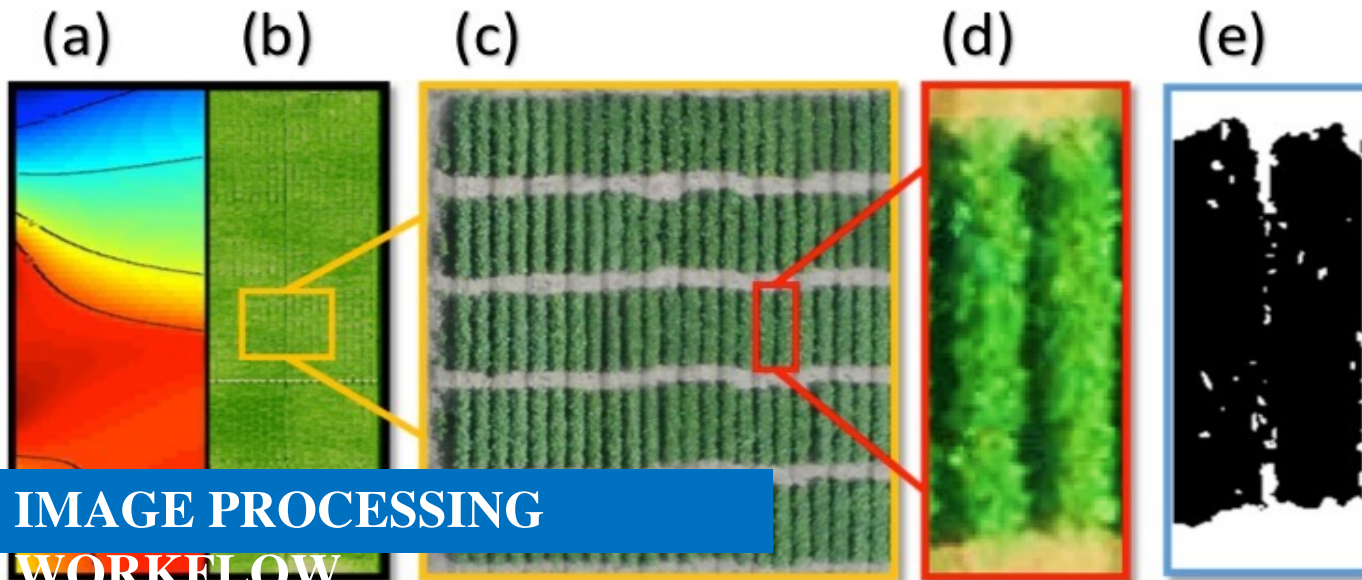


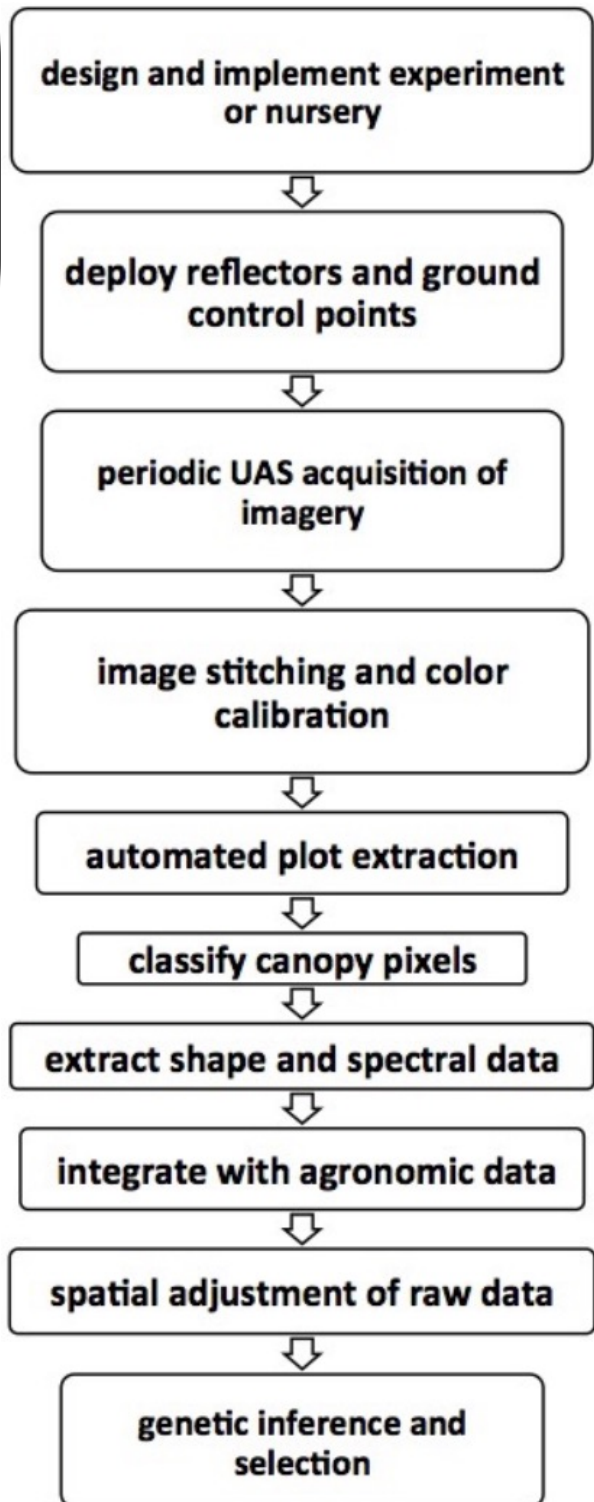
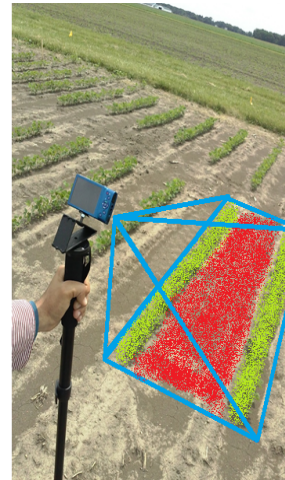
# Genetic Architecture of Phenomic-enabled Canopy Coverage in Soybean

*Alencar Xavier, Benjamin Hall, Anthony Hearst,  
Keith Cherkauer, Katy Martin Rainey  
Agronomy , Agricultural & Biological Engineering  
Purdue University*

# Soybean Canopy Phenotyping



(A) FIELD VARIATION (B) FIELD IMAGE (C) BLOCK OF PLOTS (D) INDIVIDUAL PLOT (E) CLASSIFIED PLOT.



# UAS Image Acquisition

Slide from A. Hearst

2013-2015:  
20,000 plots  
80,000 images  
SoyNAM primary target



Flight Dates



# Canopy Coverage and Light Interception

“Soybean canopy coverage and light interception measurements using digital imagery”

*Larry C. Purcell (2001)*

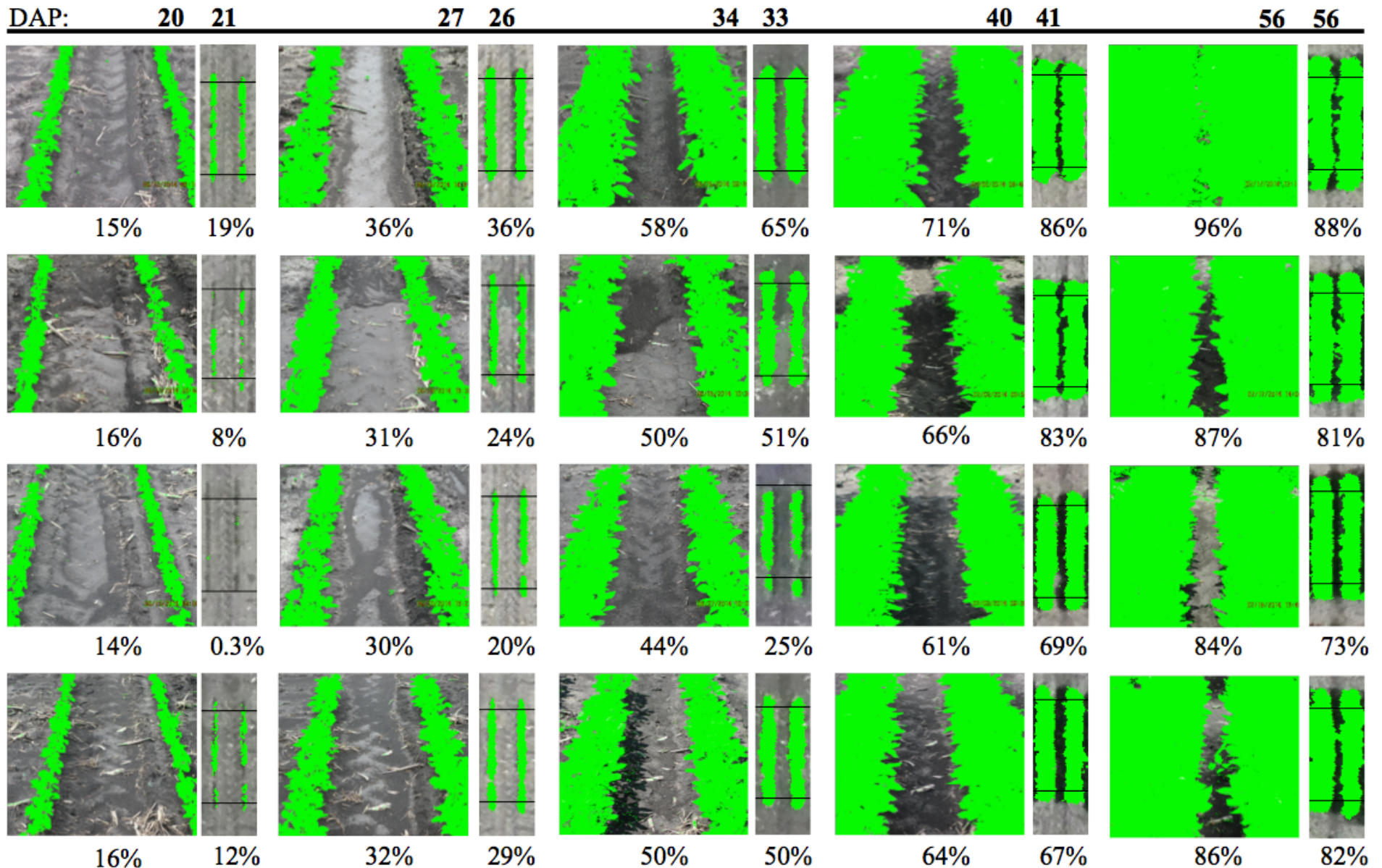
Crop Science, Vol. 40 No. 3.

Comparisons of canopy coverage values with LI measured near solar noon indicated that there was a one-to-one relationship



# Correspondence Between Ground & Air

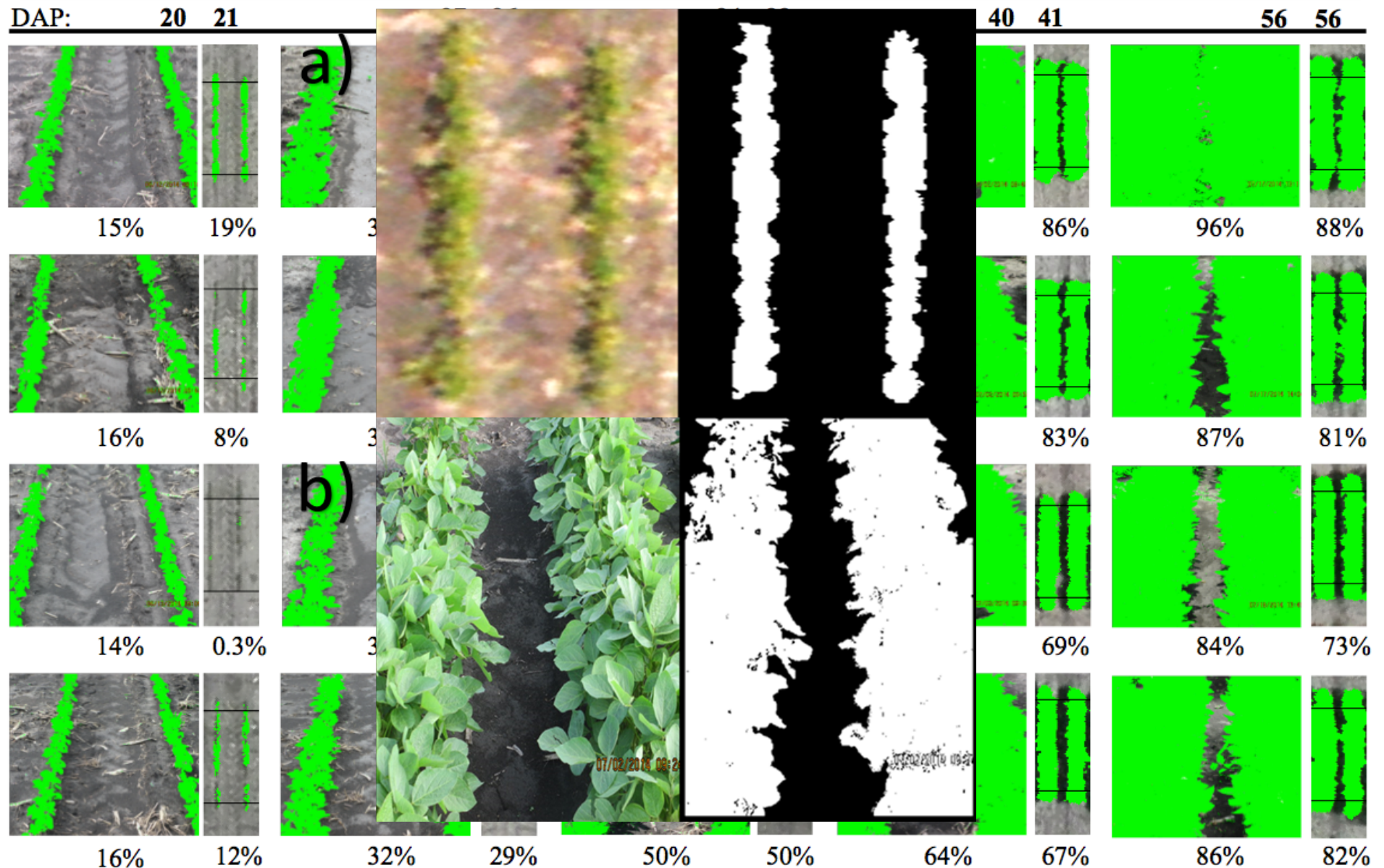
B. Hall (2015)





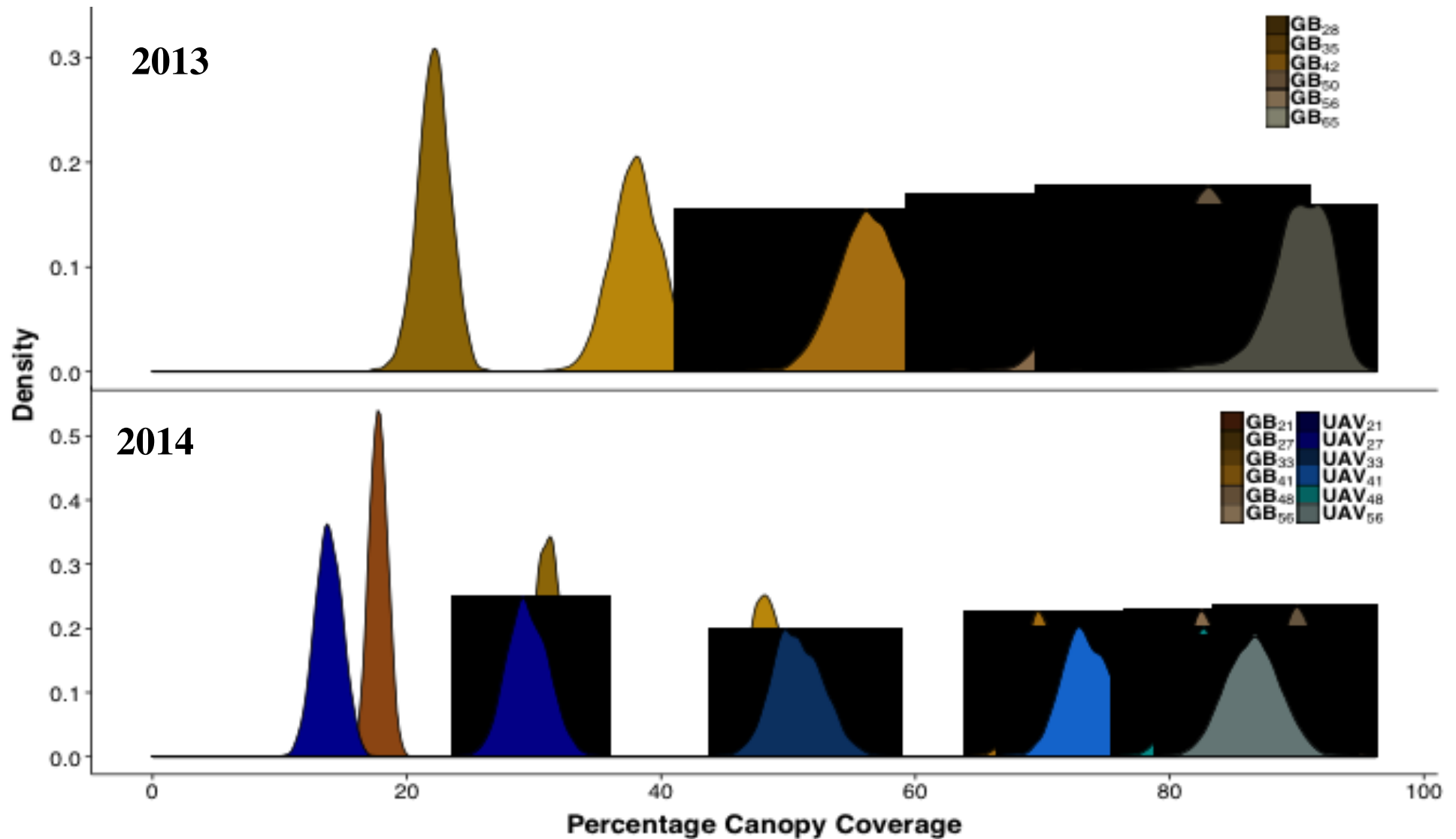
# Correspondence Between Ground & Air

B. Hall (2015)



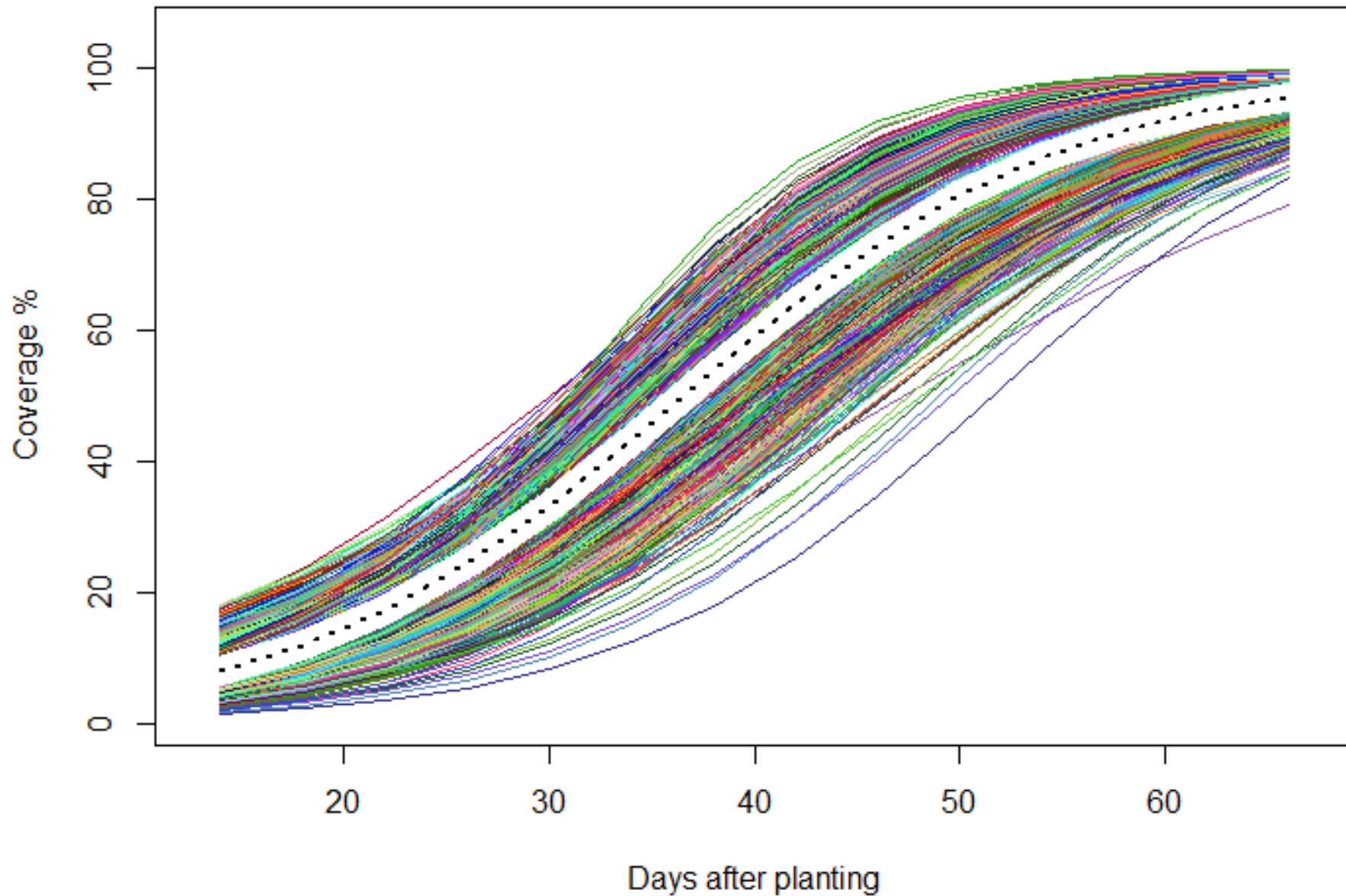
# Distribution of Seasonal Canopy Coverage in SoyNAM

B. Hall (2015)



# Canopy Development in SoyNAM

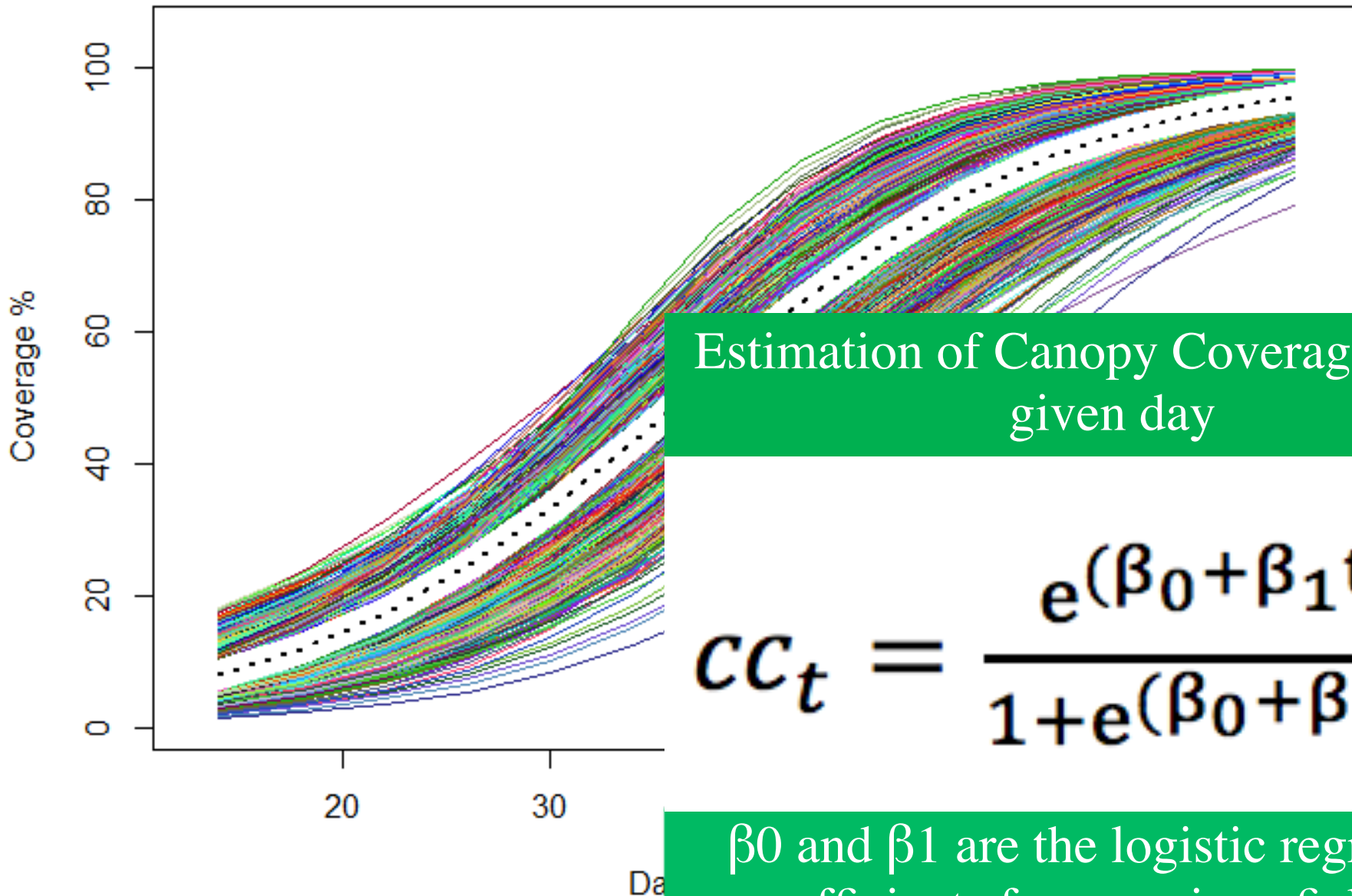
Xavier et al. (2017)





# Canopy Development in SoyNAM

Xavier et al. (2017)



$$CC_t = \frac{e^{(\beta_0 + \beta_1 t)}}{1 + e^{(\beta_0 + \beta_1 t)}}$$

$\beta_0$  and  $\beta_1$  are the logistic regression coefficients for any given field plot

# Calculation of Genotypic Values

Xavier et al. (2017)

$$y = 1\mu + f(x) + Zu + Wg + e$$

$y$  is the vector of observed phenotypes

$\mu$  is the intercept

$f(x)$  is a non-linear function that accounts for the spatial heterogeneity of field variation, where  $f(x)$  is computed as the average phenotypic value of neighbor plots

$Z$  is the incidence matrix of environment

$u$  is the vector of regression coefficients of environment effects

$W$  is the incidence matrix of genotypes

$g$  is the vector of genetic values

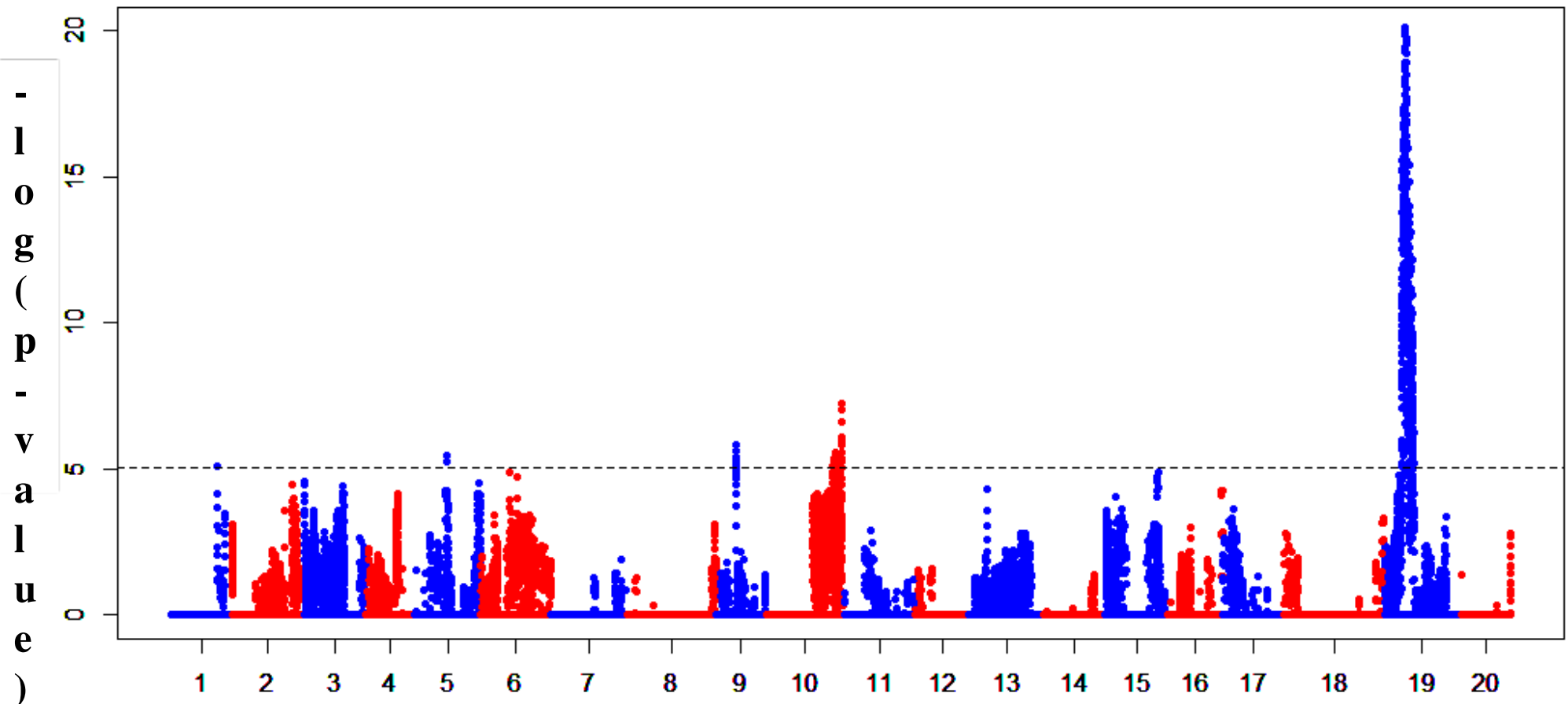
$e$  is the vector of residuals.

$$u \sim N(0, I\sigma_u^2) \quad g \sim N(0, I\sigma_g^2) \quad e \sim N(0, I\sigma_e^2).$$

# Genetic Architecture of CC

Xavier et al. (2017)

## GWAS Canopy Coverage



Empirical Bayes GWAS:

NAM: association studies in multiple populations.

A Xavier, S Xu, WM Muir, & KM Rainey (2015)

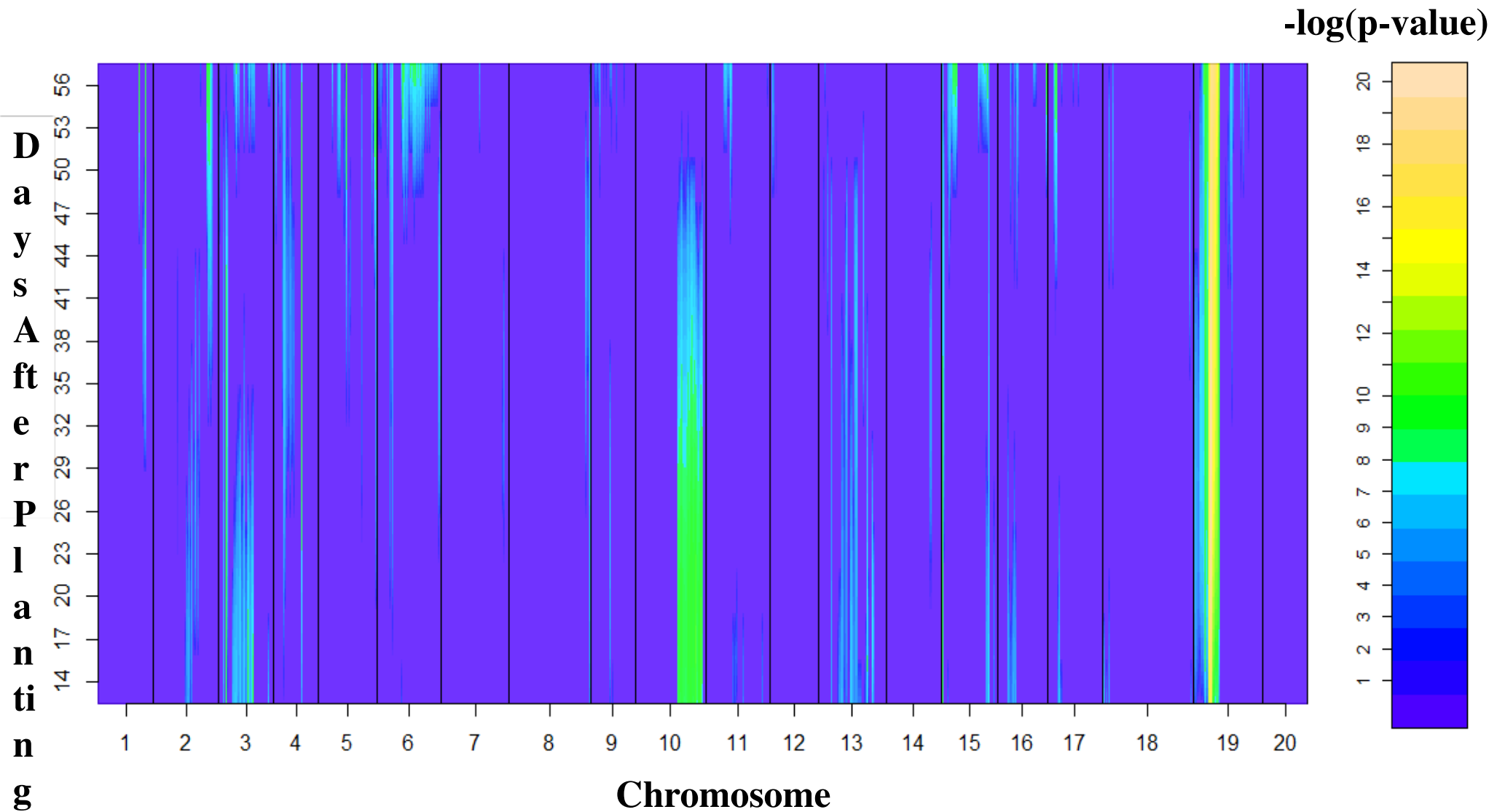
Bioinformatics.

**Chromosome**



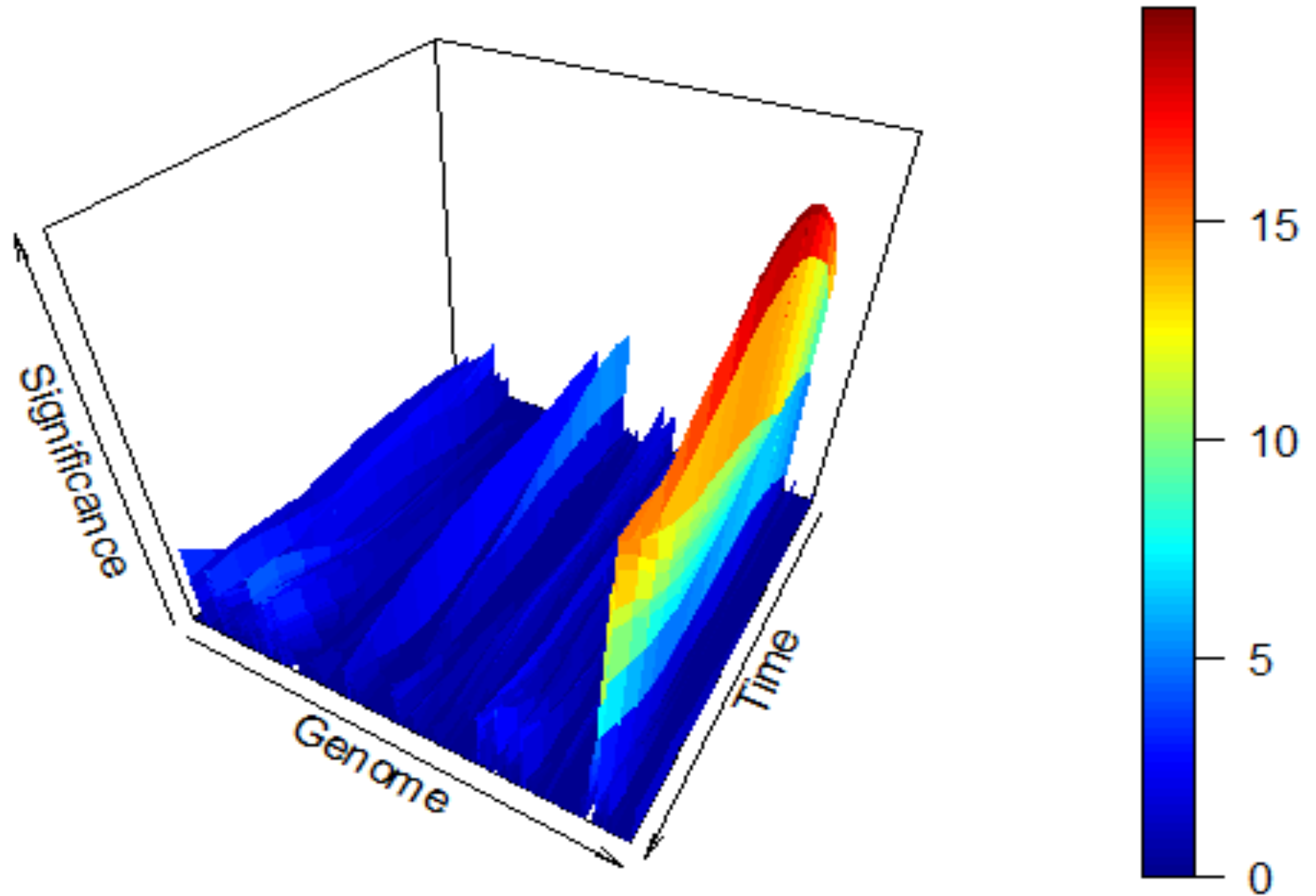
# Genetic Architecture of CC Over Time

Xavier et al. (2017)



# Genetic Architecture of CC Over Time

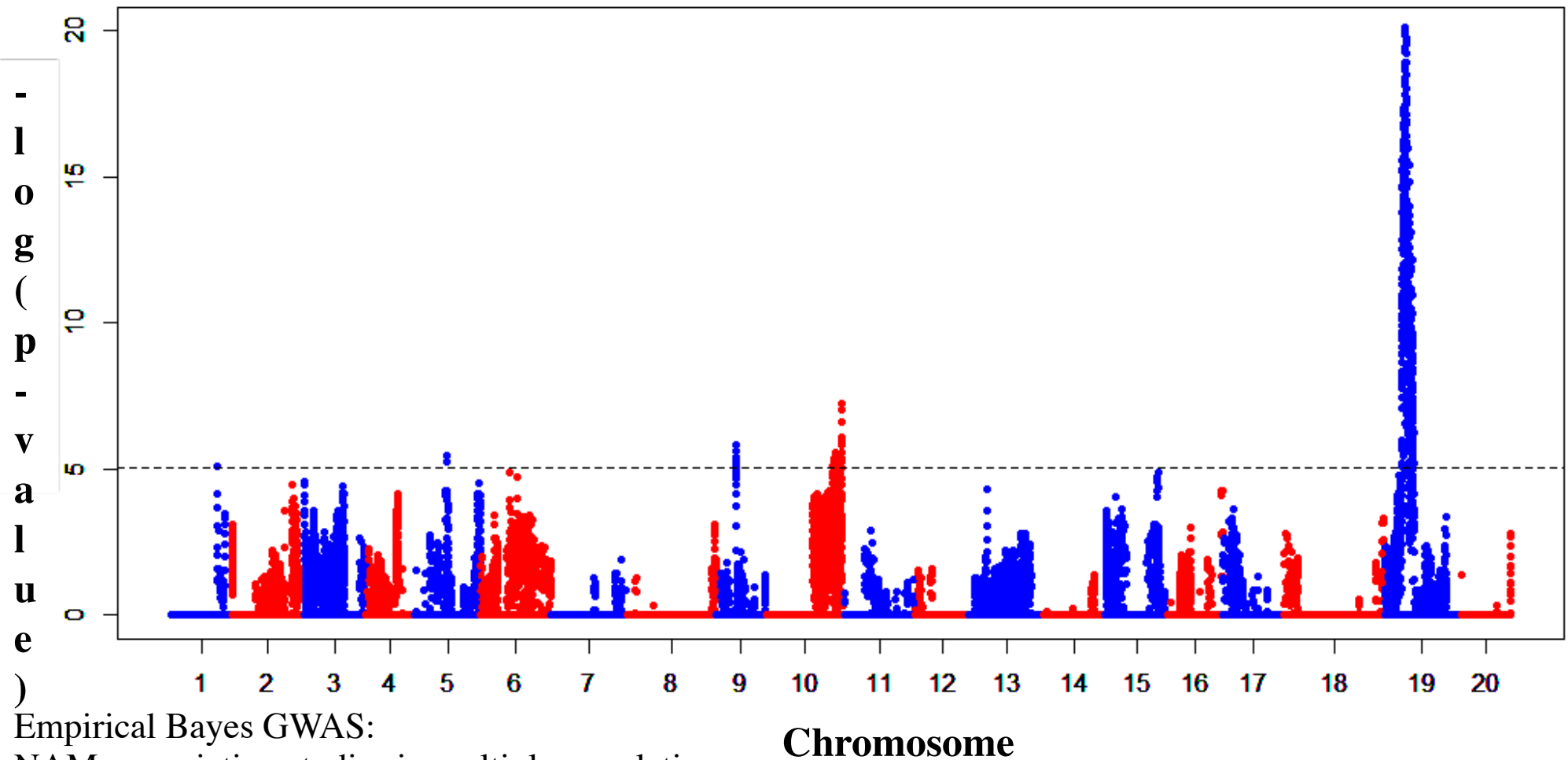
Slide from Alencar Xavier



# Genetic Architecture of CC

Xavier et al. (2017)

## GWAS Canopy Coverage



Empirical Bayes GWAS:

NAM: association studies in multiple populations.

A Xavier, S Xu, WM Muir, & KM Rainey (2015)

Bioinformatics.



# Average Canopy Coverage (ACC)

Xavier et al. (2017)

ACC value is an arithmetical mean of multiple seasonally observed values of canopy coverage.

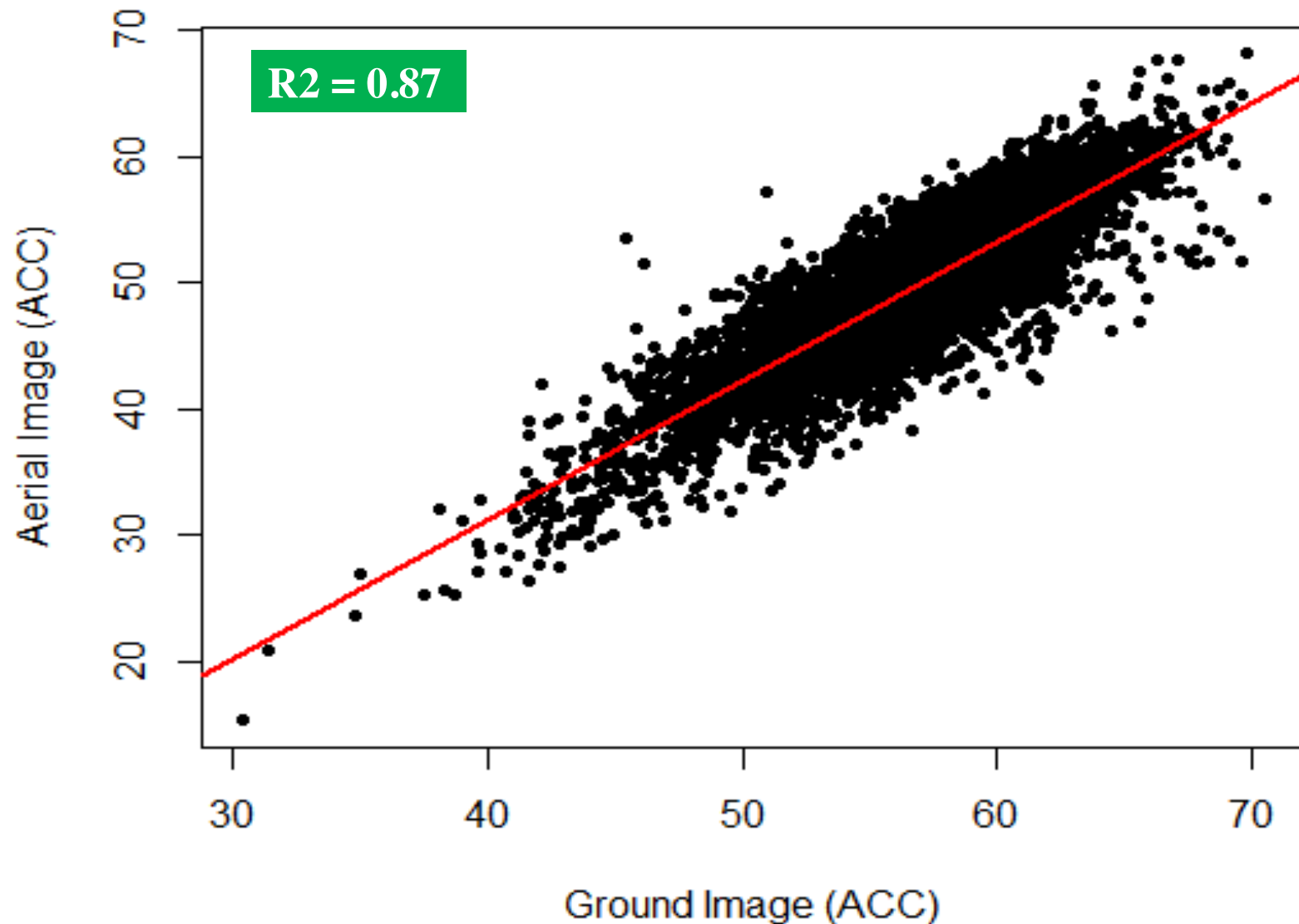
$$ACC = \int_{t_0}^{t_n} f(cc_t) \partial(cc_t) = \frac{1}{N} \sum_{t_0}^{t_n} cc_t$$

N = number of observed days

t<sub>0</sub> and t<sub>n</sub> = the first and last day when canopy coverage was assayed

# ACC: Correlation Between Ground & Air

Xavier et al. (2017)



# Canopy Coverage QTL

Xavier et al. (2017)

SNP	Period of Significant Association (DAP)	Allelic Effect		
		GY (kg.ha-1)	R8 (DAP)	ACC (%)
Gm01_50911939_C_T	56	-26.04	-1.12	-0.36
Gm05_37467797_A_G	53-56	-44.71	0.08	-0.65
Gm06_14104090_T_C	56	99.58	0.50	0.55
Gm09_4034850_C_T	14-35, 59-64	12.24	-0.60	-0.44
Gm010_44120764_T_ C	14-23	5.95	-0.59	0.04
Gm010_44630777_C_ A	14-26	51.61	1.06	0.33
Gm019_1586092_T_C	14-64	47.30	-0.24	1.34



# Quantitative Properties of ACC

Xavier et al. (2017)

Genetic (G) and environmental (E) variances and heritabilities for grain yield (GY), average canopy coverage (ACC) and days to maturity (R8).

	GY	R8	ACC
Var (G)	78.80	208.36	25.54
Var (E)	56.12	9.39	8.10

Phenotypic (P), Spearman (S), genetic (G) and environmental (E) correlations among grain yield (GY), average canopy coverage (ACC) and days to maturity (R8).

	GY-ACC	R8-ACC	GY-R8
Cor(P)	0.63	0.31	0.42
Cor(S)	0.70	0.38	0.46
Cor(G)	0.88	0.77	0.72
Cor(E)	0.18	-0.06	0.23

# Quantitative Properties of ACC

Xavier et al. (2017)

Genetic (G) and environmental (E) variances and heritabilities for grain yield (GY), average canopy coverage (ACC) and days to maturity (R8).

	GY	R8	ACC
Var (G)	78.80	208.36	25.54
Var (E)	56.12	9.39	8.10

Phenotypic (P), Spearman (S), genetic (G) and environmental (E) correlations among grain yield (GY), average canopy coverage (ACC) and days to maturity (R8).

	GY-ACC	R8-ACC	GY-R8
Cor(P)	0.63	0.31	0.42
Cor(S)	0.70	0.38	0.46
Cor(G)	0.88	0.77	0.72
Cor(E)	0.18	-0.06	0.23

**POSTER SESSION: FABIANA MOREIRA**

# Acknowledgements

## Three graduate students projects:

1. **Alencar Xavier**, Quantitative Genetics, Dow AgroSciences, **Analyses**
2. **Ben Hall**, Ph.D. student Soybean Agronomy, Purdue, **Canopy Traits from M.S.**
3. **Anthony Hearst**, Ph.D. student, NSF Fellow, Purdue Agricultural & Biological Engineering, **UAS Flights + Image Analysis**, advised by **Dr. Keith Cherkauer**

**Chris Hoagland** and **Curtis Brackett** (Rainey Lab Technical Support)

**Bill Muir**, **Shizhong Xu** and **Bill Beavis** (Quantitative Genetics)

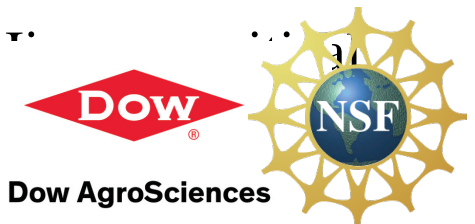
**Brian Diers**, **Jim Specht**, **Bill Beavis** (SoyNAM Developers & reviewer)

**Qijian Song** and **Perry Cregan** (Genotypic Data)

**Concept of Canopy Coverage**

**Shaun Casteel**, Purdue Soybean Agronomy

**Larry Purcell**, Arkansas Soybean Agronomy + consultation



# PHENOMICS WORKSHOP AT PURDUE



**Module-based instruction**



**Expert panel discussion**



**Networking opportunities**

**March 13 - 14, 2017**

**\$500 registration fee**

Open to first 30 professionals

Lunch provided

Accommodations and logistics coming soon

[ag.purdue.edu/plantsciences/phenomics-curricula](http://ag.purdue.edu/plantsciences/phenomics-curricula)

Instruction, demonstration and discussion  
to enable effective incorporation of  
**phenomic tools** and **approaches** into  
plot-based crop research, and to **educate**  
engineers on relevant agricultural and  
scientific topics.

[krainey@purdue.edu](mailto:krainey@purdue.edu)

# Lack of Variation in Historical Cultivars

Historical gains in soybean (*Glycine max* Merr.) seed yield are driven by linear increases in light interception, energy conversion, and partitioning efficiencies.

Koester *et al.*

J Exp Bot. 2014 65(1).

-there was difference in the rate of canopy closure in older or newer cultivars, and most cultivars approached 90% closure by ~60 d after planting

-the time to canopy closure did not change in historical soybean cultivars

