

The logo consists of the letters 'UF' in white, bold, sans-serif font, centered within an orange square. The background of the slide is a blue-tinted photograph of the University of Florida campus, featuring the prominent Spire tower and other Gothic-style buildings. A faint, light blue molecular or network diagram is overlaid on the right side of the image.

UF

Big data applications in strawberry breeding

Zhen Fan, PhD, Postdoctoral Associate

Co-authors: Vance Whitaker, Sujeet Verma, Luis Osorio, Ronald Tapia, Cheryl Dalid

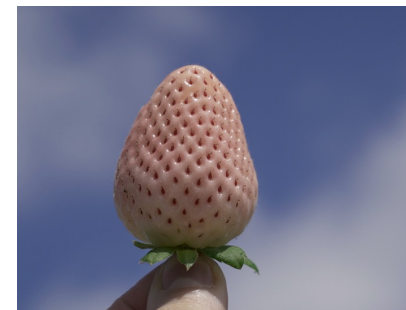
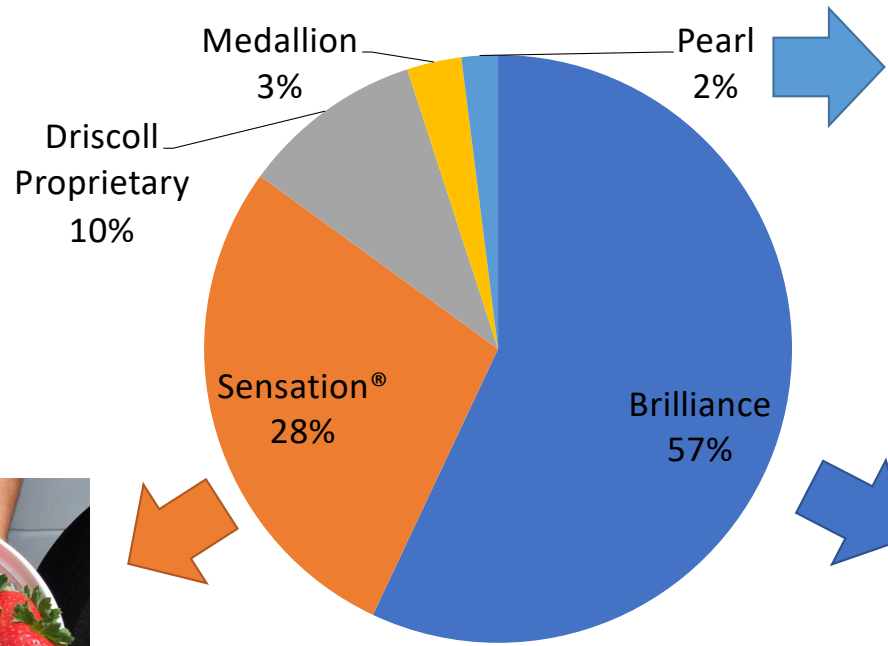
Horticultural Sciences Department

Gulf Coast Research and Education Center

University of Florida



The UF strawberry breeding program



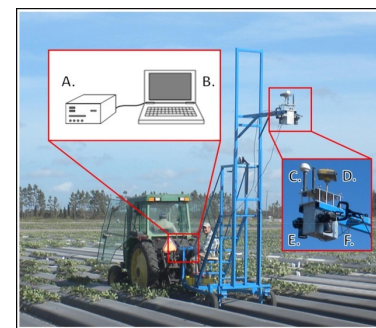
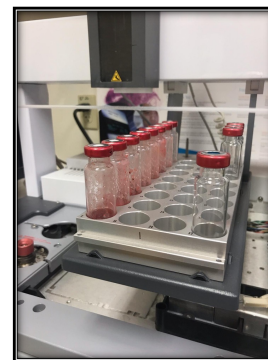
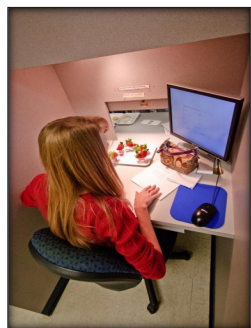
Categories of big data in strawberry breeding

Survey/Field data
 Sensory evaluation
 Field phenotyping

Genomics data
 Short reads (Illumina®)
 Long reads (Pacbio® Hifi)
 SNP array

Other omics data
 Transcriptome
 Metabolites (Volatile)

Remote sensing data
 Canopy reflectance
 High resolution image



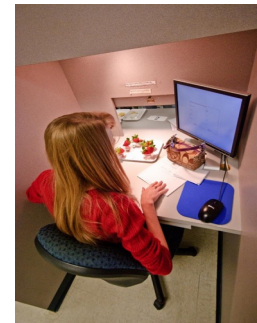
Volatiles modulating eating experience

Objectives:

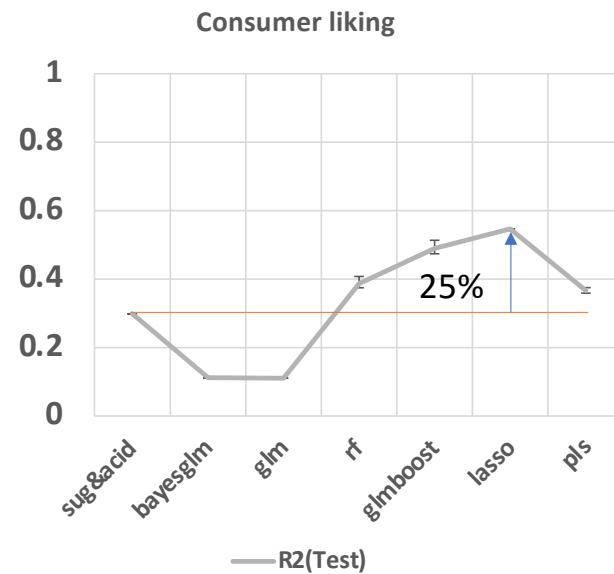
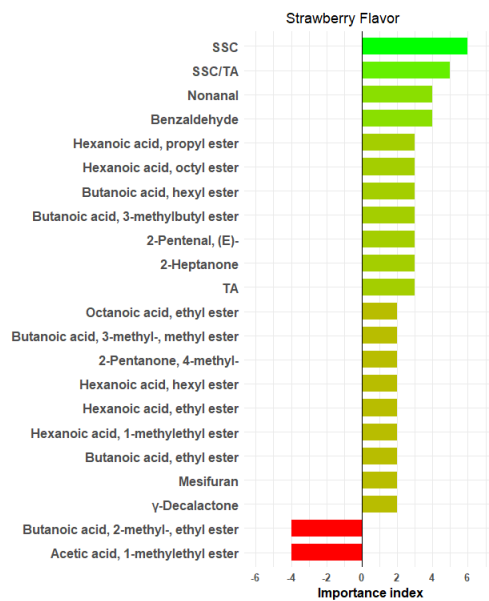
- Identification of volatiles **important for consumer liking**
- **Prediction of consumer liking** with metabolites data

Materials and methods:

- **154** genotype/harvest date combinations, over **15,400** sensory/liking evaluations (consumer panel)
- **213** genotype/harvest date combinations, over **2,130** sensory evaluations (descriptive panel)



Volatiles modulating eating experience



Article | Open Access | Published: 01 April 2021

Strawberry sweetness and consumer preference are enhanced by specific volatile compounds

Zhen Fan, Tomas Hasing, Timothy S. Johnson, Drake M. Garner, Michael L. Schwieterman, Christopher R. Barbey, Thomas A. Colquhoun, Charles A. Sims, Marcio F. R. Resende & Vance M. Whitaker

Horticulture Research 8, Article number: 66 (2021) | Cite this article

ORIGINAL RESEARCH article

Front. Plant Sci., 17 March 2021 | <https://doi.org/10.3389/fpls.2021.640704>



Volatiles Influencing Sensory Attributes and Bayesian Modeling of the Soluble Solids–Sweetness Relationship in Strawberry

Zhen Fan¹, Anne Plotto², Jinhe Bai³ and Vance M. Whitaker^{1*}

Discovering flavor genes and their regulatory elements

Objectives:

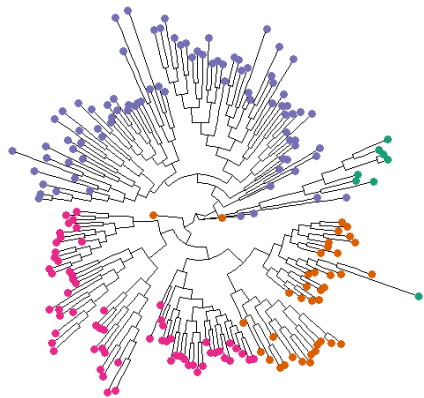
- Identification of **biosynthetic genes** for volatile production
- Exploration of **natural variation** in their **regulatory regions**

Materials and methods:

- RNAseq data for **196** breeding accessions
- Pacbio Hifi reads for a UF variety; short reads for parents
- Volatile quantification for a GWAS panel with over **300** individuals

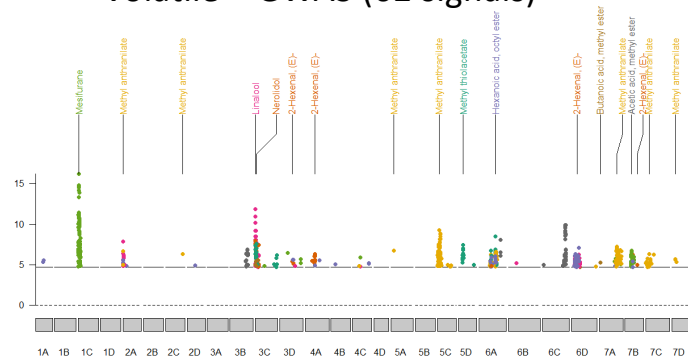
Discovering flavor genes and their regulatory elements

eQTL

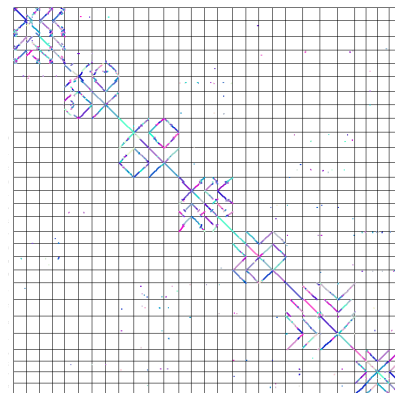


Expressed gene	62181
# eQTL/cis-eQTL/trans-eQTL	68535/23452/45083
eGenes with cis-eQTL	22731
eGenes with dis-eQTL	20650

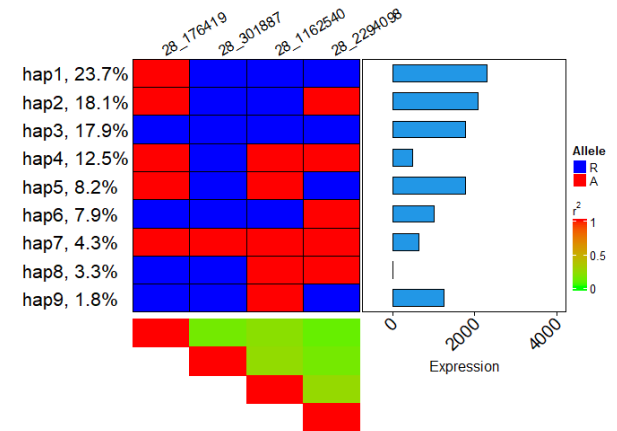
Volatile – GWAS (62 signals)



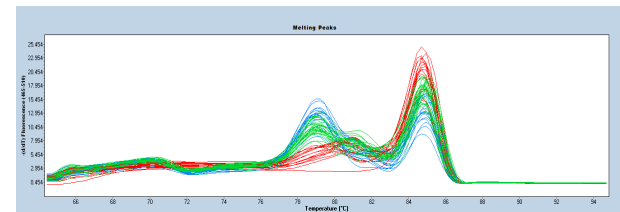
Phased genome (88 & 79 gaps)



Favorable alleles



Functional markers



Paper in review

SNP array + GS

Genomic Selection

Objectives

1. Predict parents from the seedling population
2. Predict parents from current Advanced selection trials

GS Methodology

- GP Methods: GBLUP, Bayes B, RKHS, Deep learning models
- Software: ASReml-R, BGLR, Tensorflow
- GP Models: Additive
- GBVs: Phenotype Ranking & Selection Index



Dr. Luis Osorio




Dr. Sujeet Verma



Predictive Ability

Predictive Ability: $corr(y, \hat{y})$

Trait	PBLUP	GS Methods		
		GBLUP	Bayes B	RKHS
AWT	0.44	0.49	0.49	0.52
SSC	0.37	0.43	0.44	0.45
EMY	0.14	0.29	0.3	0.3
TMY	0.24	0.31	0.35	0.33
TC	0.14	0.32	0.35	0.32

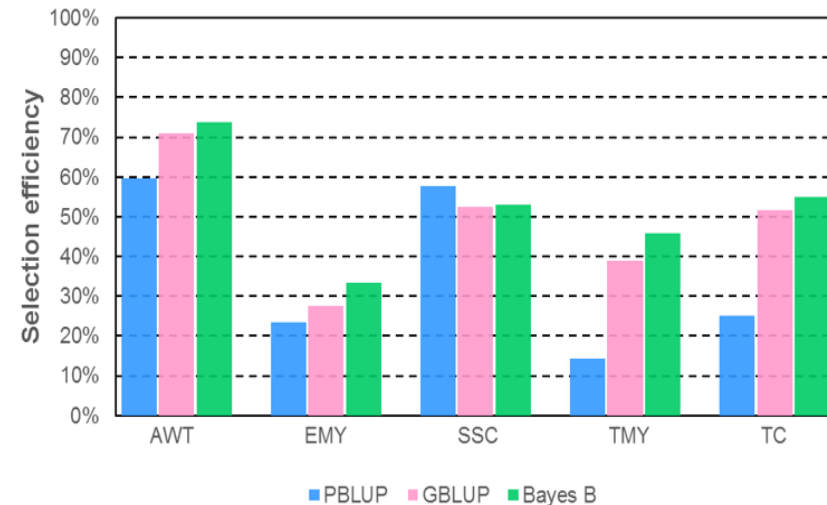
An experimental validation of genomic selection in octoploid strawberry 

Salvador A Gezan, Luis F Osorio, Sujeet Verma, Vance M Whitaker 

Horticulture Research, Volume 4, 2017, 16070, <https://doi.org/10.1038/hortres.2016.70>

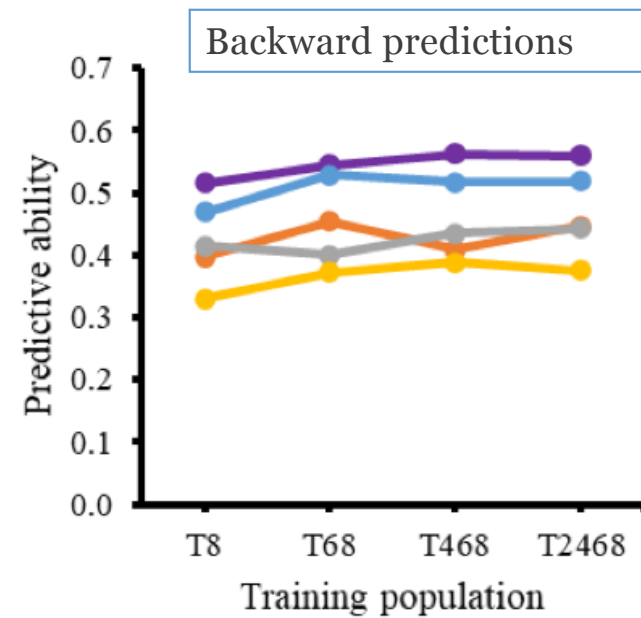
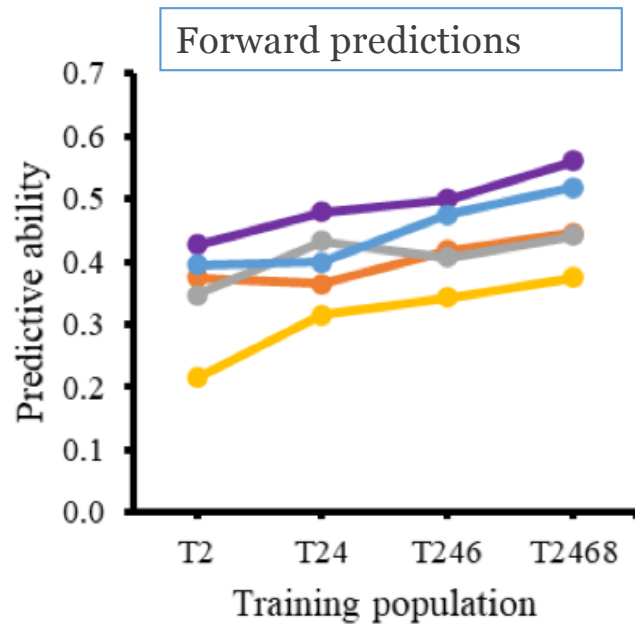
Published: 11 January 2017 [Article history](#) ▼

Parent Selection Efficiency (PSE)



$$PSE = GG_{PRED} / GG_{TOTAL}$$

Validation of GS over multiple cycles



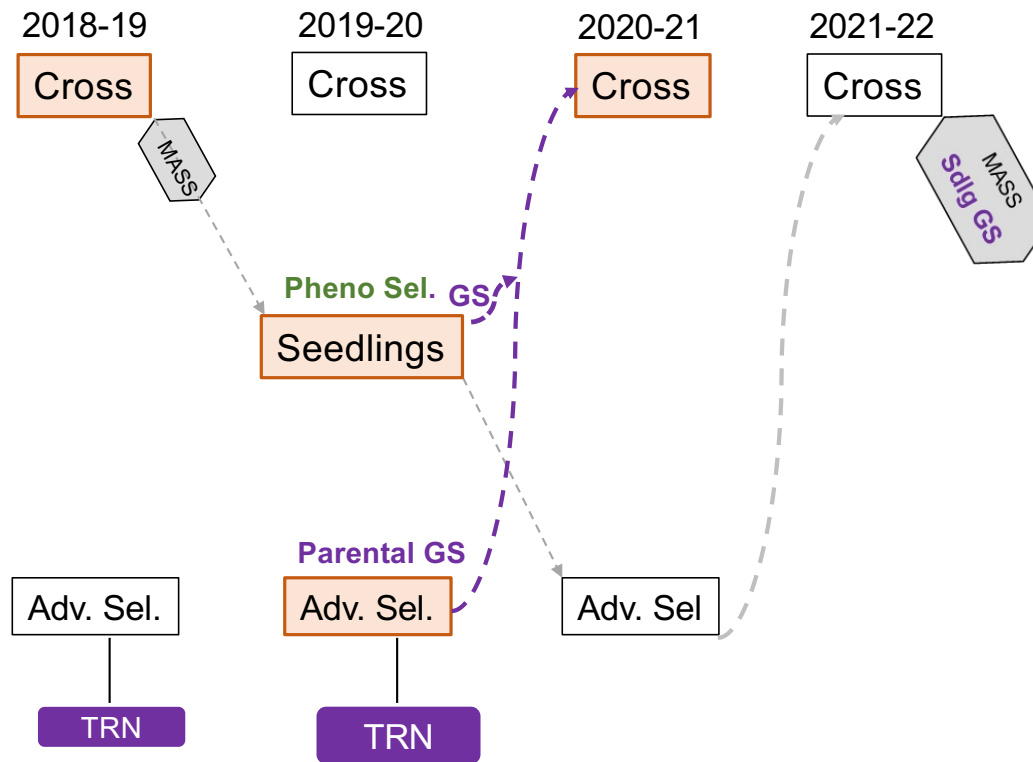
Independent Validation of Genomic Prediction in Strawberry Over Multiple Cycles

Luis F. Osorio¹, Salvador A. Gezan^{2*}, Sujeet Verma¹ and Vance M. Whitaker^{1*}

¹Gulf Coast Research and Education Center, University of Florida, Wimauma, FL, United States

²School of Forest Resources and Conservation, University of Florida, Gainesville, FL, United States

Strawberry breeding cycles



High-resolution images

UF

Modeling strawberry biomass and leaf area

Objectives:

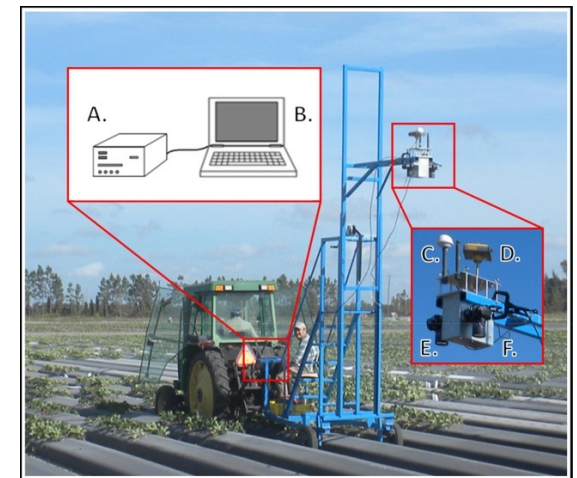
- Modeling strawberry **biomass** and **leaf area** using object-based analysis of high-resolution images

Materials and methods:

- Vegetation Mobile Mapping System (VMMS) consisted of two digital cameras, a GPS and a GNSS receiver
- Total leaf area and dry biomass of the plants were measured using destructive methods.



DR. AMR ABD-ELRAHMAN



Modeling strawberry biomass and leaf area

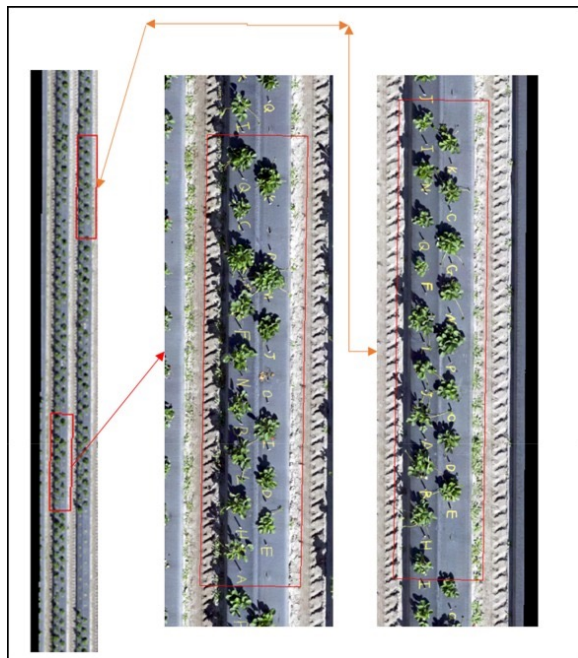


Table 8. LOOCV and ten-fold CV results for the dry biomass and leaf area models.

	Dry Biomass Models			Leaf Area Models		
	R ²	Cross-validation RSME (g)	Cross-validation RSME (%)	R ²	Cross-validation RSME (m ²)	Cross-validation RSME (%)
LOOCV	0.82	7.96	8.75	0.79	5.80E-02	9.28
Ten-fold CV	0.84	7.72	8.48	0.80	5.60E-02	8.96

Modeling strawberry biomass and leaf area using object-based analysis of high-resolution images

Zhen Guan ^{a, b}, Amr Abd-Elrahman ^{a, b}, Zhen Fan ^c, Vance M. Whitaker ^c, Benjamin Wilkinson ^a

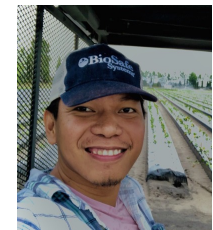
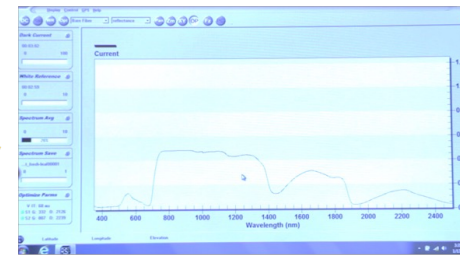
Prediction of powdery mildew resistance in strawberry

Canopy reflectance measurement

ASD Field Spec 4

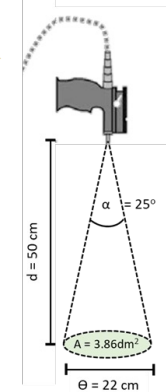


Real-time spectral data recording



Ronald Tapia
PhD student

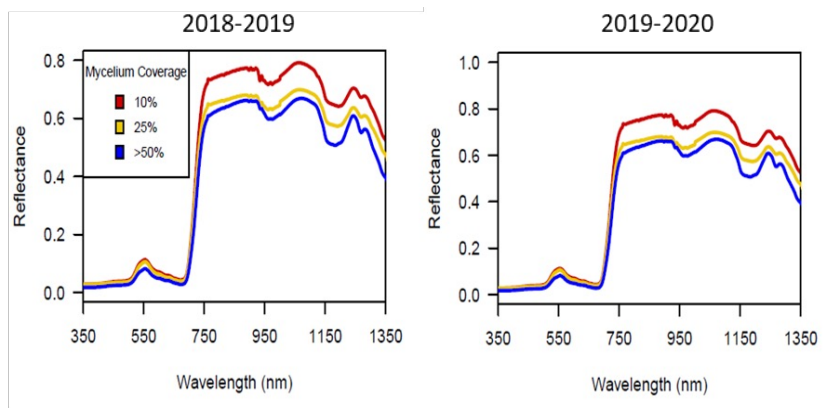
Field of view



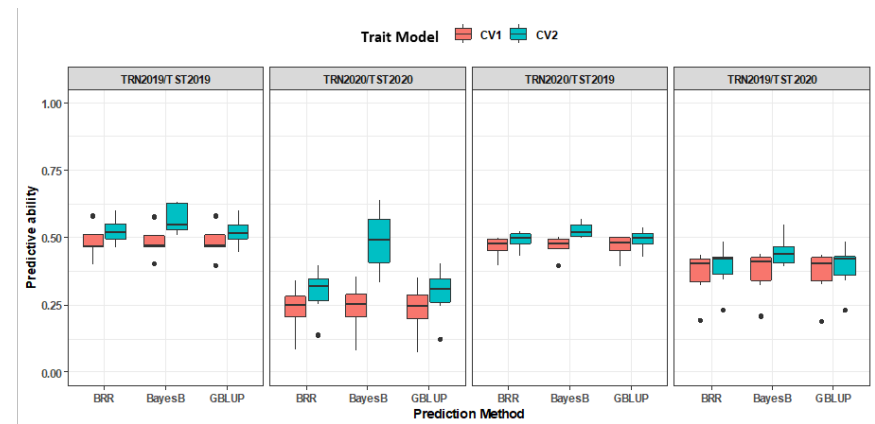
Canopy reflectance acquisition:

- 350-2500 nm with 1 nm resolution
- 10 AM – 2 PM with full sunlight
- Calibration every 10 mins with white board panel
- Capture 3 times/plant where each is an average of 10 recordings
- >8,000,000 data points for 450 individuals

Prediction powdery mildew resistance in strawberry



Canopy-based spectral signatures of powdery mildew disease of strawberry.



PA using simple univariate method with only genomic information, and a combination of genomic and spectral data as predictors.

Combining canopy reflectance spectrometry and genome-wide prediction to increase response to selection for powdery mildew resistance in cultivated strawberry

Ronald Tapia^{1,2}, Amr Abd-Elrahman^{1,3}, Luis Osorio^{1,2}, Vance M. Whitaker^{1,2,*} and Seonghee Lee^{1,2,*}

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UC Davis strawberry breeding lab
Dr. Thomas Colouhoun's research program
Dr. Charlie Sims sensory lab



Dr. Vance Whitaker

