

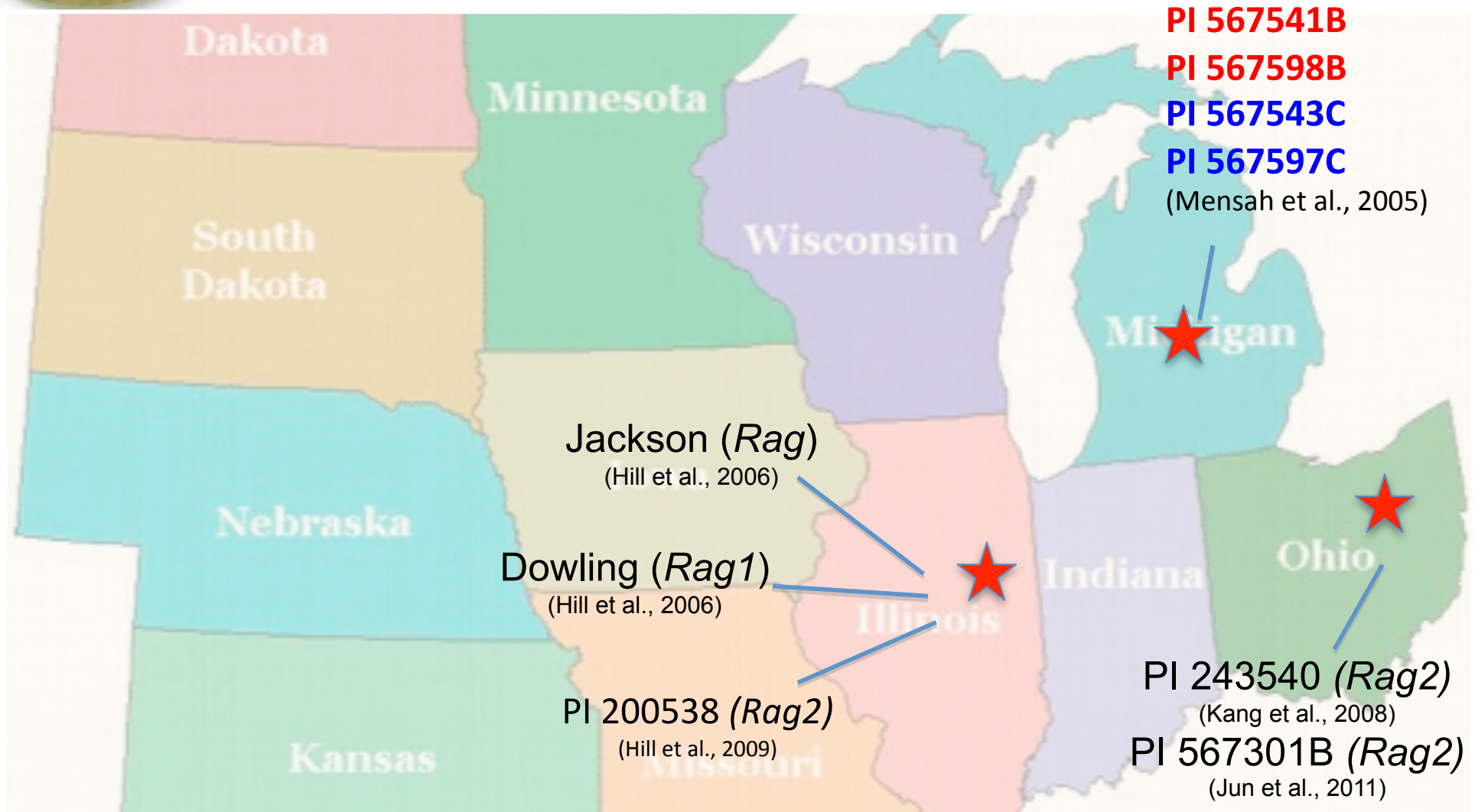
# Fine mapping of aphid resistance genes in soybean PI567543C PI567598B and PI567541B

- *Rag3*
- *rag1b* and *rag3*
- *rag1c* and *rag4*



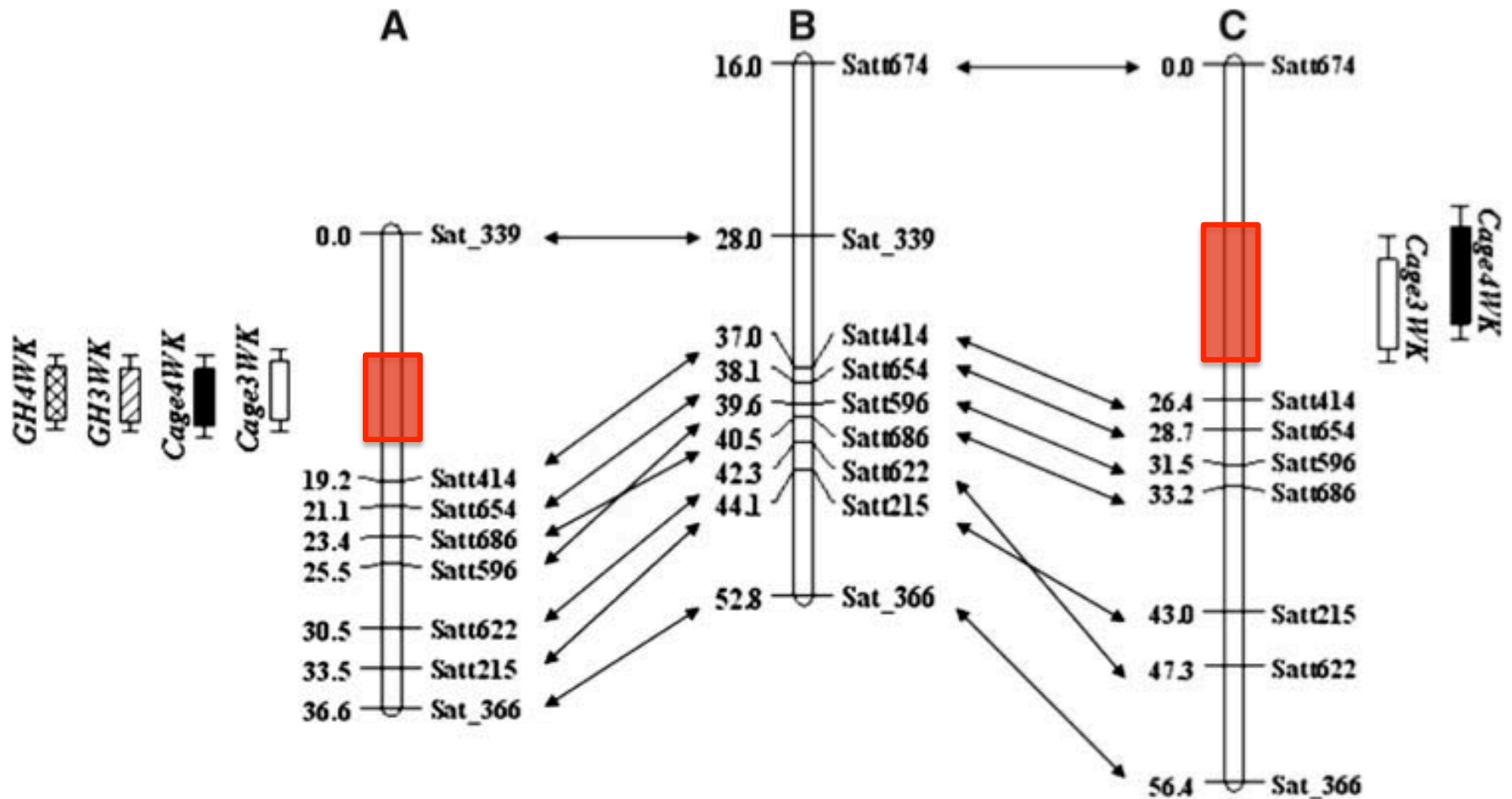


# SA resistant sources in the Midwest



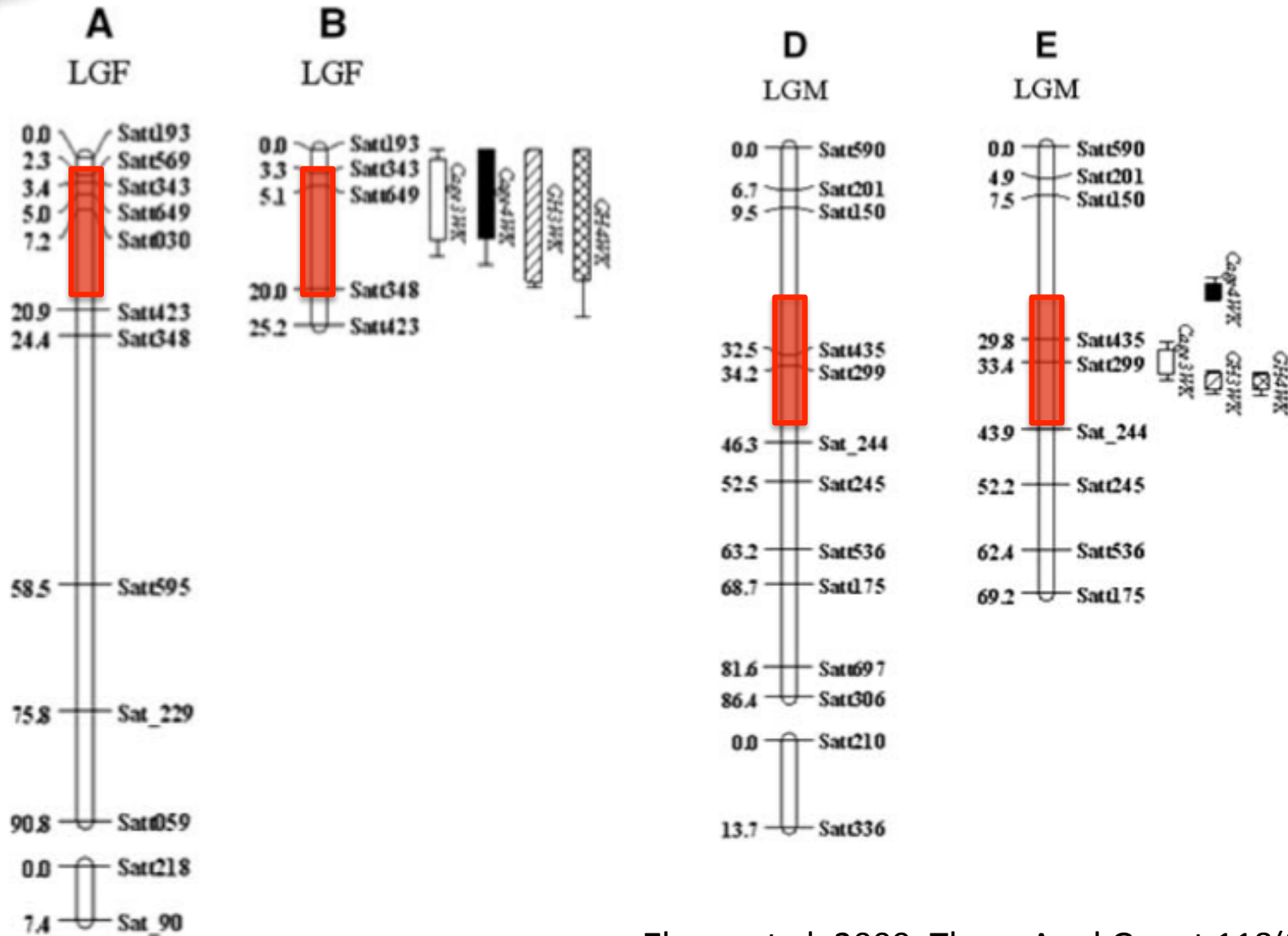


# QTL mapping of *Rag3* using 249 F<sub>4</sub>-derived lines in PI 567543C





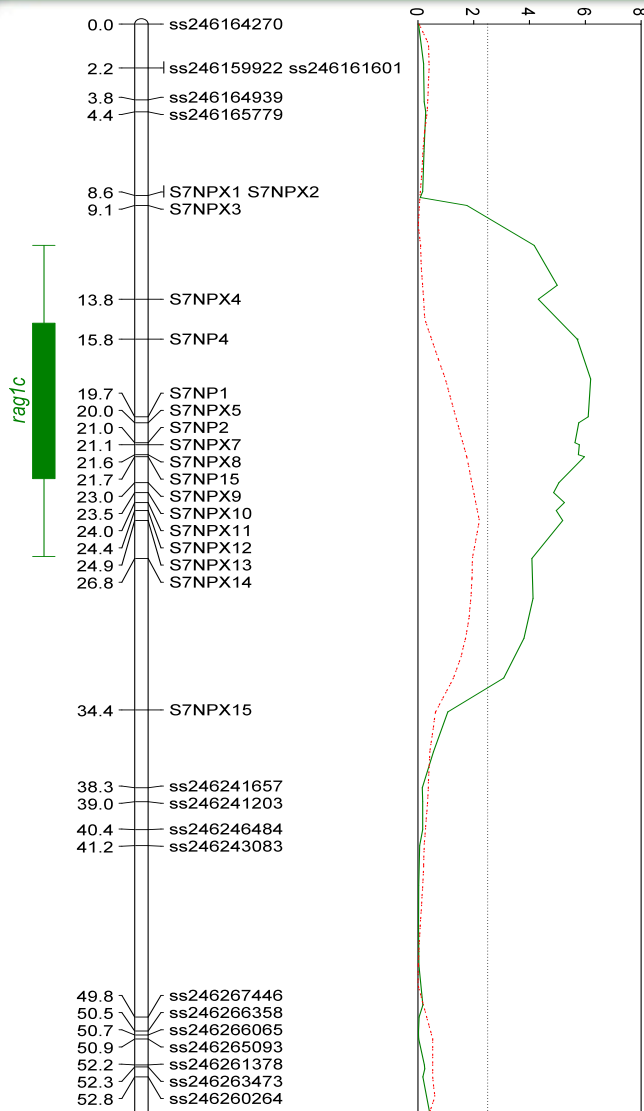
# QTL mapping of *rag1c* and *rag4* using 228 F<sub>3</sub>-derived lines in PI 567541B



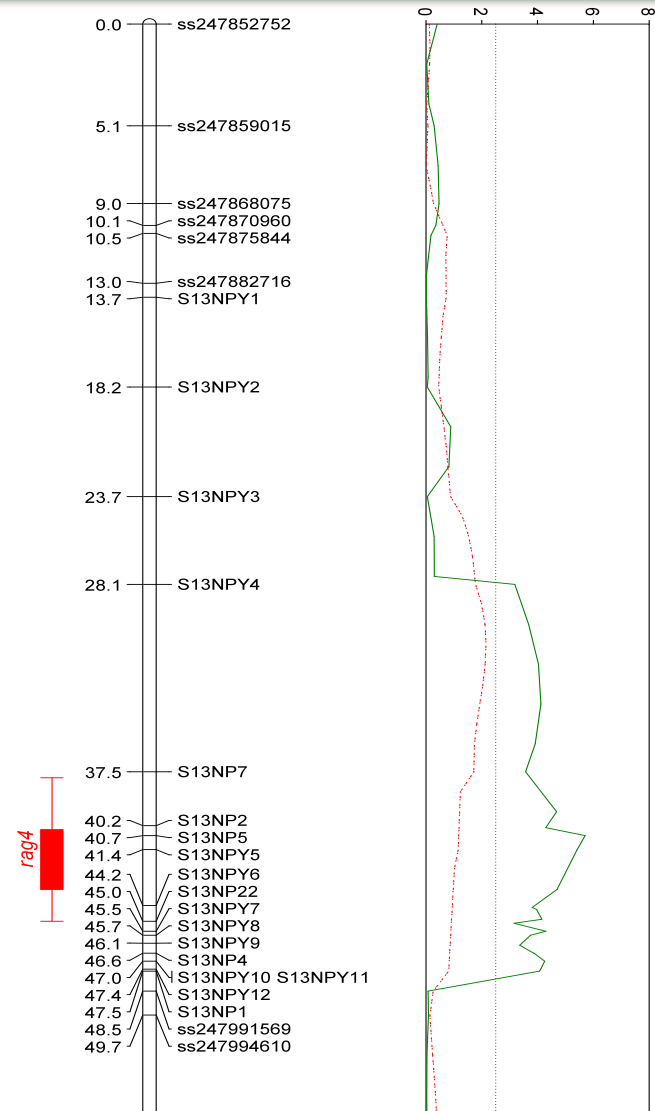


# Updated QTL mapping of *rag1c* and *rag4* in PI 567541B

**R<sup>2</sup> 19.9**



**Chromosome 7**



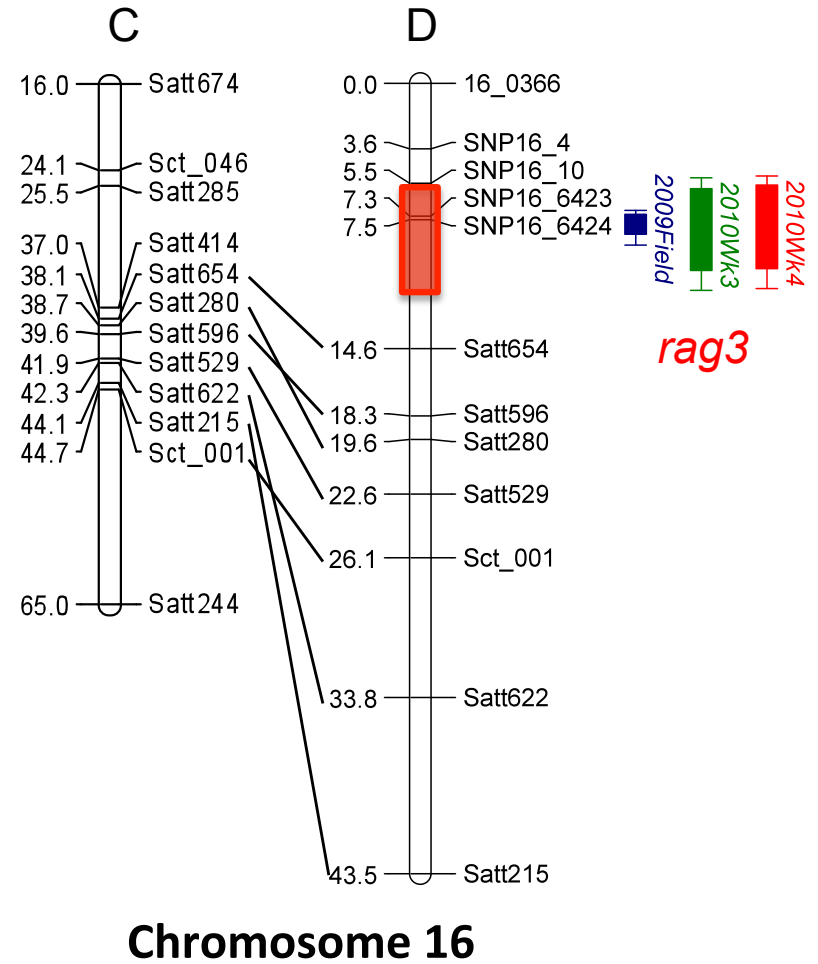
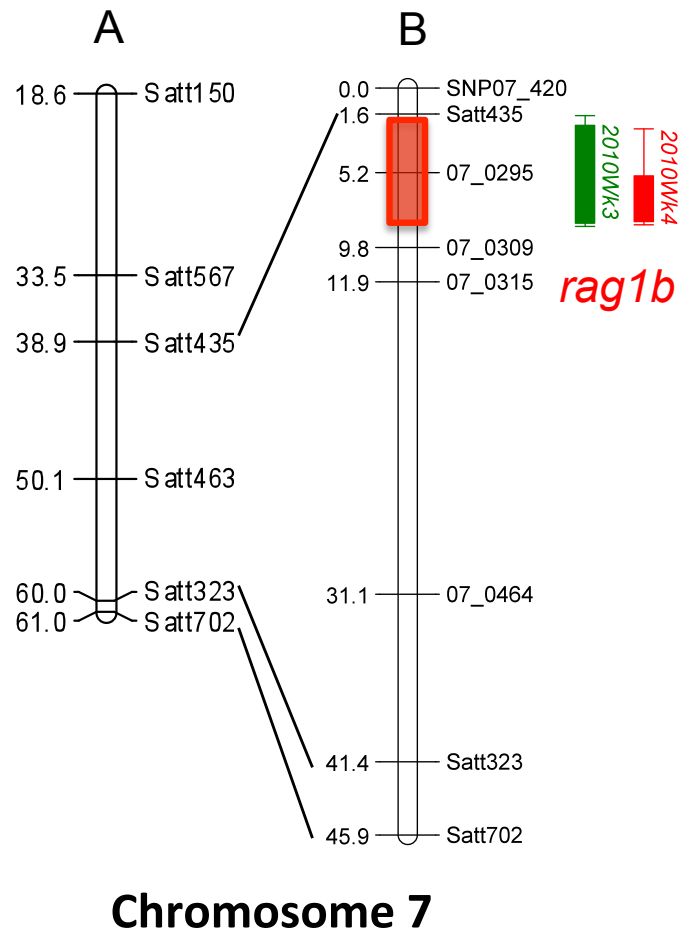
**Chromosome 13**

**R<sup>2</sup> 17.2**

(unpublished data)



# QTL mapping of *rag1b* and *rag3* using 282 F<sub>4</sub>-derived lines



(unpublished data)

# Phenotyping Methods

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# Phenotyping Methods

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- Inoculation: two aphids/plant at the V1 stage;
- Resistance level scale:
  - 0; 0.5; 1.0; 1.5; 2.0; 2.5; 3.0; 3.5; 4.0



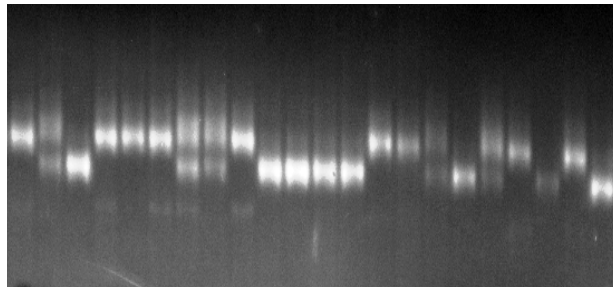
Mensah et al., 2005





# Genotyping Methods

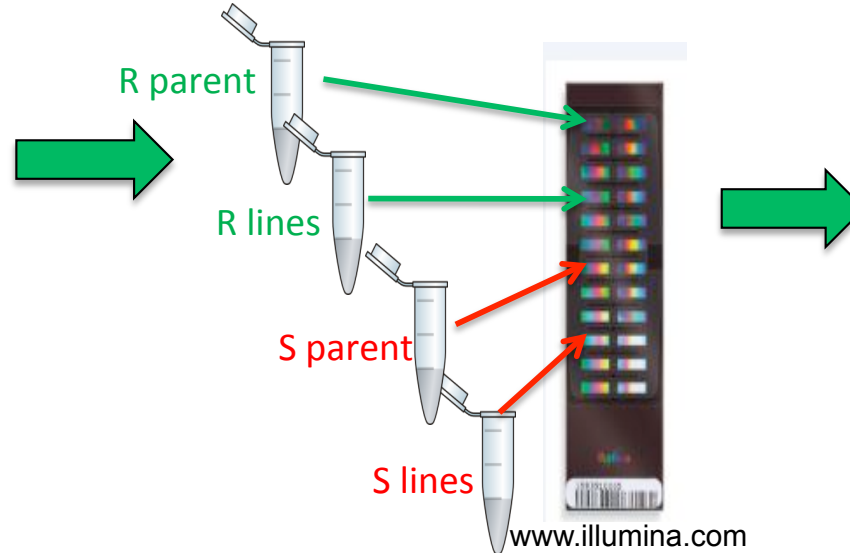
## SSR markers



1,015 markers(Song et al., 2004)  
33,065 markers(Song et al., 2010)

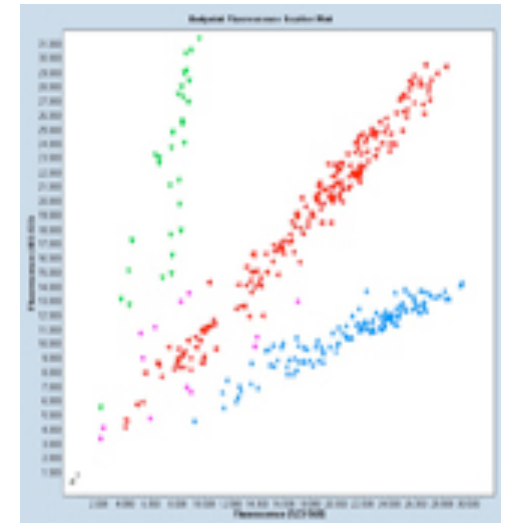
- Bulk segregant analysis
- QTL mapping using RILs

## SoySNP50 iSelect SNP Chip (Song et al., 2011)



- Bulk segregant analysis
- Fine mapping using identified recombinants

## TaqMan® Assay (Endpoint Genotyping)



Lightcycler 480®, Roche Applied Science

- Screening for recombinants
- QTL mapping



# *Rag3* Fine Mapping Populations

090089

[E07048 x E10902]

1,889 F<sub>2</sub>

Quick extracted DNA

SNP 1  
SNP 5  
SNP 20

65 Lines with recombination events

F<sub>2:3</sub> plants for SA test in greenhouse 2011

27 Recombinants to Infinium assay

1 Segregating line for SA test of F<sub>3:4</sub> plants in Spring 2012

090104

[E10905 x E07048]

1,913 F<sub>2</sub>

Quick extracted DNA

SNP 16-4  
SNP 16-10  
SNP 16\_6424

37 Lines with recombination events

F<sub>2:3</sub> plants for SA test in greenhouse 2011

33 Recombinants to Infinium assay

1 Segregating line for SA test of F<sub>3:4</sub> plants in Spring 2012



# *rag3* Fine Mapping Populations

090004

[E00003 x (E00003 x PI98B)]

2,184 BC<sub>1</sub>F<sub>2</sub>

Quick extracted DNA

SNP 16-4  
SNP 16-10  
SNP 16\_6424

107 Lines with recombination events

F<sub>2:3</sub> plants for SA test in field 2011

34 Recombinants to Infinium assay

3 Segregating lines for SA test of F<sub>3:4</sub> plants in Spring 2012

090068

[Skylla x (Skylla x PI98B)]

1,784 BC<sub>1</sub>F<sub>2</sub>

Quick extracted DNA

SNP 16-4  
SNP 16-10  
SNP 16\_6424

94 Lines with recombination events

F<sub>2:3</sub> plants for SA test in field 2011

21 Recombinants to Infinium assay

3 Segregating lines for SA test of F<sub>3:4</sub> plants in Spring 2012

070063

[IA2070 x (Titan x PI98B)]

282 F<sub>4:6</sub>

Initial mapping markers

13 Lines with recombination events

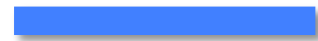
SA resistance testing of progenies

Infinium assay

# SNP Markers from SoySNP50 iSelect Infinium Assay

Lines	Gen	Phenotype	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP	SNP
			1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19
2-499-2	F <sub>2:3</sub>	Resistant	S	S	S	S	S	S	S	S	S	S	S	S	R	R	R	R	R	R	R
2-277-6	F <sub>2:3</sub>	Resistant	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	S	S	S

*Rag3*



04-2-653	F <sub>2:3</sub>	Segregating	S	S	H	H	H	H	H	H	H	H	H	H	H	H	H	H	H	H	
63-5-64	F <sub>4:7</sub>	Susceptible	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S
04-2-229	F <sub>2:3</sub>	Segregating	R	R	R	R	R	H	H	H	H	H	H	H	H	H	H	H	H	H	H
04-2-471	F <sub>2:3</sub>	Resistant	H	H	H	H	H	R	R	R	R	R	R	R	R	R	R	R	R	R	S
04-2-742	F <sub>2:3</sub>	Resistant	H	H	H	H	H	R	R	R	R	R	R	R	R	R	R	R	R	R	S
63-4-29	F <sub>4:7</sub>	Resistant	S	S	S	S	S	R	R	R	R	R	R	R	R	R	R	R	R	R	R
63-1-18	F <sub>4:7</sub>	Susceptible	R	R	R	R	R	R	R	S	S	S	S	S	S	S	S	S	S	S	S
04-2-466	F <sub>2:3</sub>	Resistant	H	H	H	H	H	H	H	H	R	R	R	R	R	R	R	R	R	R	S
68-1rem-39	F <sub>2:3</sub>	Segregating	H	H	H	H	H	H	H	H	H	H	H	H	H	H	H	H	R	R	R
68-1rem-168	F <sub>2:3</sub>	Resistant	H	H	H	H	H	H	H	H	H	R	R	R	R	R	R	R	R	R	R
68-5-146	F <sub>2:3</sub>	Segregating	R	R	R	R	R	R	R	R	R	R	H	H	H	H	H	H	H	H	H

*rag3*





# Acknowledgement

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