



The University of Georgia

# De-bugging soybean

## A story of nano-spears and QTLs

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# Leaf-chewing insects in Soybean

## Lepidoptera



## Coleoptera



# Defoliation Thresholds in Soybean



30%



15%

**>20% defoliation**  
during pod formation and filling  
**Yield reduction**



**Vegetative**

**Reproductive**

[extension.entm.purdue.edu/](http://extension.entm.purdue.edu/)  
Haile et al., 1998

# Resistance Mechanisms: **Antixenosis**

**Experiment:** choice

**Resistance:** non preference

**Phenotype:** reduced plant defoliation



**Greenhouse plots**

**Susceptible**

**Resistant**



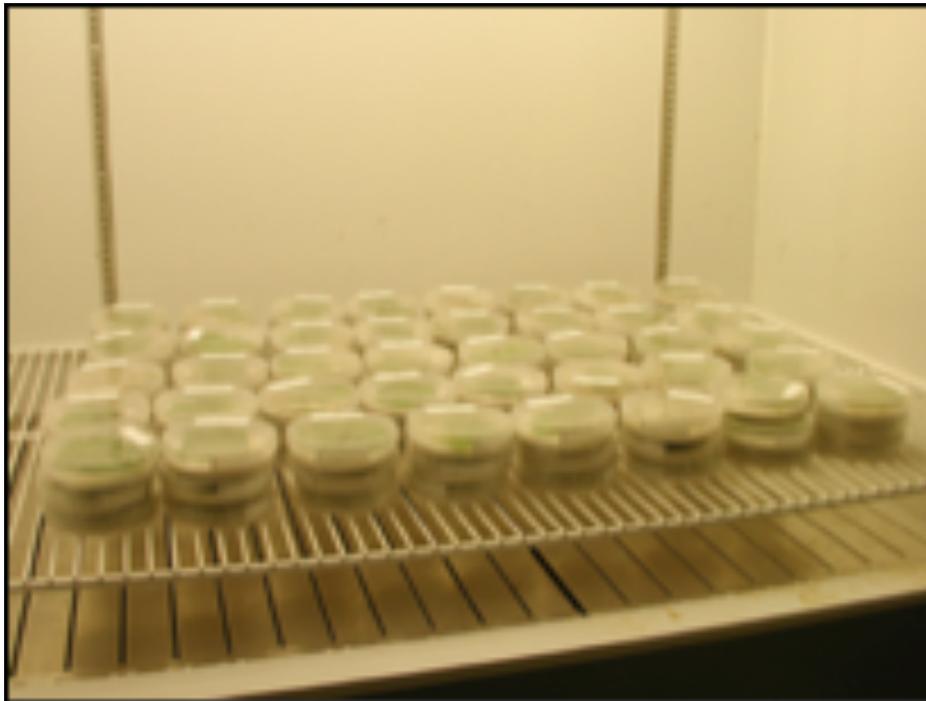
**Visual rating: % defoliation**

# Resistance Mechanisms: **Antibiosis**

**Experiment:** non choice

**Phenotype:** larval weight

**Growth chamber**

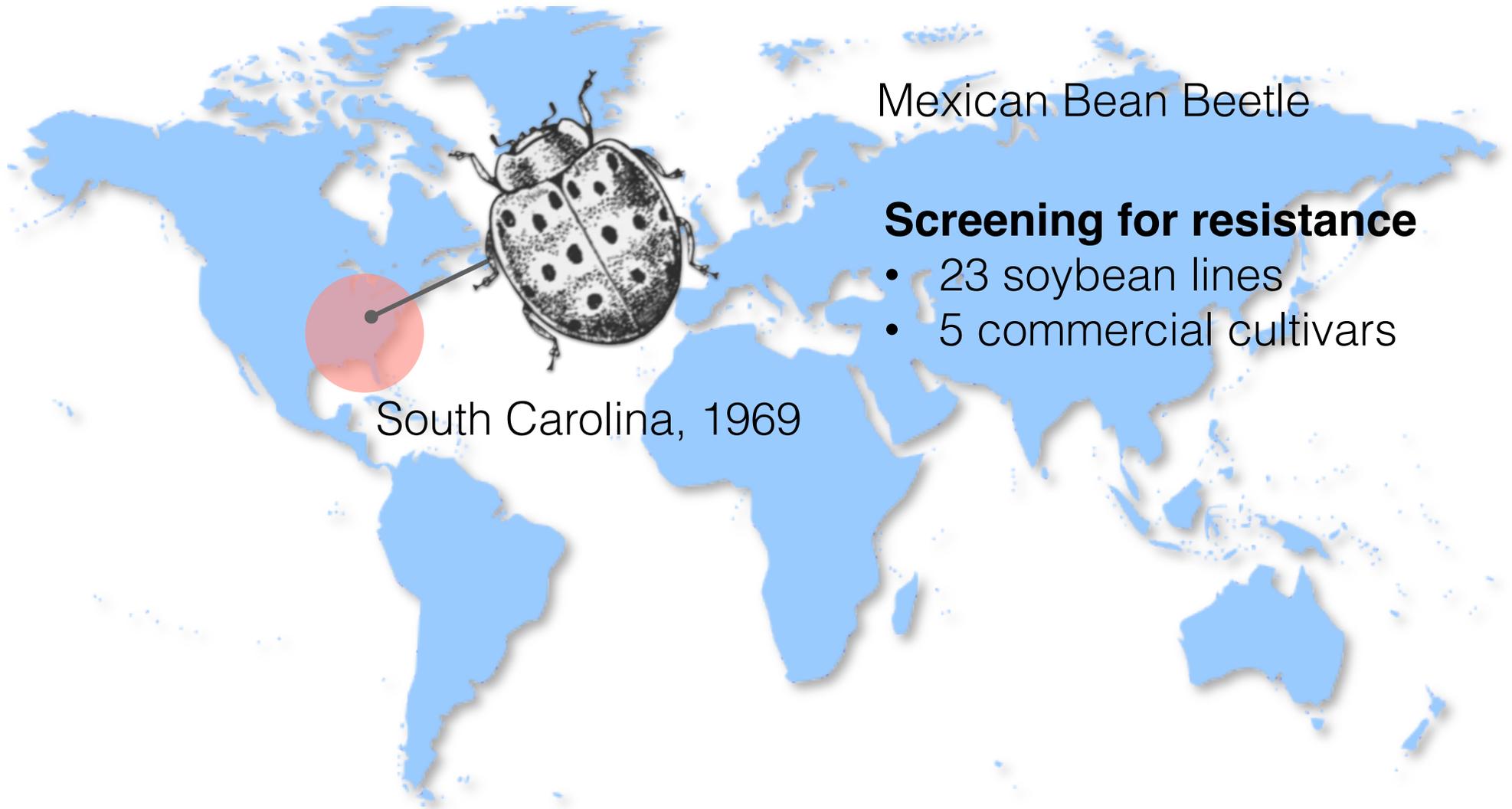


**Susceptible**

**Resistant**



# Insect Resistant Soybean



Mexican Bean Beetle

## **Screening for resistance**

- 23 soybean lines
- 5 commercial cultivars

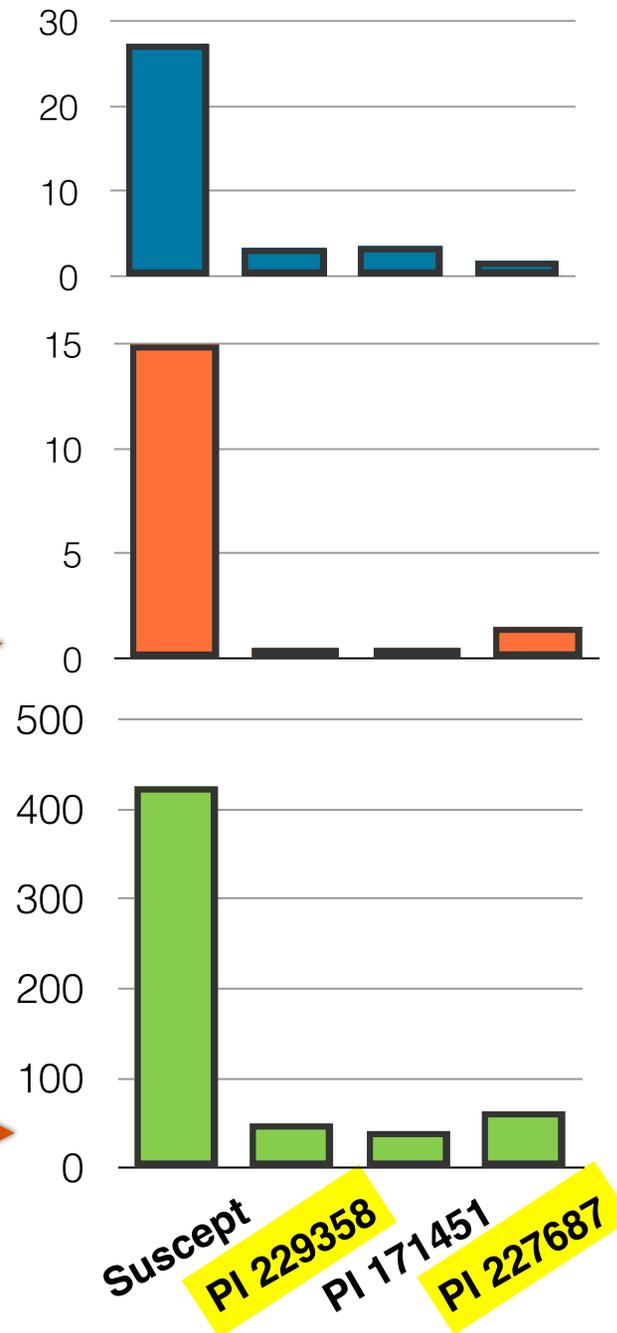
# Insect Resistant Soybean



# beetles / plant

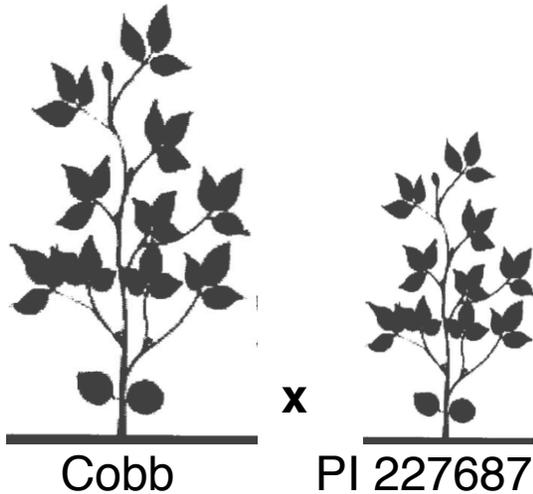
# egg masses / plant

Larvae weight / plant



Van Duyn et al., 1971

# Resistance in PI 227687



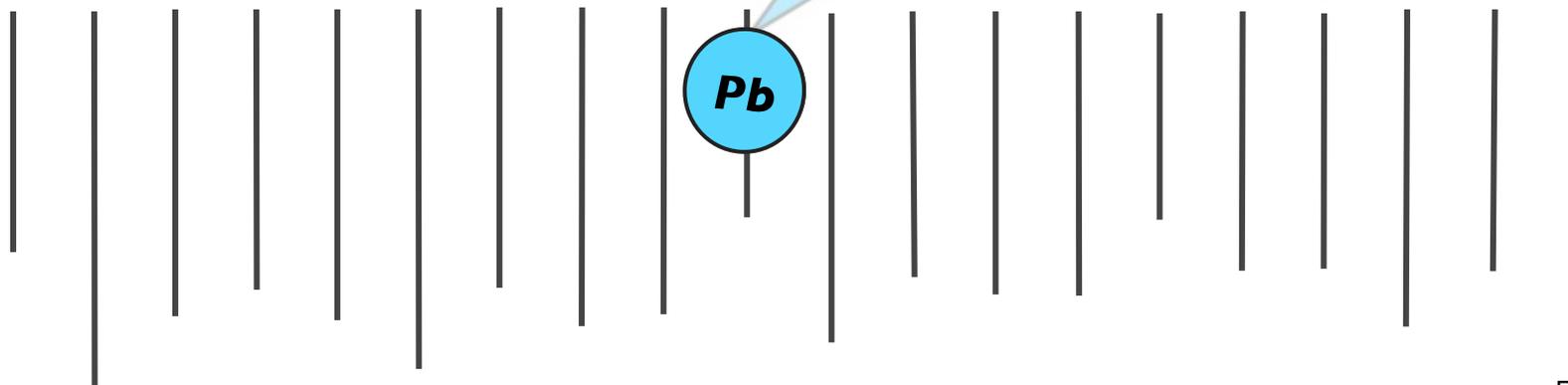
F2:3 population



Corn earworm

## Soybean Linkage Groups

RFLP map



QTL-E explained

26% Antibiosis

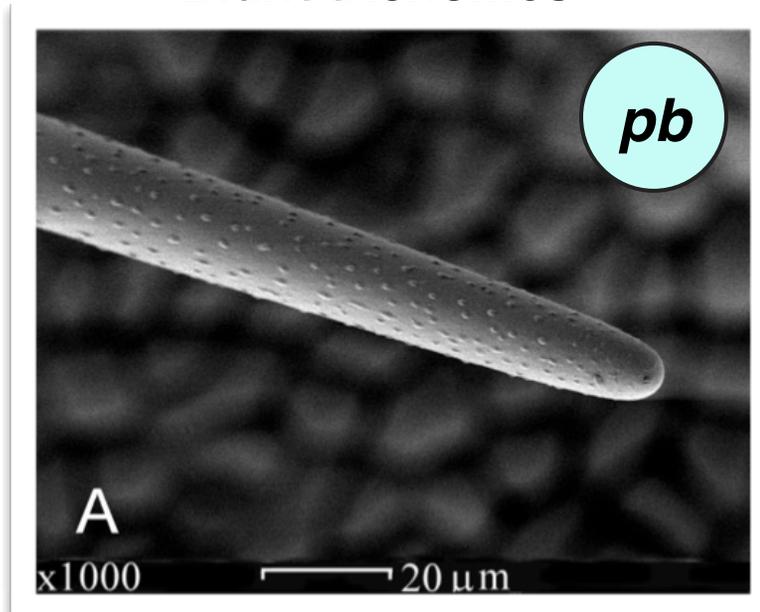
20% Antixenosis

Rector et al., 1998  
Hulburt et al., 2004  
Boerma & Walker, 2005  
Palmer, 2008

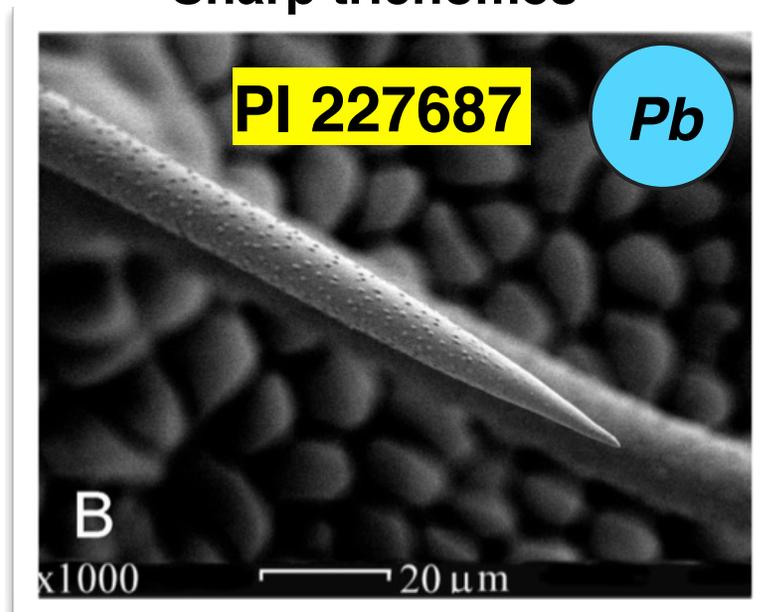
# Sharp Pubescence locus: *Pb*



Blunt trichomes



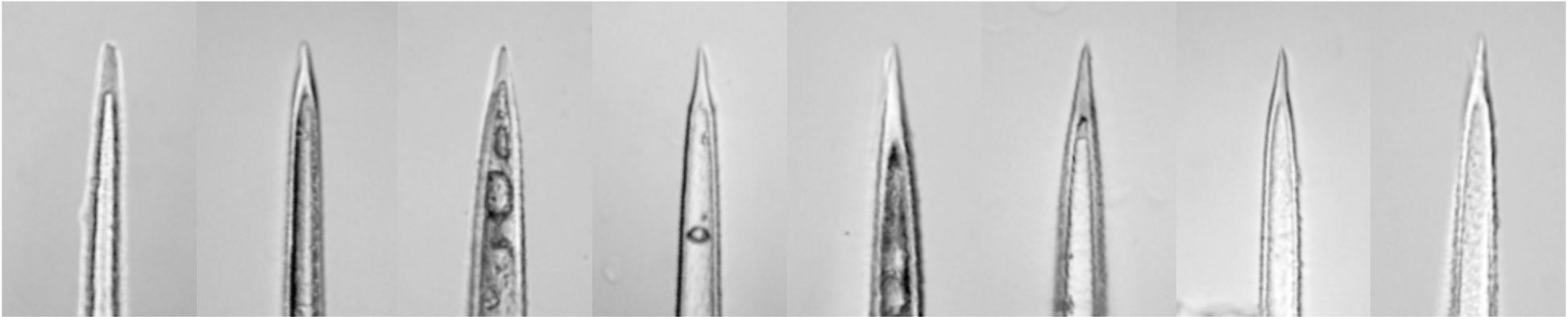
Sharp trichomes



# Sharp vs. Blunt Pubescence

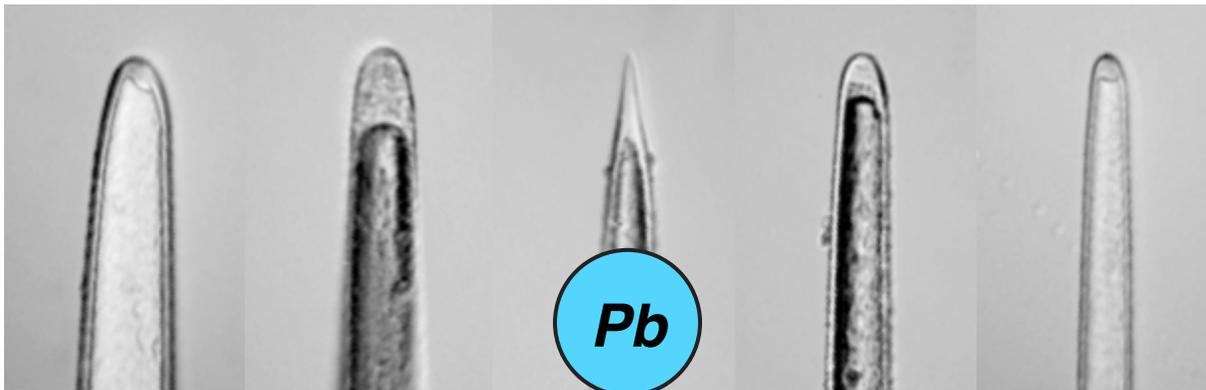
***Pb***

**Wild soybean**



***pb***

**Cultivated soybean**



# CHOICE IOWA SEEDS

56 ANNUAL  
CATALOGUE

1921

ONLY ONE  
THE BEST  
QUALITY

Soy  
Beans

Iowa  
Gold Mine  
Corn



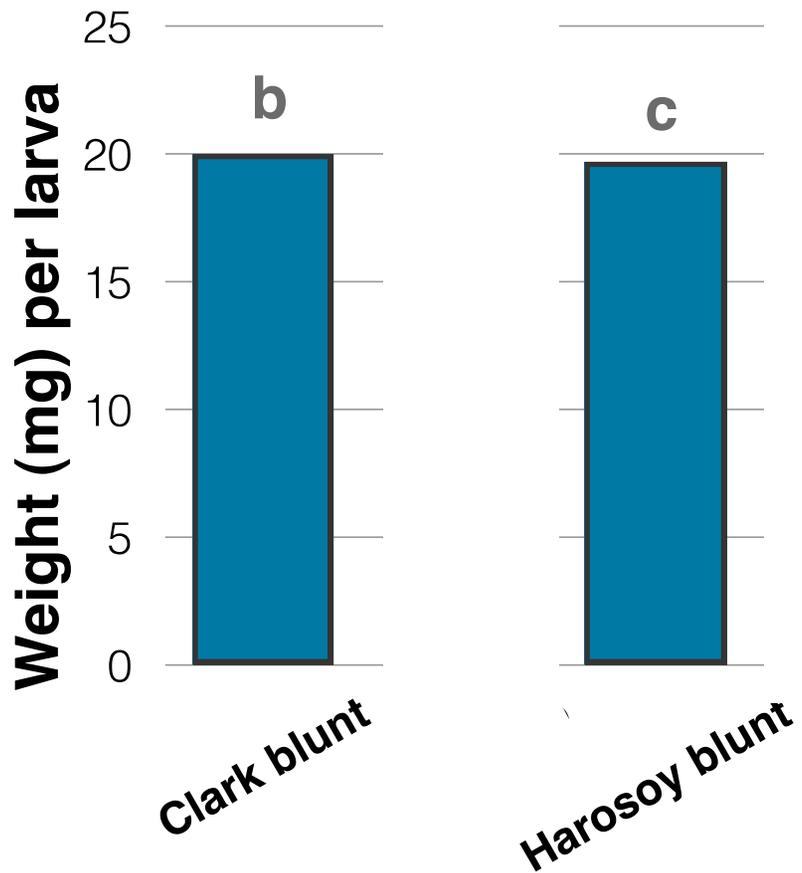
SEEDSMEN TO THE AMERICAN PEOPLE

IOWA SEED CO. DES MOINES  
IOWA U.S.A.

Smithsonian Museum  
Catalog Collection

# Sharp vs. Blunt Pubescence

**Phenotype:** larval weight



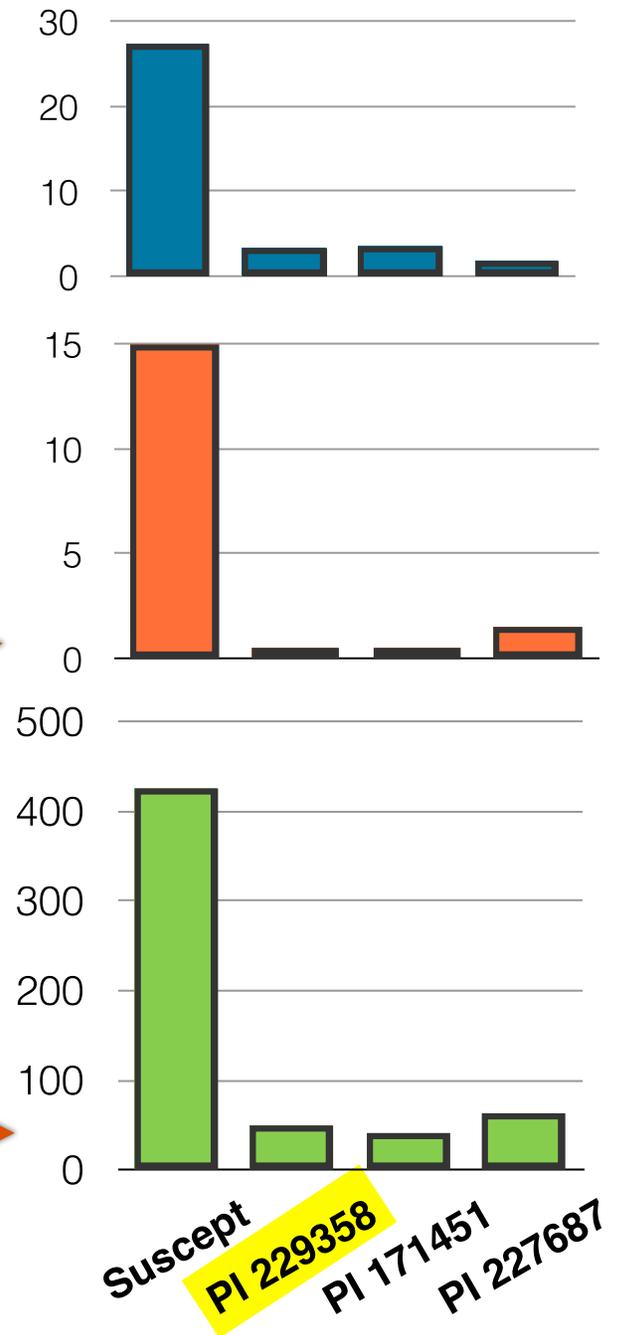
# Insect Resistant Soybean



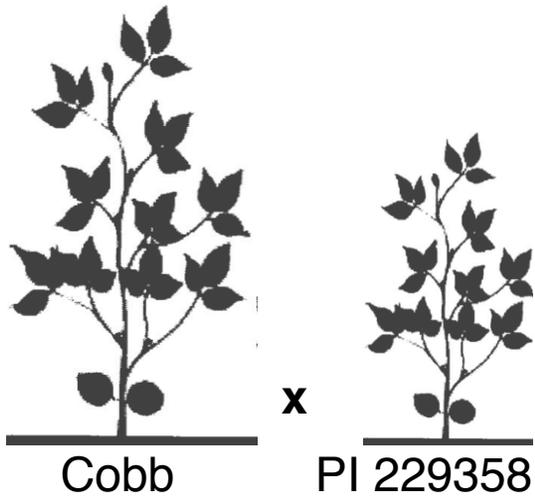
# beetles / plant

# egg masses / plant

Larvae weight / plant



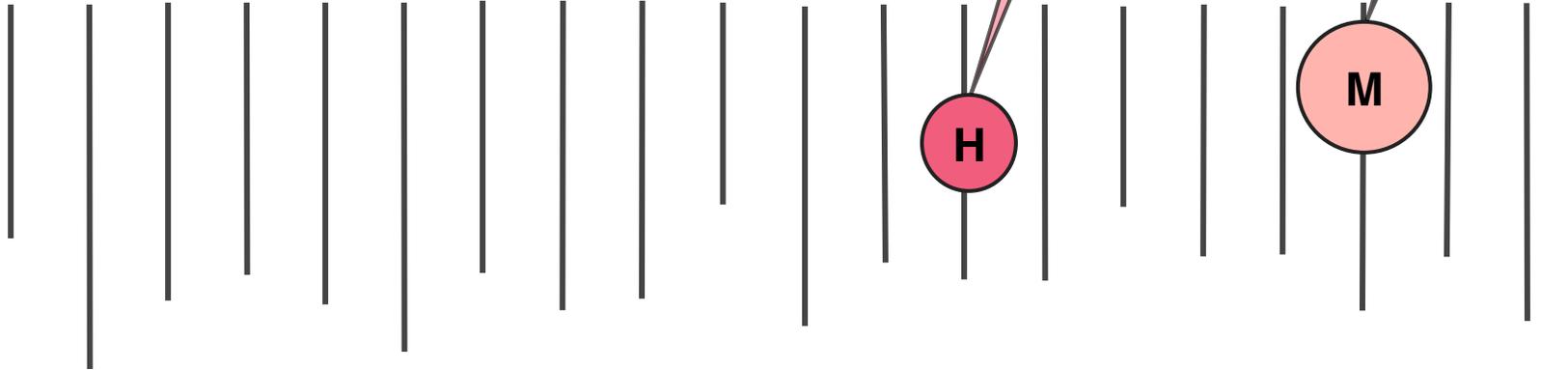
# QTLs in PI 229358 - Antixenosis



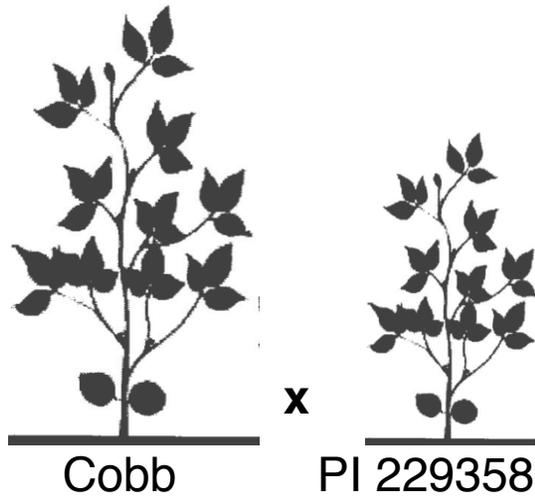
F2:3 population



Soybean Linkage Groups  
RFLP map



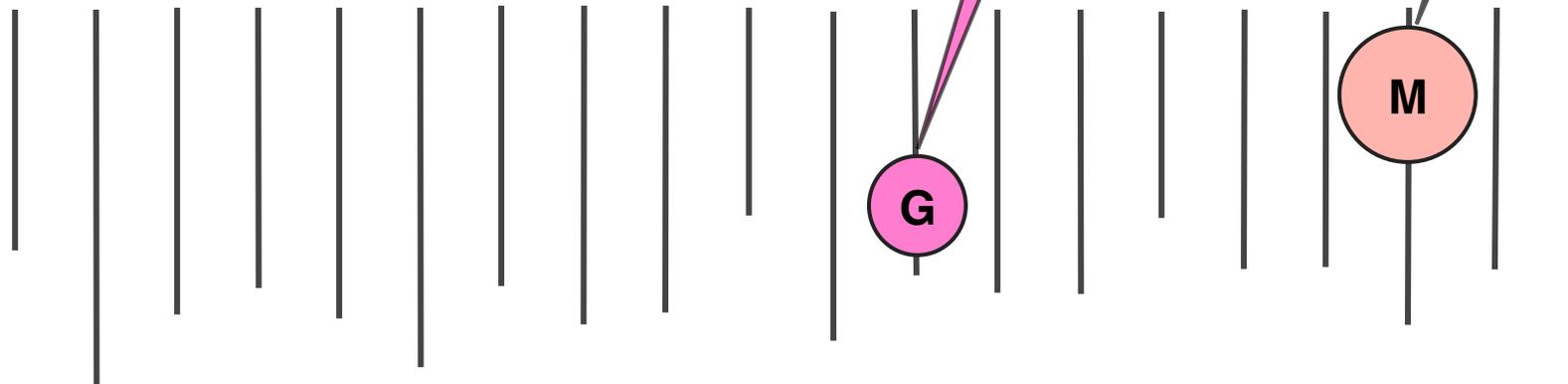
# QTLs in PI 229358 - Antibiosis



F2:3 population



Soybean Linkage Groups



# QTL-M **activates** the other QTLs

## Benning BC6F2:3 Near Isogenic Lines

### % Defoliation

	- M	+ M
- H	31.4 <sup>a</sup>	23.4 <sup>b</sup>
+ H	30.1 <sup>a</sup>	19.2 <sup>c</sup>

### Larval weight (mg)

	- M	+ M
- G	131.3 <sup>a</sup>	80.0 <sup>b</sup>
+ G	128.4 <sup>a</sup>	49.3 <sup>c</sup>

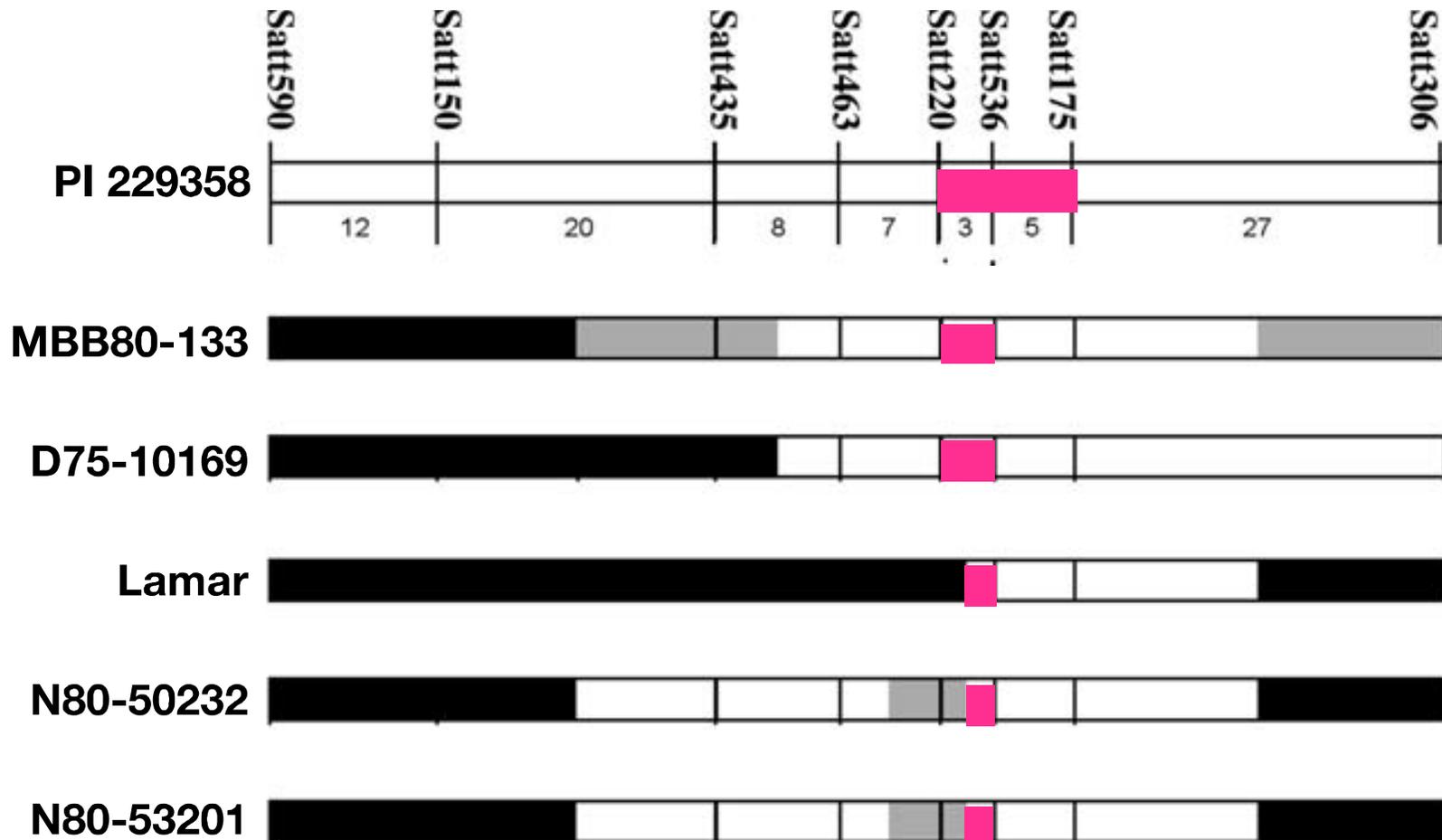
# Retrospective study of QTL-M

## 15 soybean breeding lines

Phenotypic selection  
Insect Resistance

## Graphical genotype

82 cM of Chromosome 7  
SSR flanking QTL-M



# QTL-M + Sharp Pubescence

**Benning BC<sub>6</sub>F<sub>2:3</sub>**  
Greenhouse



**Soybean looper**



ANOVA: Bars represent means from fifteen replicates using RCB design ( $p < 0.001$ )

# QTL-M + Sharp Pubescence

**Benning BC<sub>6</sub>F<sub>2</sub>:3**  
Choice test

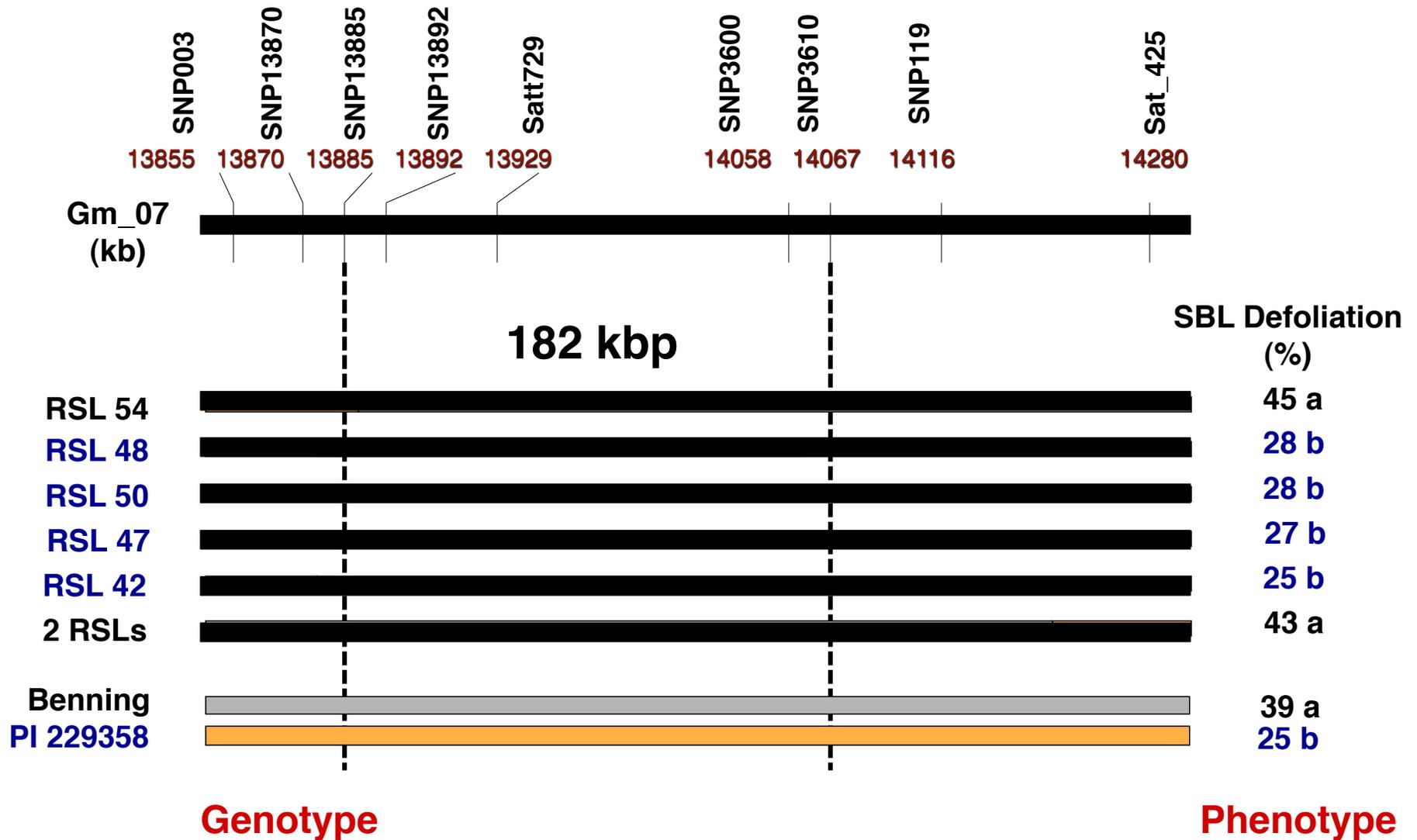


ANOVA: Bars represent means from fifteen replicates using RCB design ( $p < 0.001$ )

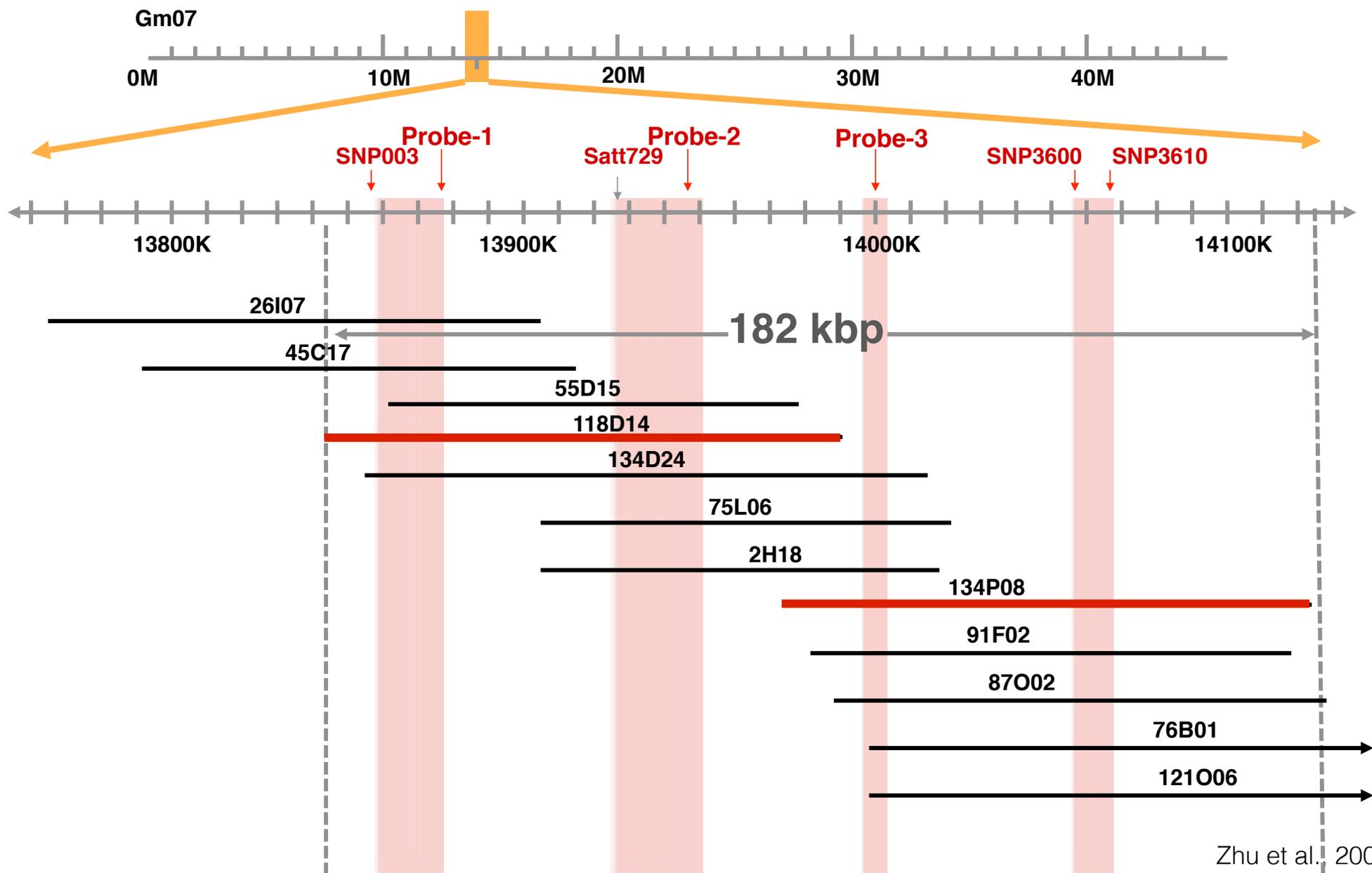
# Cloning and Characterizing **QTL**

1. Confers **antibiosis** and **antixenosis** resistance
2. Activates **QTL-G** and **QTL-H**
3. Enhances the effectiveness of **Bt**
4. Effective across **multiple** insect herbivores

# QTL-M: Recombinant Substitution Lines

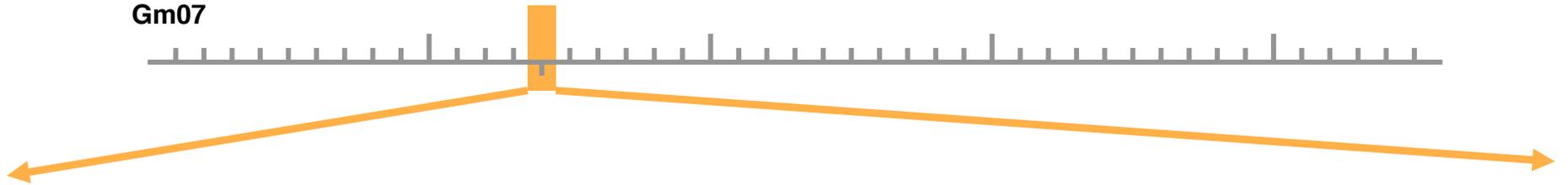


# QTL-M: BAC Contig – PI 229358



# BAC Sequence Aligned to the Soybean Genome

Gm07



13900K

14000K

14100K

182 kbp

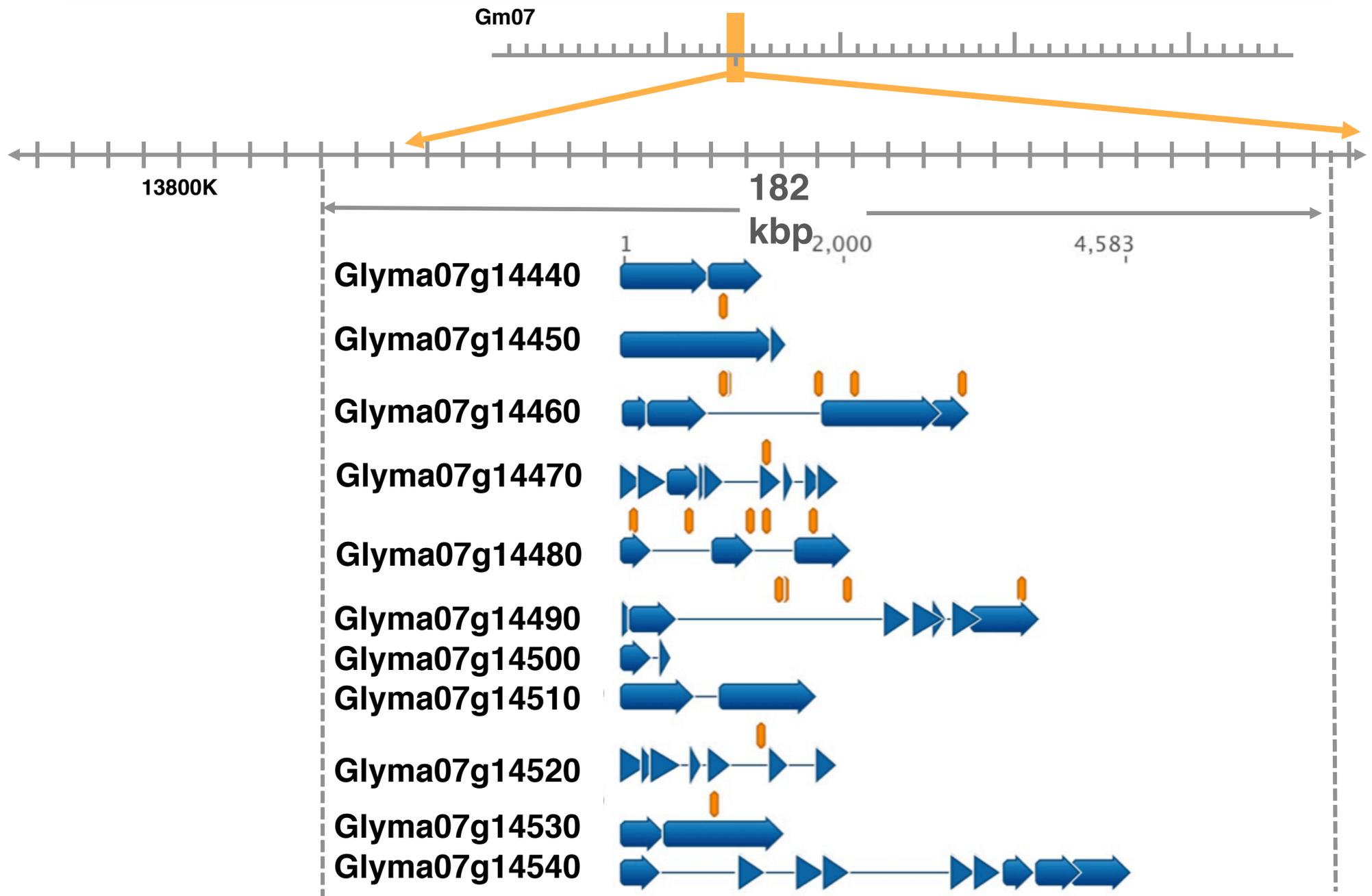
20000 40000 60000 80000 100000 120000 140000 160000 180000 200000 220000 240000 260000 280000

178 kbp – PI 229358

20000 40000 60000 80000 100000 120000 140000 160000 180000 200000 220000 240000 260000

182 kbp – Williams 82

# SNPs between Williams 82 and PI 229358



# Polymorphisms with **Susceptible** Accessions



32 Soybean Genotypes Ancestral to Modern Varieties

# SNPs **Unique** to Insect Resistant Soybeans Accession

**Glyma07g14440** Protein of unknown function



**Glyma07g14450** Protein of unknown function



**Glyma07g14460** Cytochrome P450



**Glyma07g14470** Ploop-NTPase



**Glyma07g14480** MYB related protein



**Glyma07g14490** Phosphoglycerate mutase



**Glyma07g14500** Protein of unknown function



**Glyma07g14510** Glucosyl/Glucuronosyl transferase



**Glyma07g14520** Protein of unknown function



**Glyma07g14530** Glucosyl/Glucuronosyl transferase

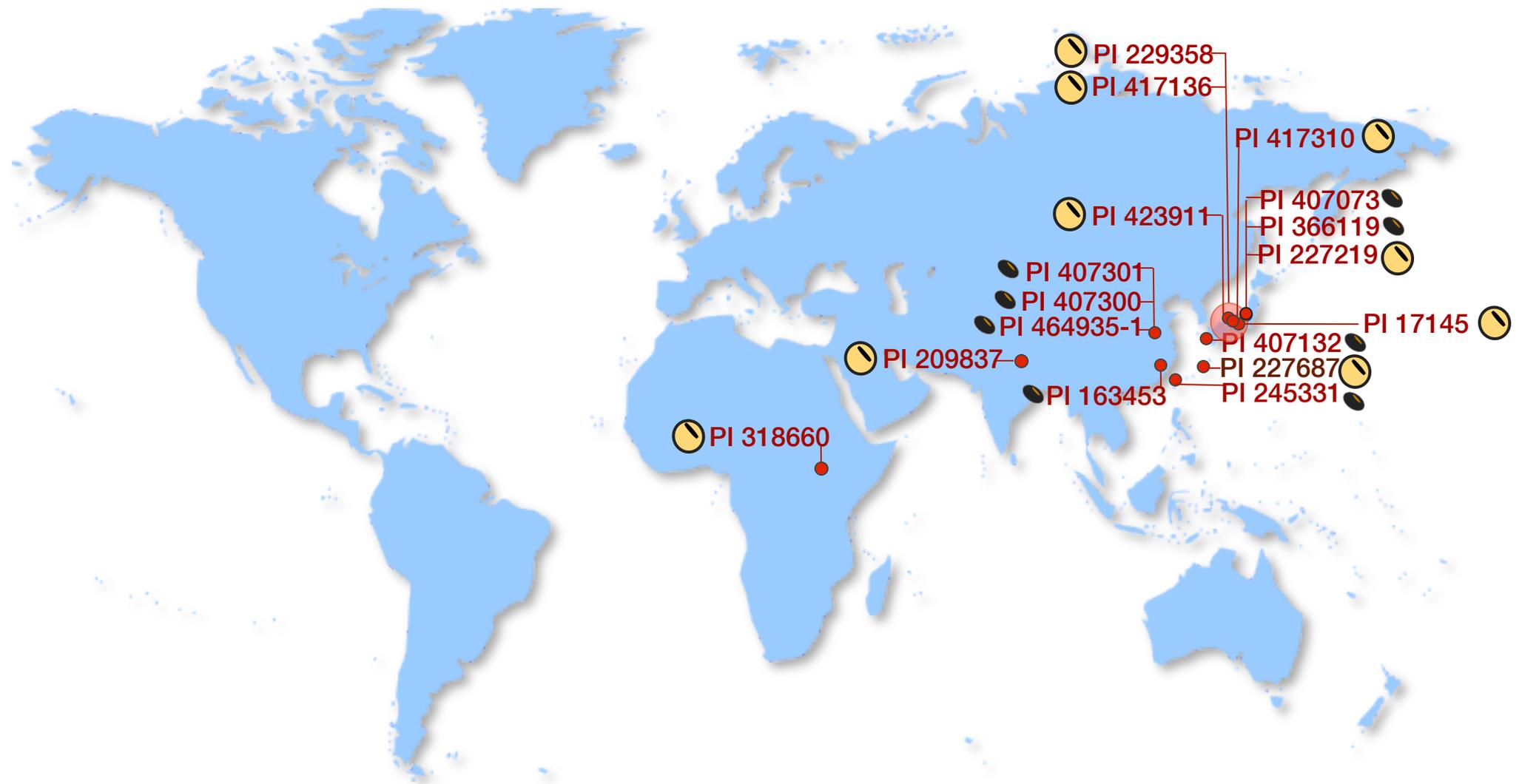


**Glyma07g14540** DNAJ/HSP40



■ Polymorphism between PI 229358 and 32 Ancestors

# Shared SNPs with Insect Resistant



16 Reported Insect-Resistant Soybean Genotypes

# SNPs **Unique** to Insect Resistant Accessions

**Glyma07g14440** Protein of unknown function



**Glyma07g14450** Protein of unknown function



**Glyma07g14460** Cytochrome P450



**Glyma07g14470** Ploop-NTPase



**Glyma07g14480** MYB related protein



**Glyma07g14490** Phosphoglycerate mutase



**Glyma07g14500** Protein of unknown function



**Glyma07g14510** Glucosyl/Glucuronosyl transferase



**Glyma07g14520** Protein of unknown function



**Glyma07g14530** Glucosyl/Glucuronosyl transferase

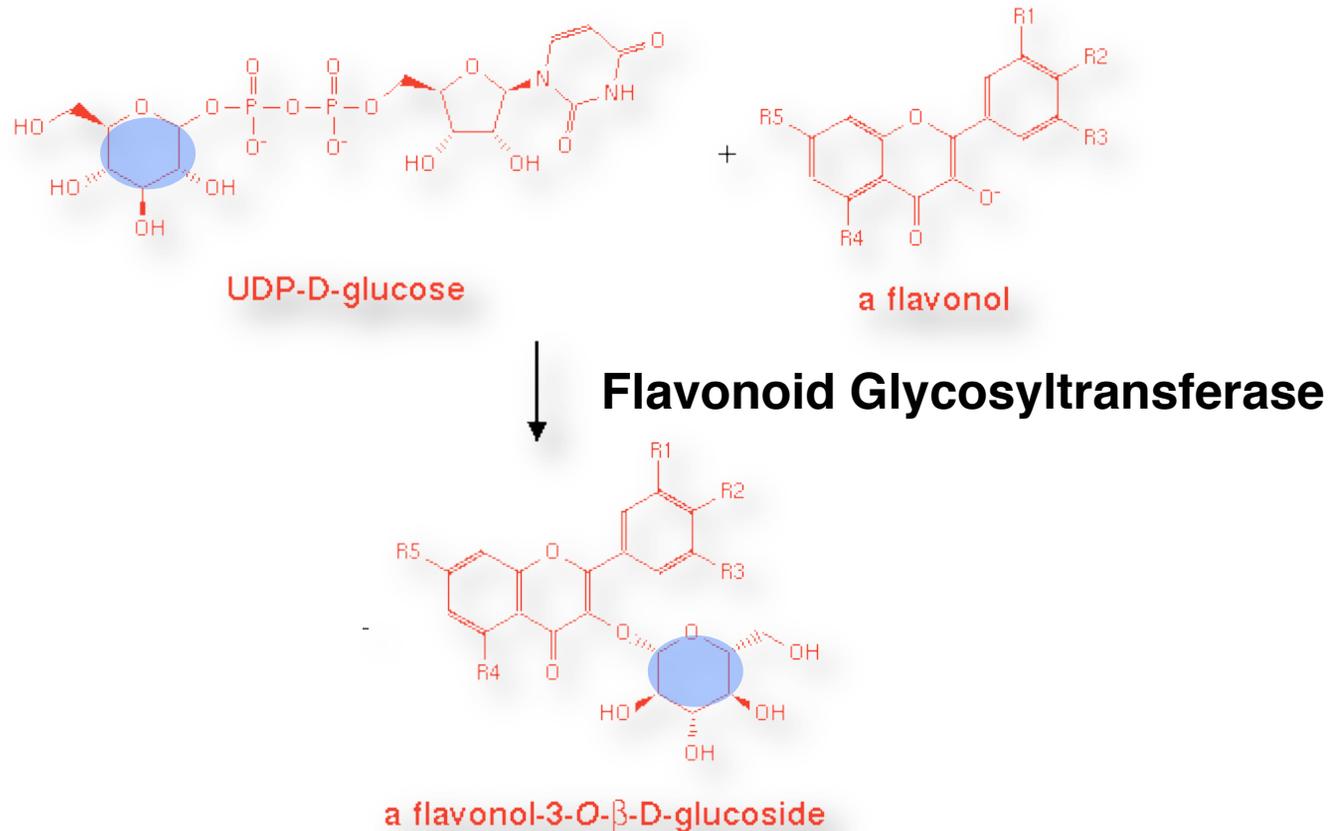


**Glyma07g14540** DNAJ/HSP40



■ SNP shared with insect resistant genotypes, excluding **PI 227687**

# Glyma07g14530 – Putative Flavonoid Glycosyltransferase



Some insects **can sequester** plant flavonoids

Other insects **are deterred** by flavonoids:

Corn earworm - **maize**

Cabbage looper – **soybean**

Bark beetle – **spruce**

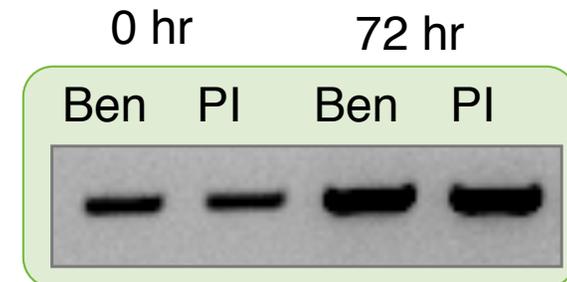
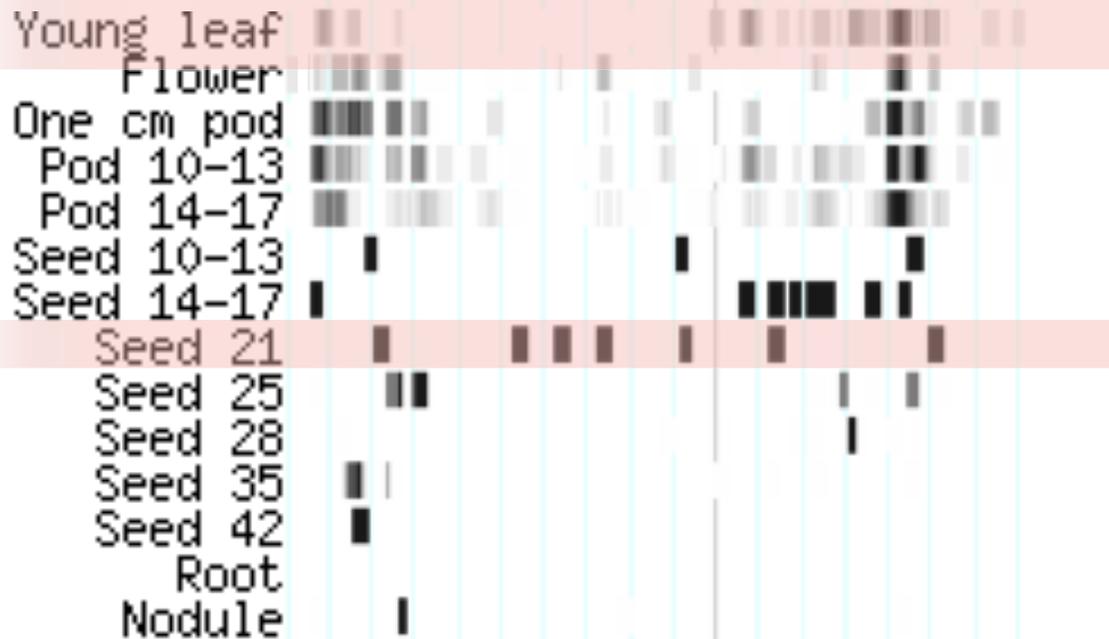
Fenny et al., 1976  
Berhow and Vaughn, 1999  
Hoffmann-Campo et al., 2001  
Widstrom and Snook, 2001

# Glyma07g14530 Expression in Leaves

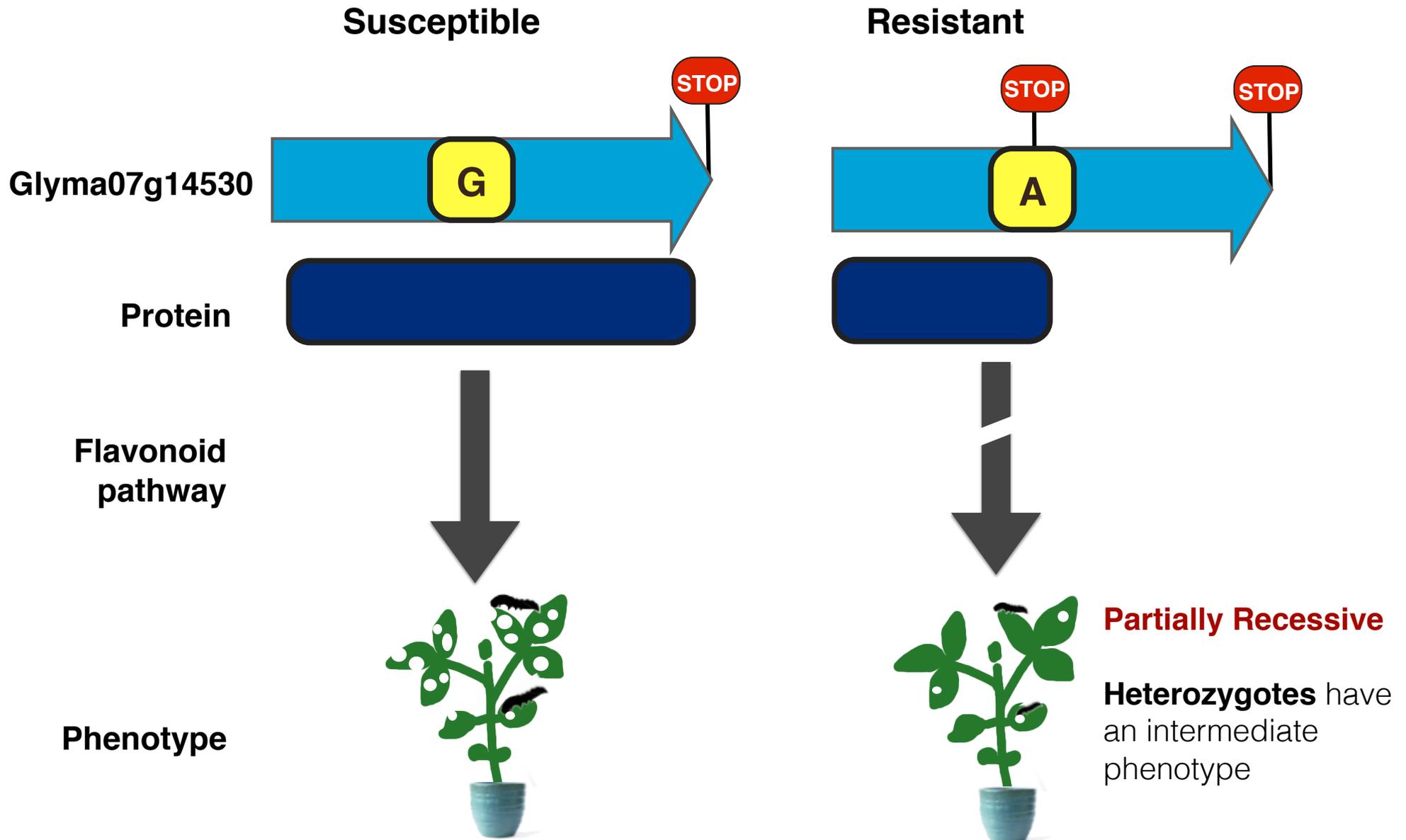
Glyma07g14530

Resistant  
1,416 bp

Susceptible  
229 bp



# Susceptible vs Resistant Soybeans



# Work in Progress

## Characterization of **Glyma07g14530**

- Complementation tests in transgenic plants
- Determine which leaf flavonoids are different between QTL-M and qtl-m plants

# Summary

- **PI 227687 resistance** is associated with sharp pubescence
- **PI 229358 resistance** is conferred by 3 QTLs:
  - **M**: antibiosis and antixenosis
  - **H**: antixenosis
  - **G**: antibiosis
- **QTL-M** activates **QTL-H** and **QTL-G**

# Summary

- **QTL-M** enhances **Bt**
  - Pyramiding these genes may provide a more durable resistance
- **QTL-M + Pb** plants are more resistant to defoliation, and caterpillars are smaller
  - Non-transgenic alternative for insect resistance
- A putative **flavonoid glycosyltransferase** is the candidate gene for QTL-M:
  - Modification of the flavonoid pathway provides a possibility to obtain insect resistance in other crops

# The De-bugging Team through the Years



**Neal Stewart**

Soybean transformation  
*Bt*

**David Hulburt**

*Pbstudies* - PI 227687

**Brian Rector**

QTL discovery  
PI 229358

**Joe Zhou**

PI 229358 BAC library  
Gene pyramiding

**Jim Narvel**

Retrospective study QTL-  
M

**Caleb Warrington**

Seed yield  
Insect Resistant  
near isogenic lines

**David Walker**

Jack near isolines

**Bo-Keun Ha**



QTL-M  
SNP discovery  
Candidate genes

# Thanks:



## Tissue culture, Transformation

- Noah Lawler
- Donna Tucker

## Entomology

- Dean Kemp
- John All
- Michael Strand

## Genotyping

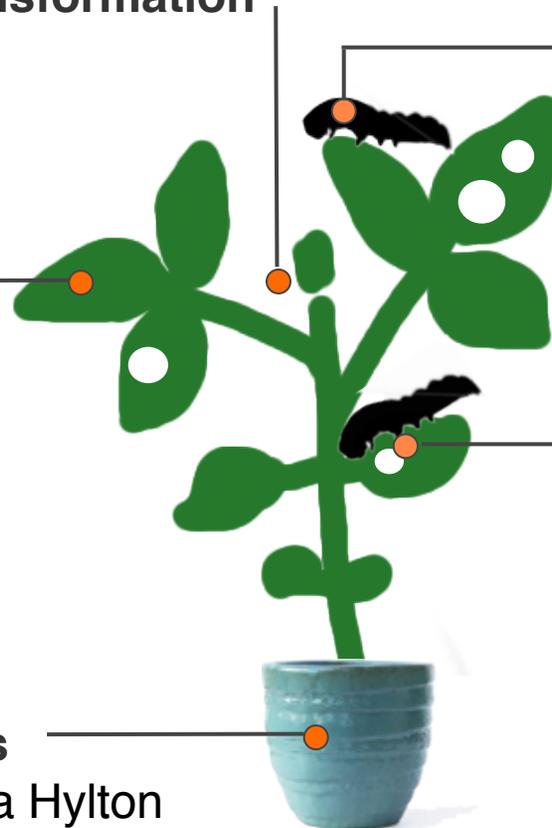
- Noah Lawler

## Gene Expression, Cloning

- Pete LaFayette

## Bioassays

- Christina Hylton



**SAVE THE DATE!**

# **“Forty years with soybeans”**

*A great*

**Symposium & Banquet**

*Celebrating the career of*

**DR. H. ROGER BOERMA**

**Distinguished Research Professor**

**Director of the Center for Applied Genetic Technologies**

**Thursday afternoon, June 28th - Friday noon, June 29th, 2012  
Athens, GA**



**Information regarding RSVP & hotel accommodations  
coming soon.**