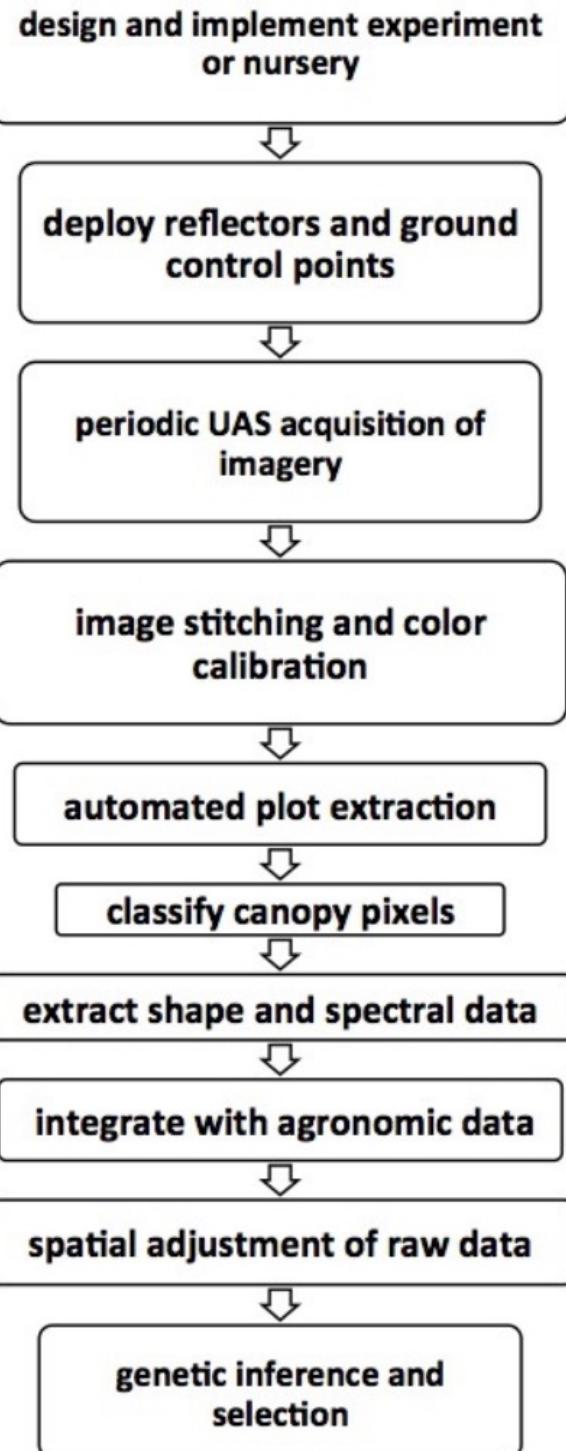
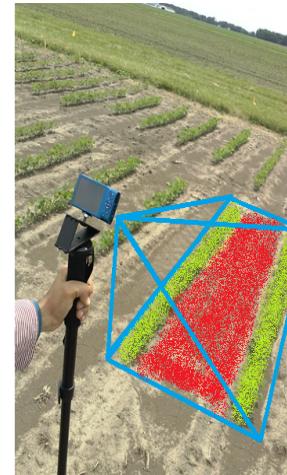
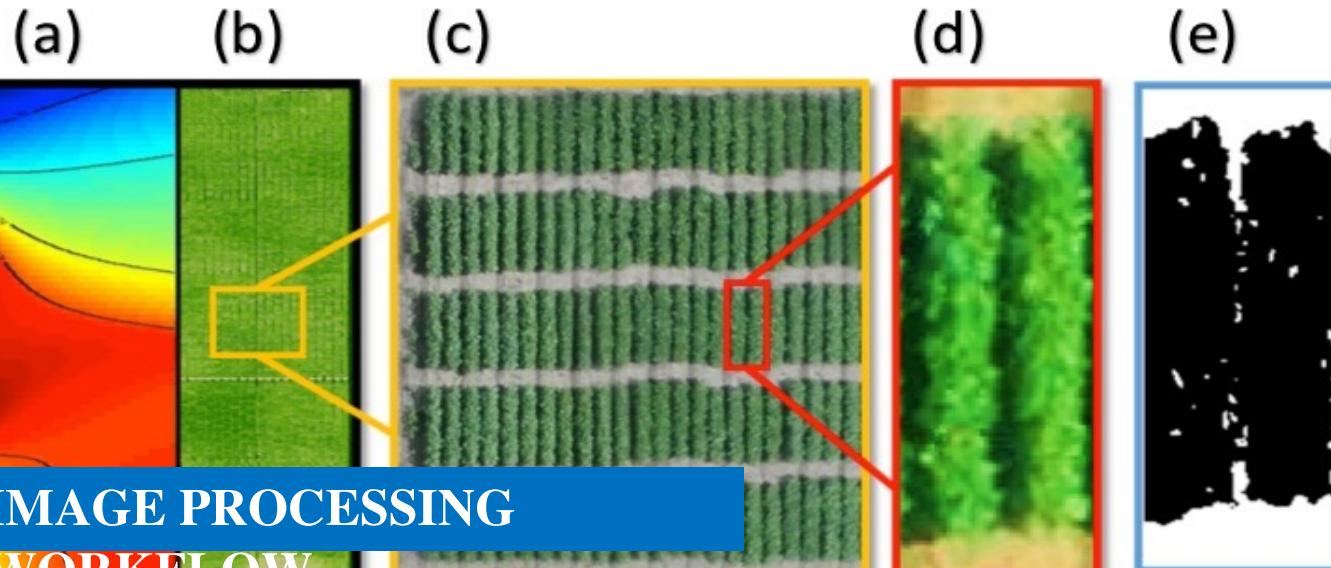


Genetic Architecture of Phenomic-enabled Canopy Coverage in Soybean

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

Soybean Canopy Phenotyping



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)

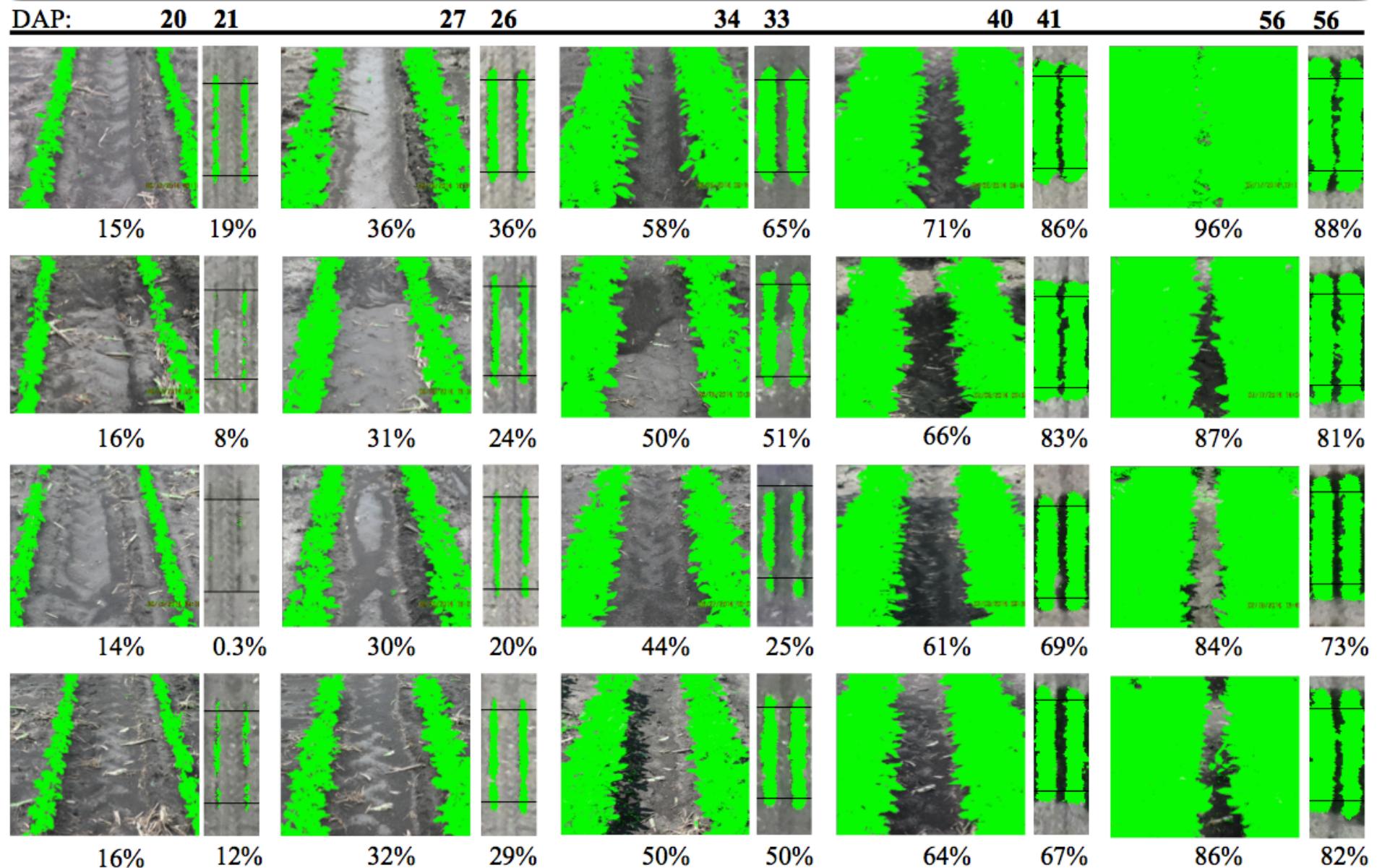
Crop Science, Vol. 40 No. 3.

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



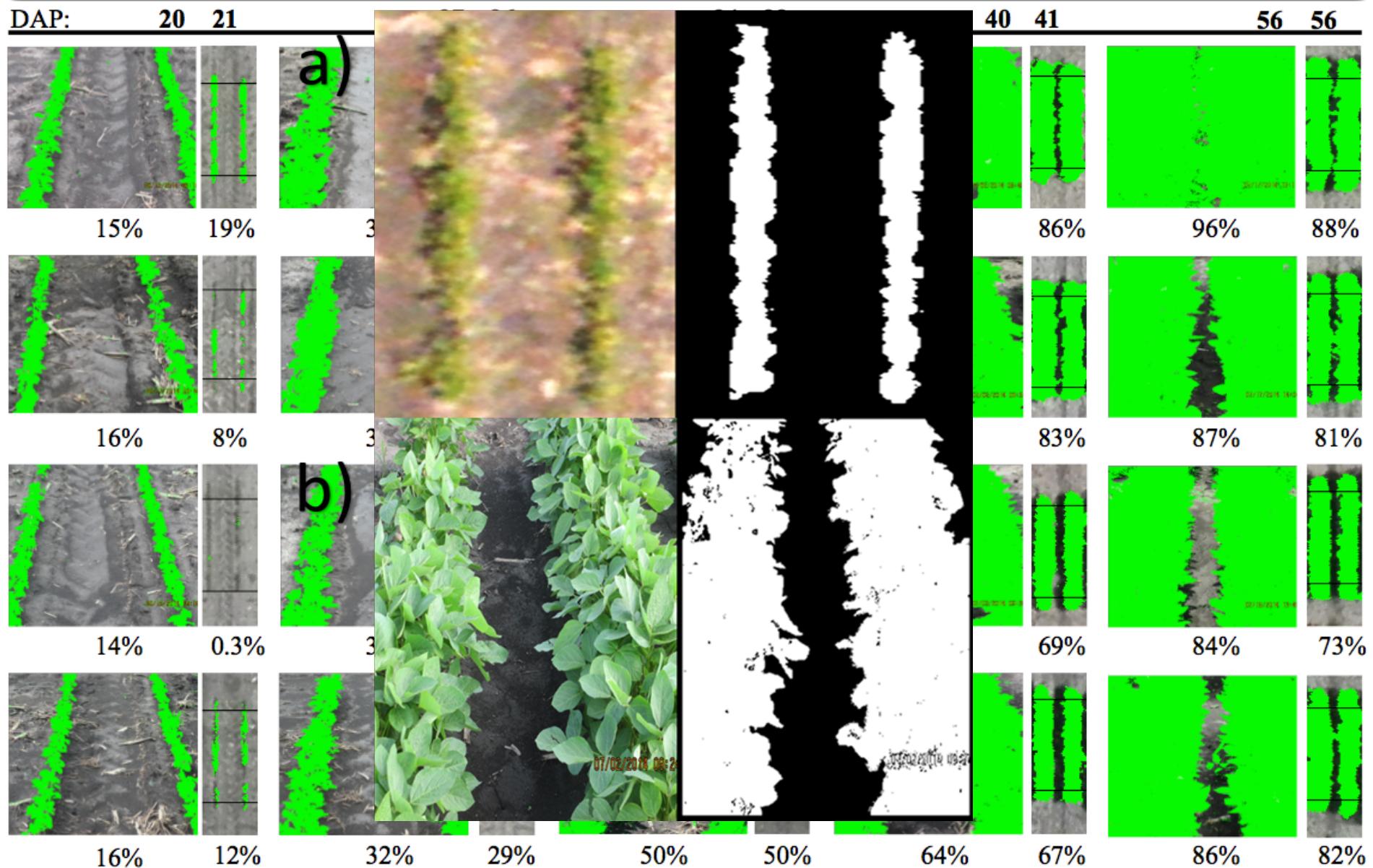
Correspondence Between Ground & Air

B. Hall (2015)



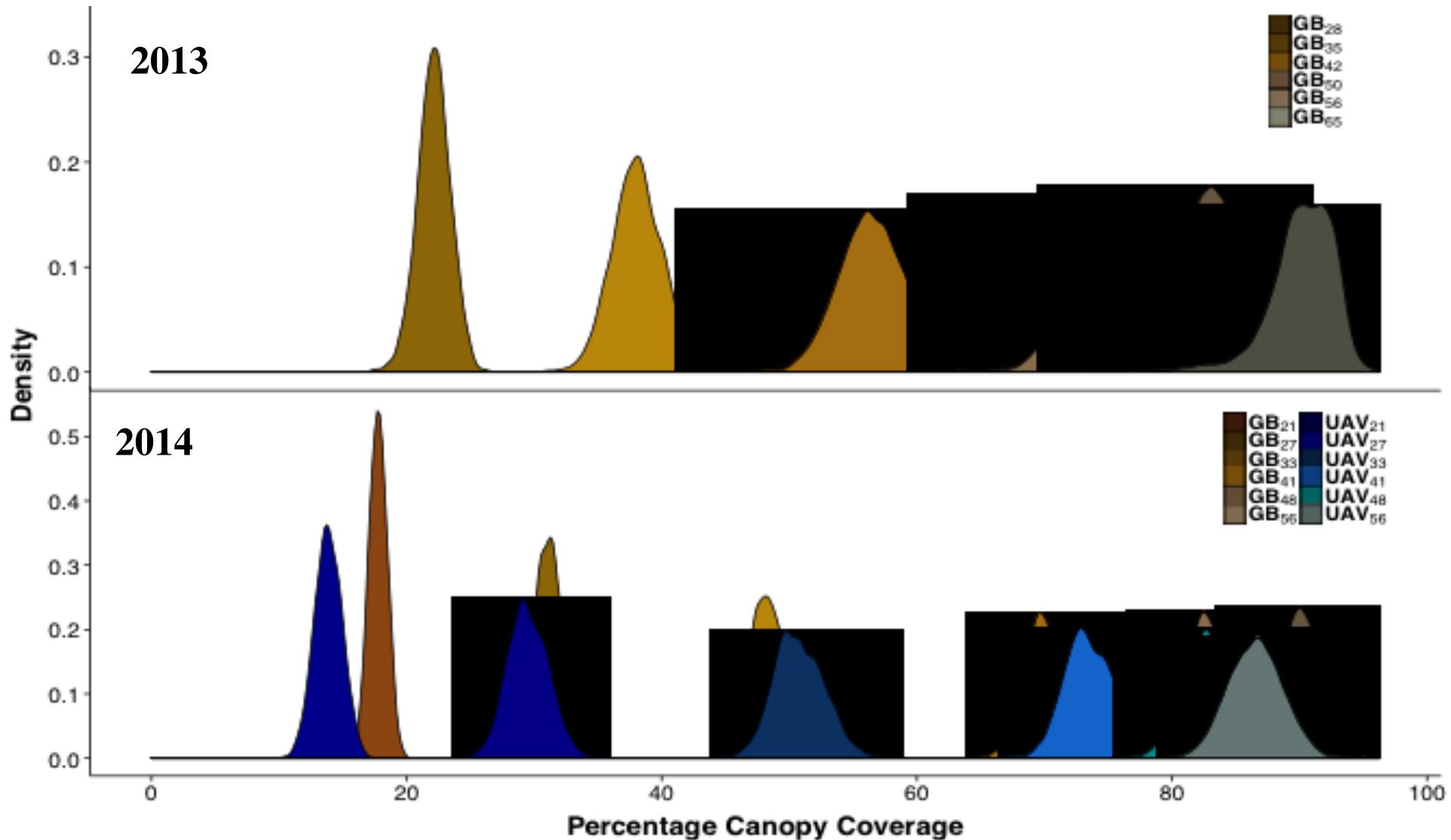
Correspondence Between Ground & Air

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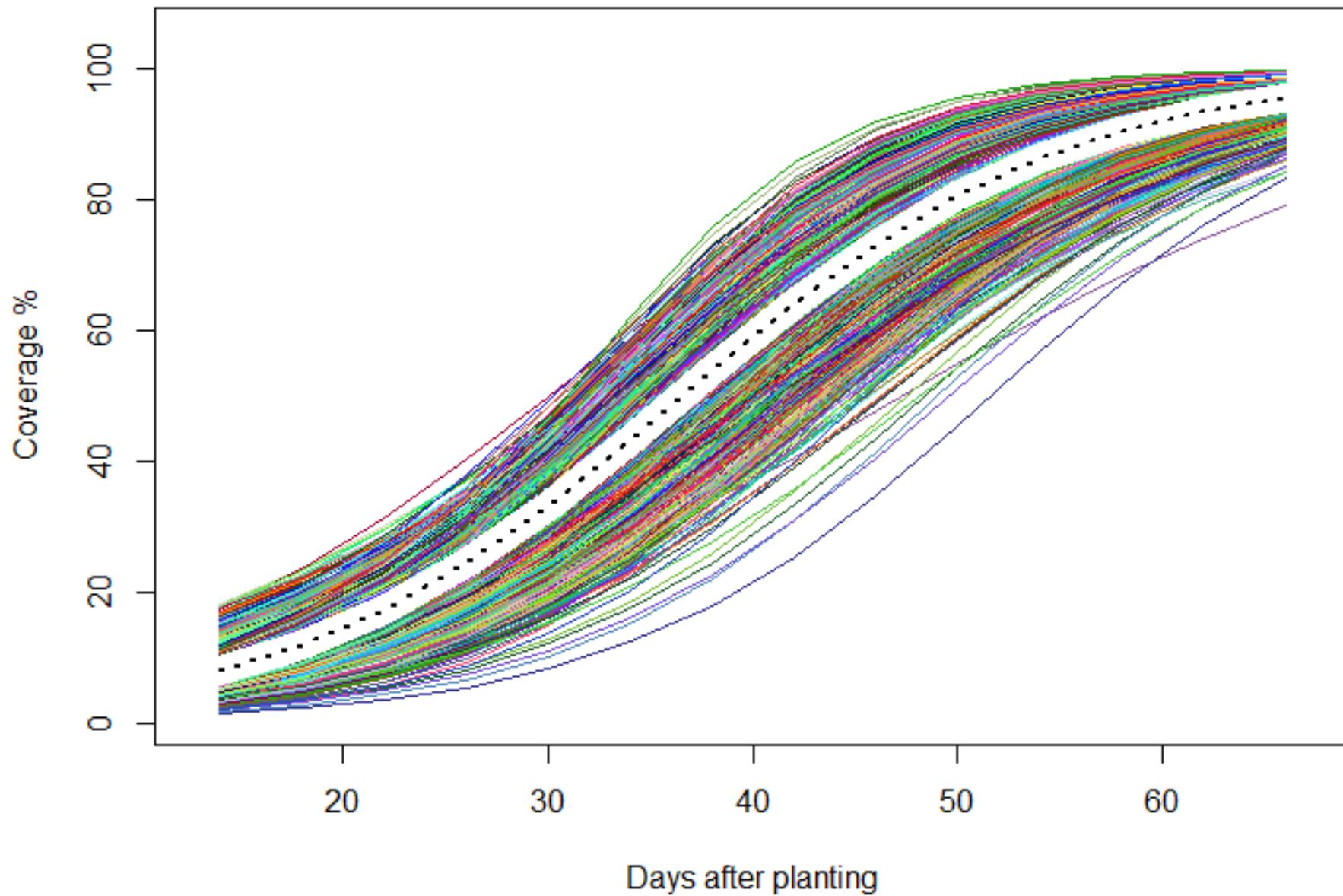
Distribution of Seasonal Canopy Coverage in SoyNAM

B. Hall (2015)



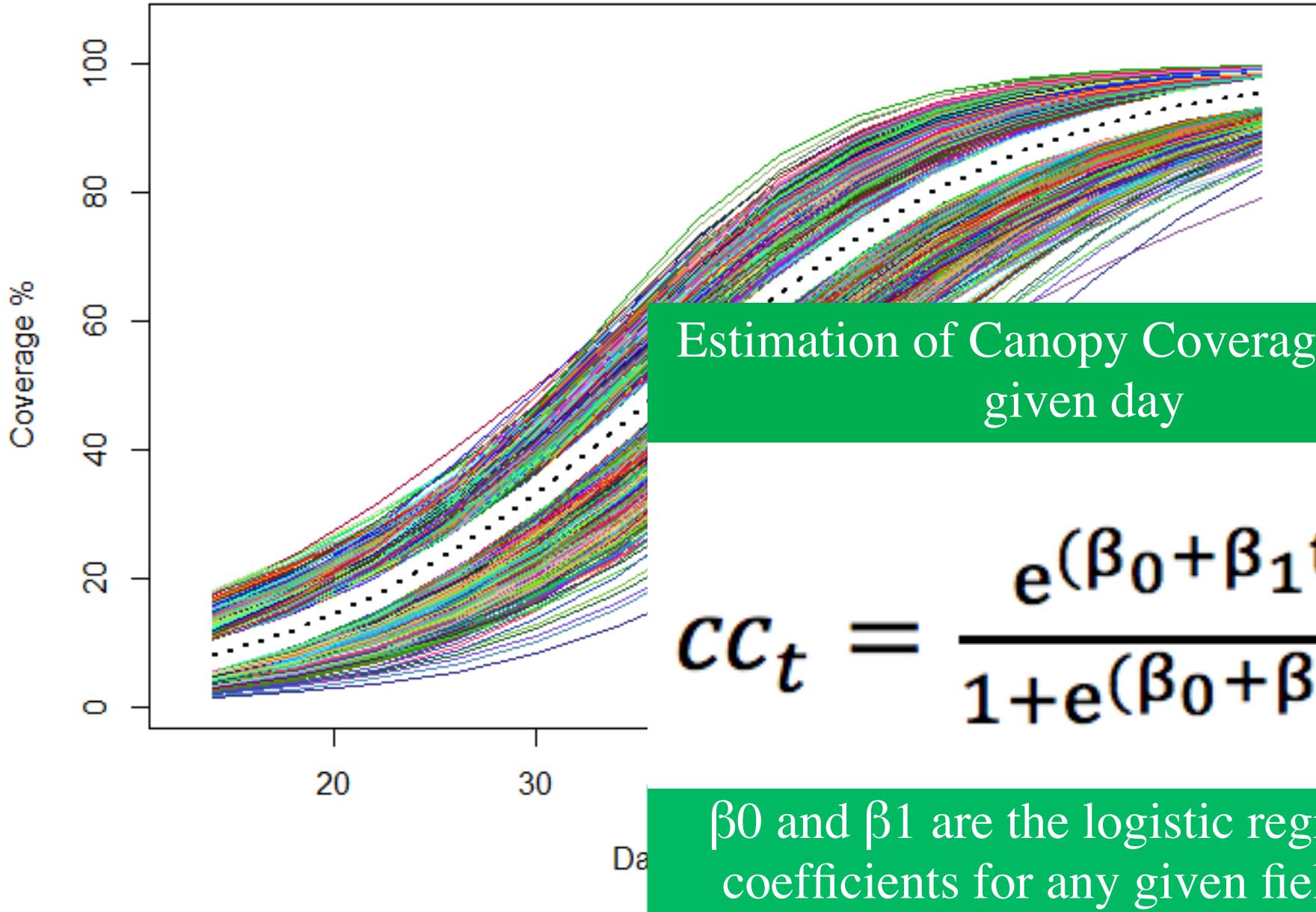
Canopy Development in SoyNAM

Xavier et al. (2017)



Canopy Development in SoyNAM

Xavier et al. (2017)



Calculation of Genotypic Values

Xavier et al. (2017)

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

y is the vector of observed phenotypes

μ 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

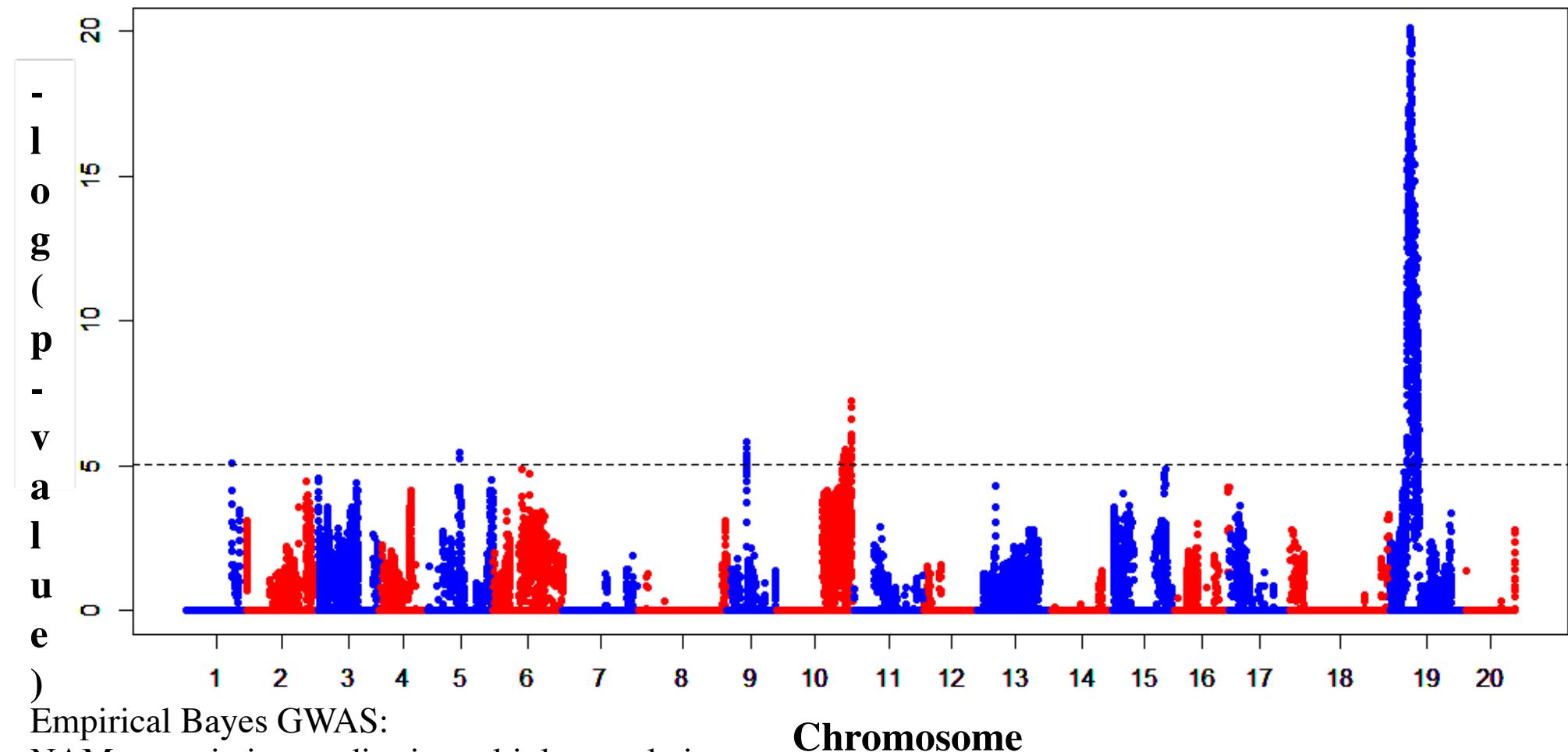
e is the vector of residuals.

$$u \sim N(0, I\sigma^2_u) \quad g \sim N(0, I\sigma^2_g) \quad e \sim N(0, I\sigma^2_e).$$

Genetic Architecture of CC

Xavier et al. (2017)

GWAS Canopy Coverage

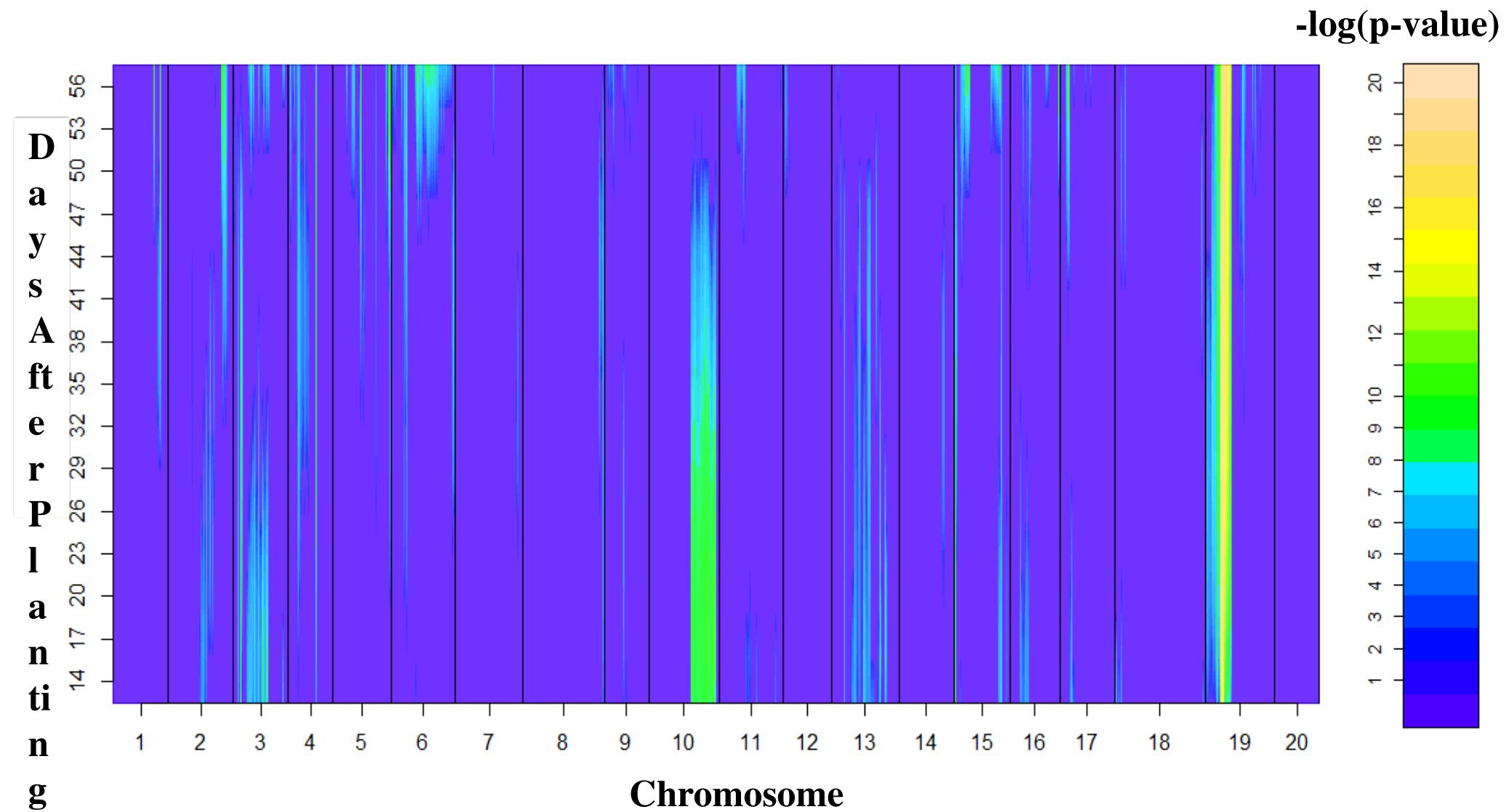


Empirical Bayes GWAS:

NAM: association studies in multiple populations.
A Xavier, S Xu, WM Muir, & KM Rainey (2015)
Bioinformatics.

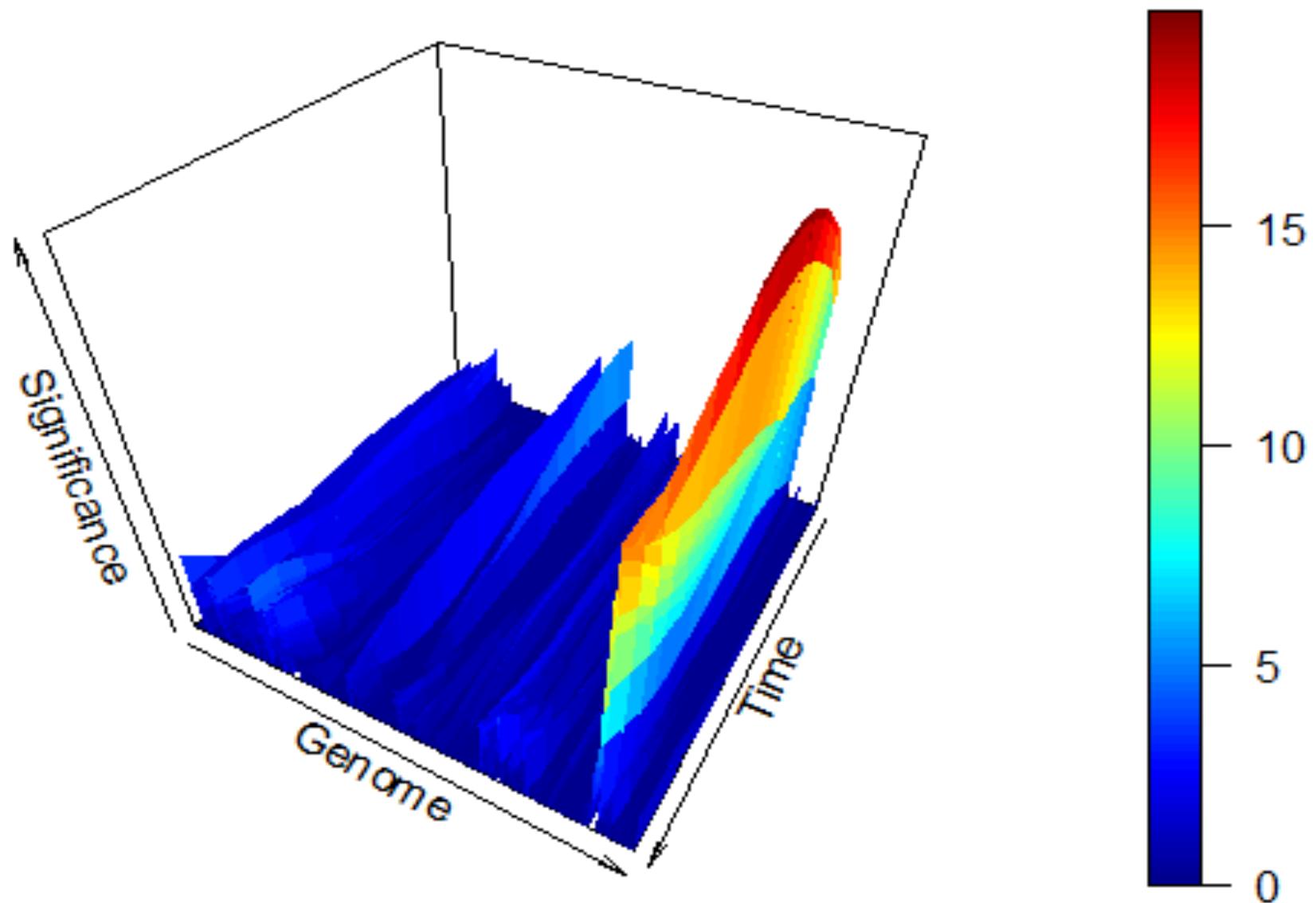
Genetic Architecture of CC Over Time

Xavier et al. (2017)



Genetic Architecture of CC Over Time

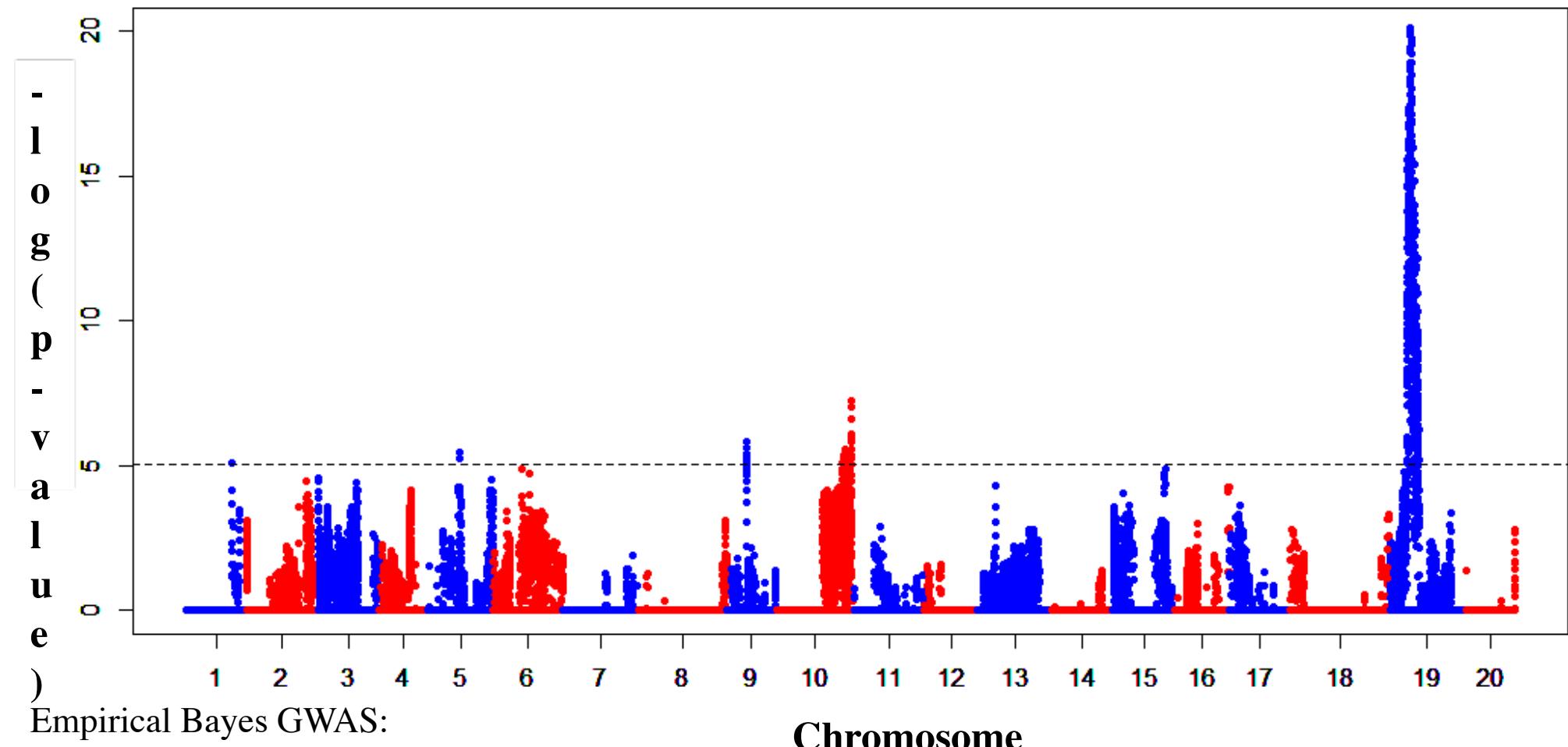
Slide from Alencar Xavier



Genetic Architecture of CC

Xavier et al. (2017)

GWAS Canopy Coverage



Empirical Bayes GWAS:

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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.

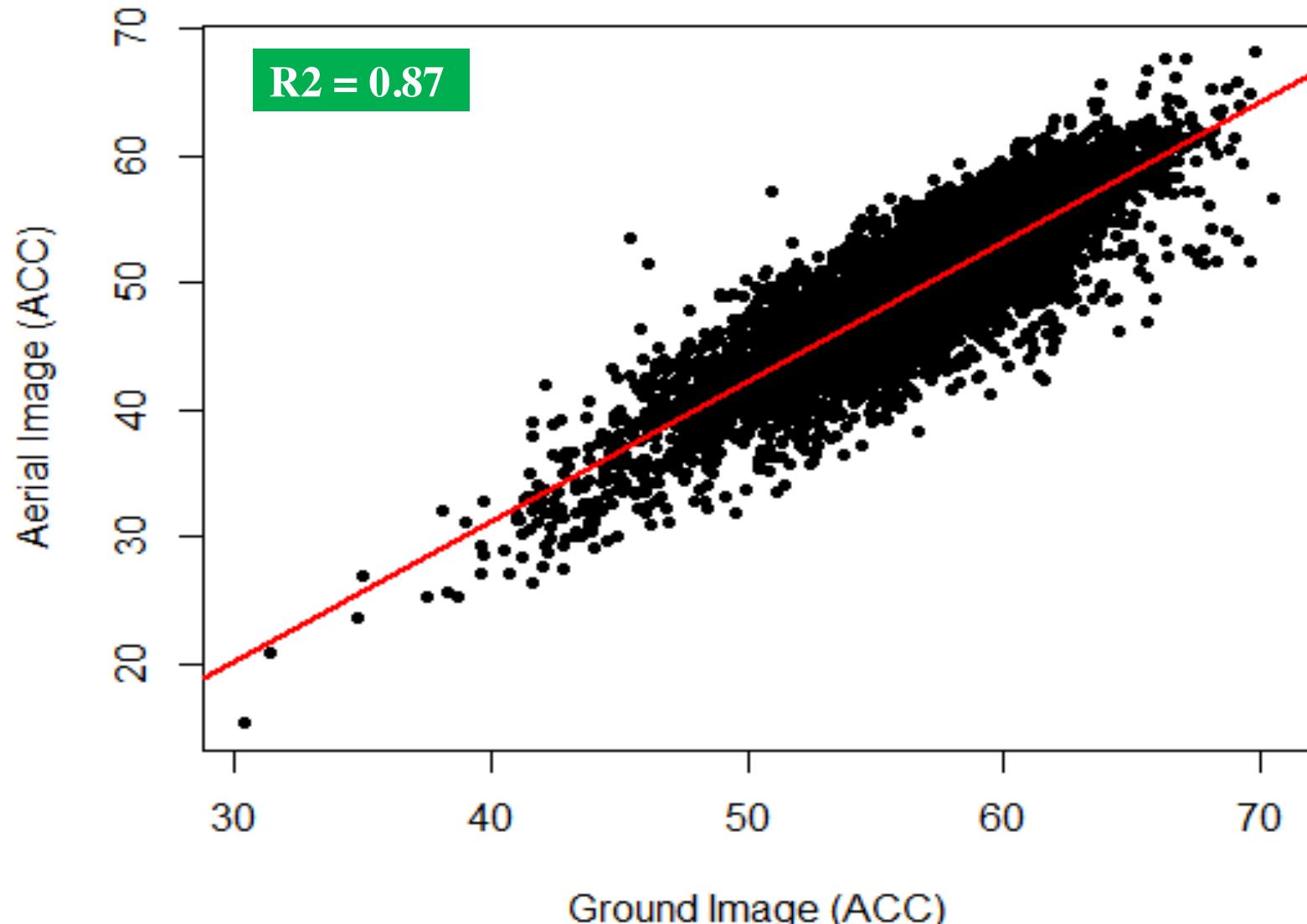
$$ACC = \int_{t_0}^{t_n} f(cc_t) \partial(cc_t) = \frac{1}{N} \sum_{t_0}^{t_n} cc_t$$

N = number of observed days

t0 and tn = 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)	GY (kg.ha ⁻¹)	Allelic Effect 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).

Phenotypic (P), Spearman (S), genetic (G) and environmental (E) correlations among 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

	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

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Xavier et al. (2017)

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POSTER SESSION: FABIANA MOREIRA

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1. **Alencar Xavier**, Quantitative Genetics, Dow AgroSciences, **Analyses**
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3. **Anthony Hearst**, Ph.D. student, NSF Fellow, Purdue Agricultural & Biological Engineering, **UAS Flights + Image Analysis**, advised by **Dr. Keith Cherkauer**

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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



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Expert panel discussion



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March 13 - 14, 2017

\$500 registration fee

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Lunch provided

Accomodations and logistics coming soon

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

Lack of Variation in Historical Cultivars

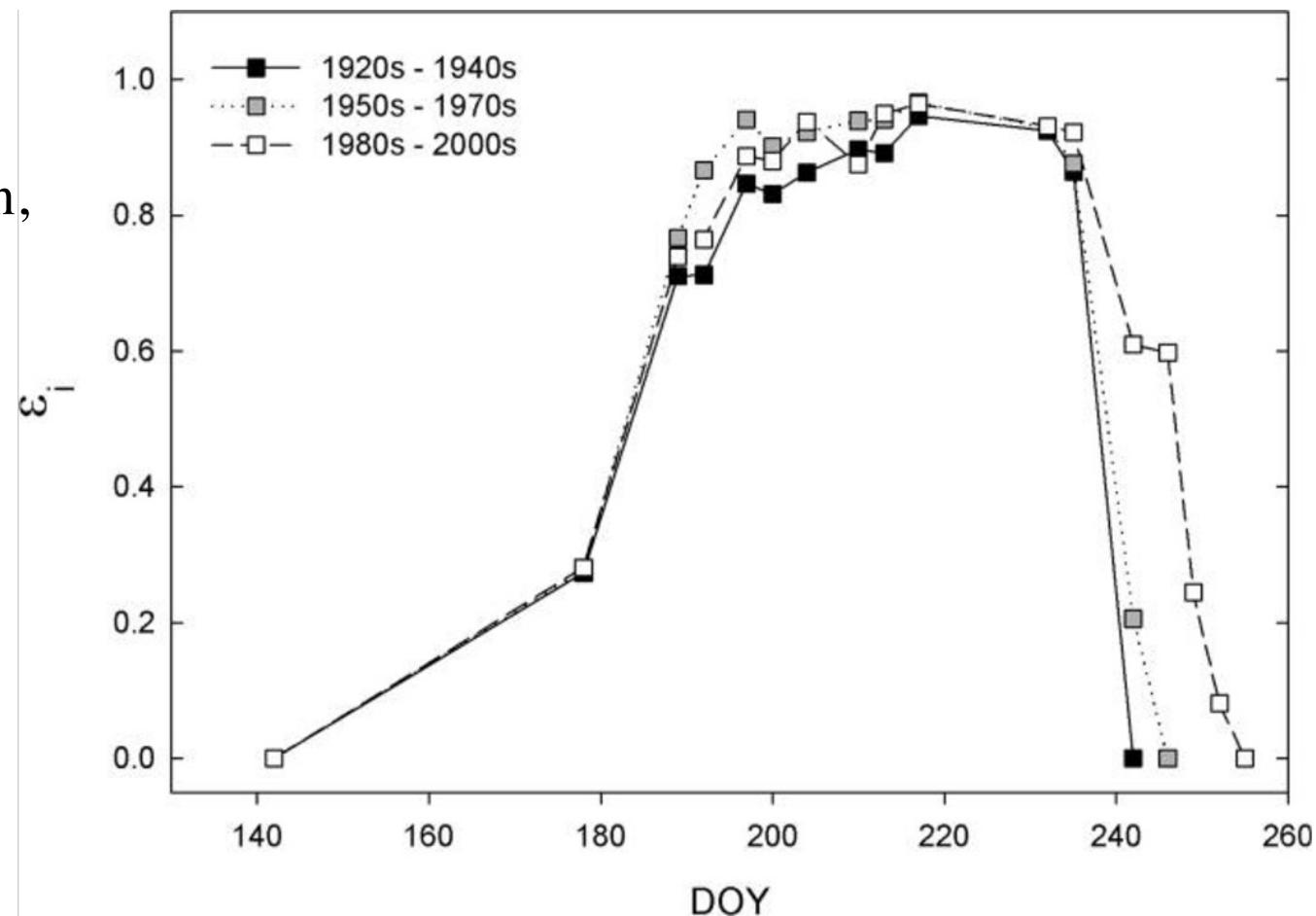
Historical gains in soybean (*Glycine max* Merr.) seed yield are driven by linear increases in light interception, energy conversion, and partitioning efficiencies.

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



Interception efficiency (ϵ_i) across the growing season in 2013 for each of the 24 soybean cultivars grouped by year of release