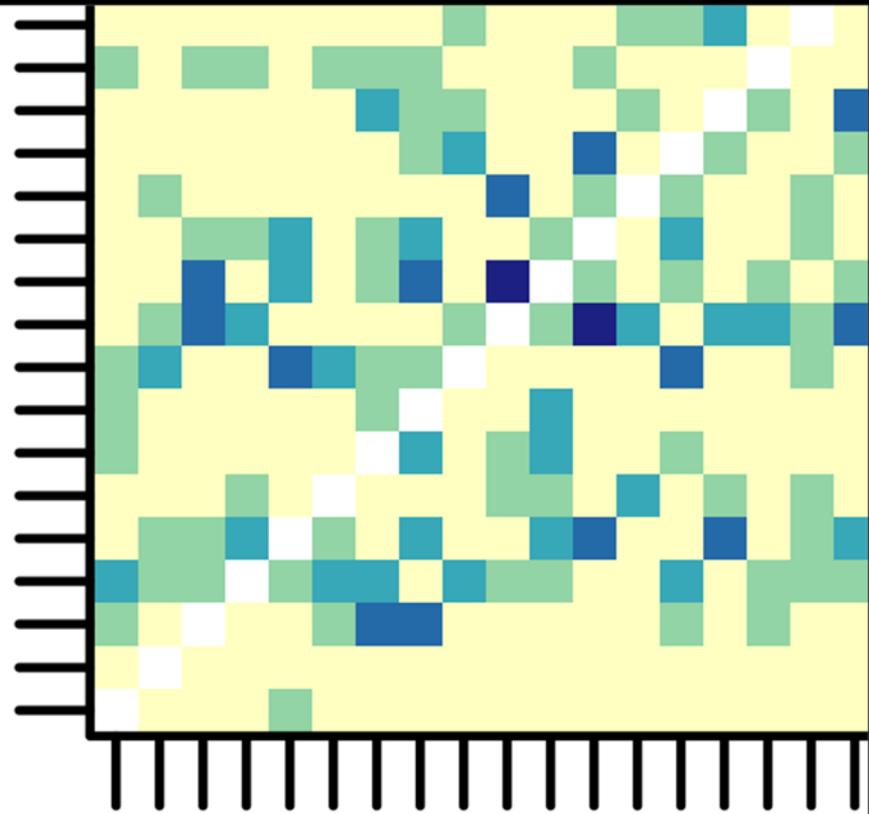
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Female ID



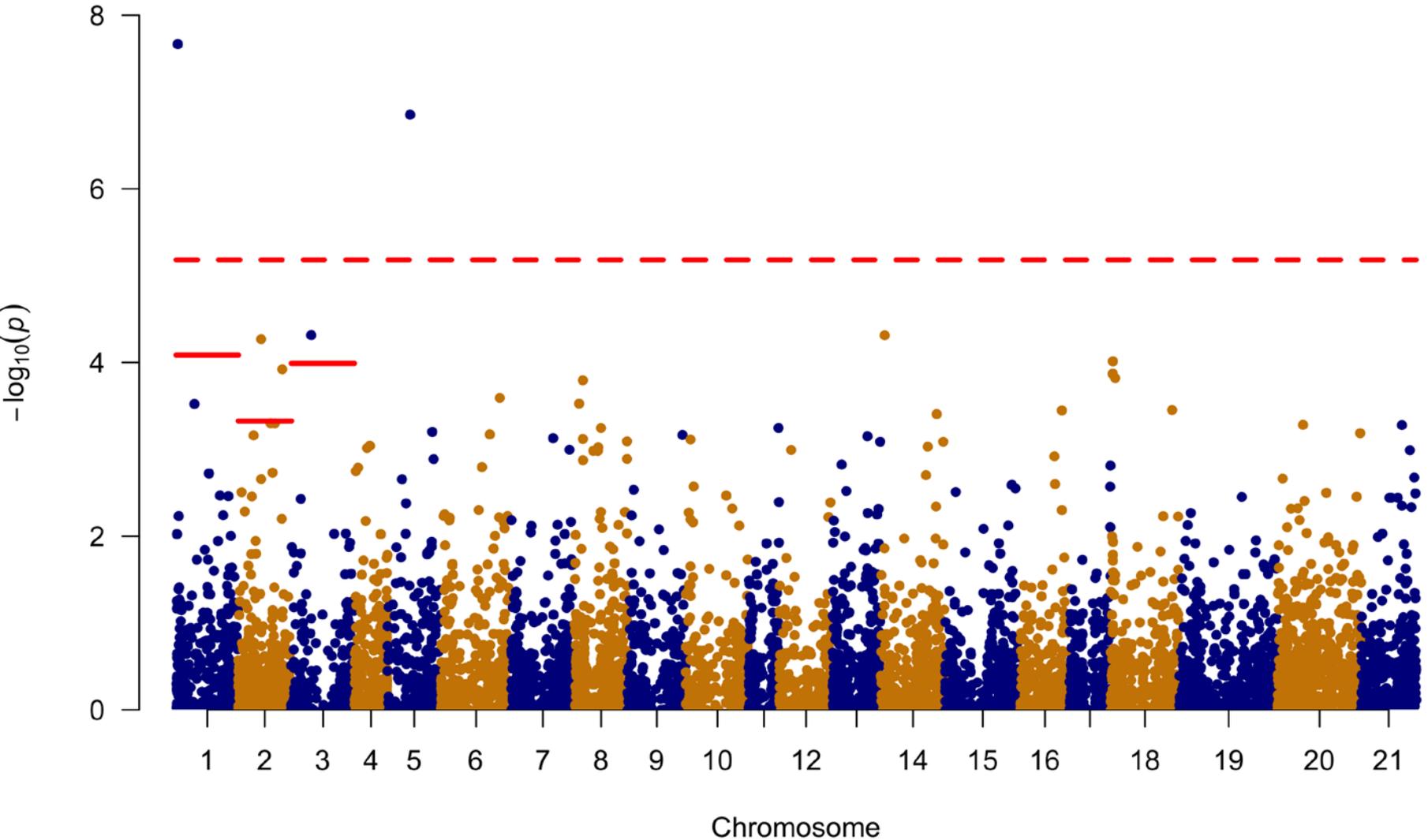
Number of progeny observed

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16SGH000001



16SGH000001  
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16SGH000018

# GWAS of Progeny Combinations



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United States  
Department of  
Agriculture

National Institute  
of Food and  
Agriculture



Forever  
Green



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