Soybean Aphid Biotypes:
Understanding Geographic
Distribution
2008-2010

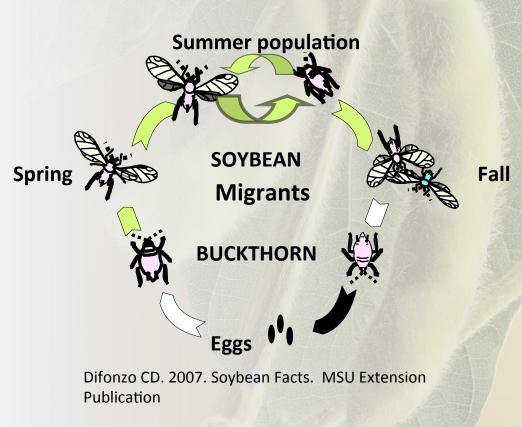
Susannah Cooper Monsanto Company



Soybean Aphid

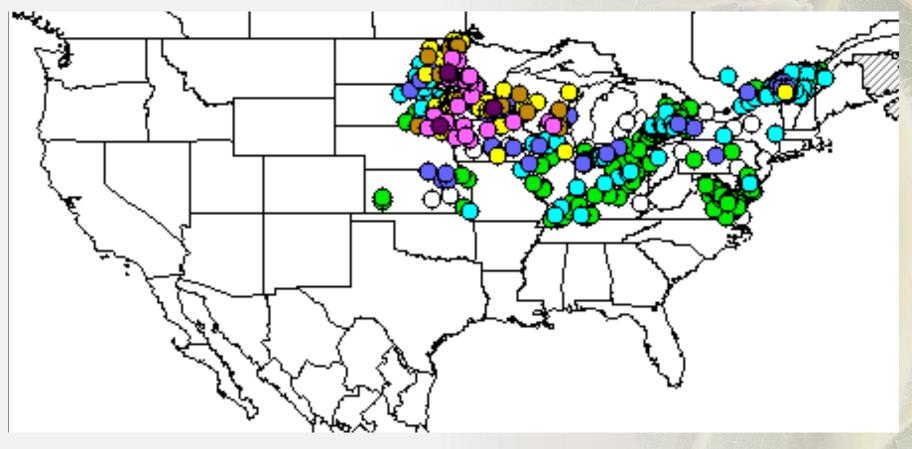
Aphis glycines Matsumura

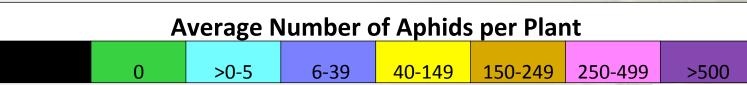
- Soybean aphids confirmed in Wisconsin in 2000 and quickly spread to the North Central region
- Much of N America suitable climate for soybean aphid
- Expansion limited by availability overwintering host, buckthorn
- Viviparous and Oviparous
 - Population can build-up rapidly
- Migrate large distances





Soybean Aphid Distribution





Source: Integrated Pest Management – Pest Information Platform for Extension and Education (PIPE) 8/9/2008 M O N S A N T Q



Management Techniques

- Biological Control
 - Lady beetles
 - Parasitic wasps (Binodoxys communis)
- Scouting
- Insecticide Treatments
- Aphid associated loci
 - Rag1, Rag2, rag3, rag4
 - Insect Biotypes



Insect Biotypes

- The performance of a genotype or group of genotypes of unknown relation on particular host (host differentials) Downie. 2010.J. Insect Science 10:176(1-18)
- Not: A population or group of individuals having the same genotypes



Understanding More about Soybean Aphid Biotypes

- Initiated in 2008
- Collaborated with with public entomologists/ breeders to design and conduct evaluations





Identification of 'Biotypes' in Soybean Aphid

- Reports of aphids proliferating on Rag1 germplasm in OH (2005) & MI (2006)
- 2008: Kim et al. Crop Sci 28:923-928

Table 3. The average number of aphids per plant 10 d after infestation and the plant damage index (PDI) 15 d after infestation with the Ohio isolate in the choice test.

	Soybean genotype	No. of aphids plant ⁻¹ 10 d after infestation	Average PDI 15 d after infestation [†]	
Rag1	Dowling	157a‡	4.2a	
	Dwight	180a	4.0a	Susceptible
	Jackson	110b	4.3a	
	PI 200538	24c	1.2d	
	Williams 82	166a	4.5a	Susceptible
	PI 567541B	38c	2.2c	TO SEE L
	PI 567597C	32c	2.2c	
Raq1	LD05-16611	172a	3.0b	19

[†]The plant damage index (PDI) ranges from 1 (no stunting and leaf distortion) to 5 (severe plant damage).



 $^{^{\}ddagger}$ Means followed by the same letters in a column are not significantly different by the least significant different test (P = 0.05).

Collaborators

- Vaino Poysna and David Hunt
 - Ag Canada: Harrow, ON
- Mike Gray and Ron Estes
 - University of Illinois
- Matt O'Neal
 - lowa State University
- Brian McCornack
 - Kansas State University
- Christian Krupke
 - Purdue
- Dechun Wang
 - Michigan State University

- David Ragsdale
 - University of Minnesota (currently at Texas A &M)
- Deirdre Prischmann-Voldseth
 - N. Dakota State University
- Guo-Liang Jiang and Marci Green
 - S. Dakota State University
- Rouf Mian
 - USDA -Wooster, OH
- Eileen Cullen (2008)/ Ag Stat (2009-10)
 - University of Wisconsin



Objectives

Understand the distribution of aphid biotypes

 Develop a panel of host differentials to characterize aphid biotypes

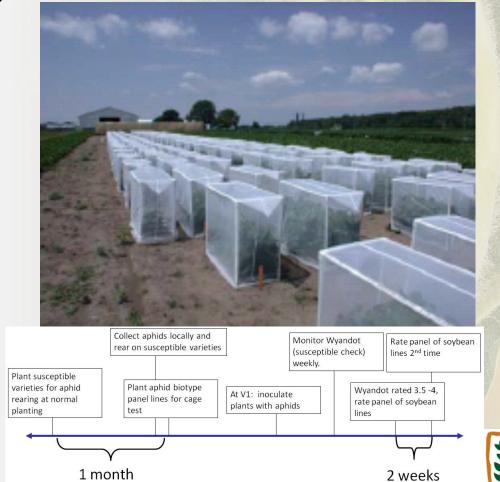


Host Differentials

Line	Institution	Gene	LG	
K1621	KSU		F	
PI567598B	MSU	rag1_b, rag3	M, J	
PI567541B	MSU	rag1_c, rag4	M, F	
Dowling	Univ IL	Rag1	M	
Jackson	Univ IL	Rag	M	
PI243540	USDA-OH	Rag2	F	
PI200538	Univ IL	Rag2	F	
UGA-MON PI	UGA-MON			
CNS	Public	Variable Rxn		
Wyandot	USDA-Ohio	Susceptible Check		ANTO

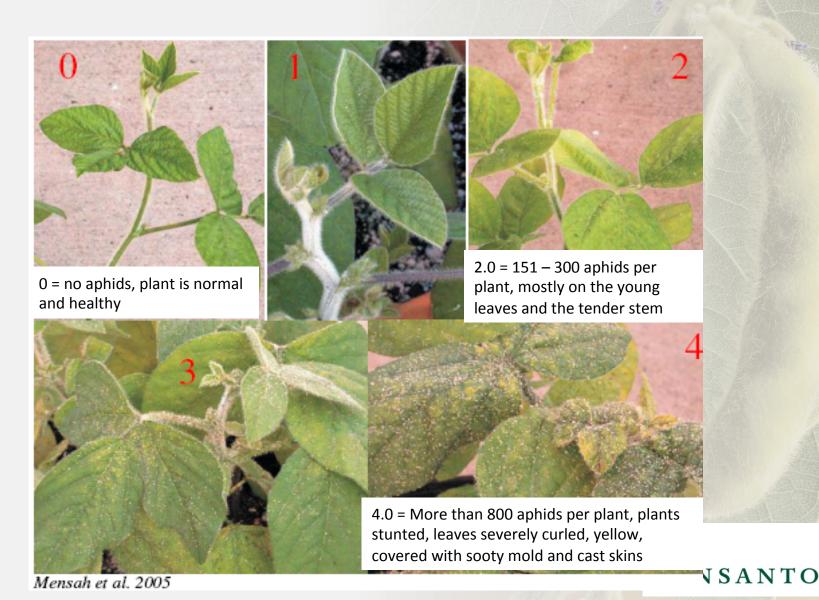
Design

- Protocol based on Mensah et al. 2005
- No-choice evaluation in small field cages
- 15 seed/ cage
- 3 replications/ entry
- 10 entries
- Field collected aphids reared on Wyandot or susceptible variety
- Plant inoculated at V1 and monitored weekly
 - 5 wingless aphids/ plant
- Rating (0-4):
 - Evaluate entire panel when Wyandot reaches 3.5-4
- 8-11 locations

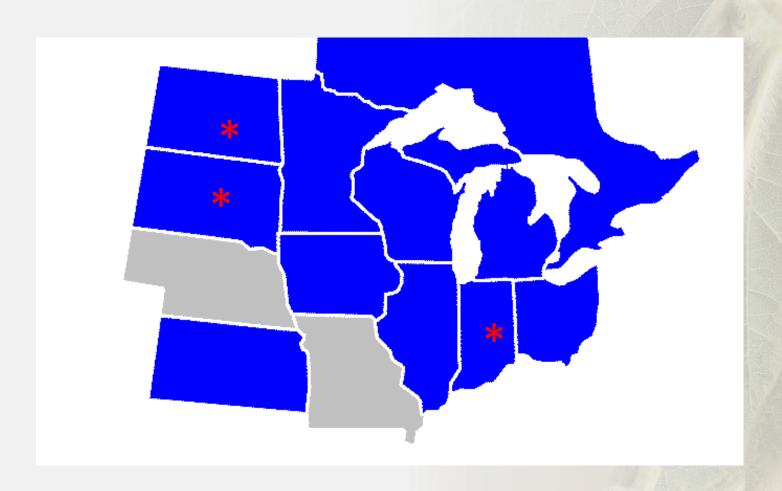


Rating Scale

Mensah et al. 2005



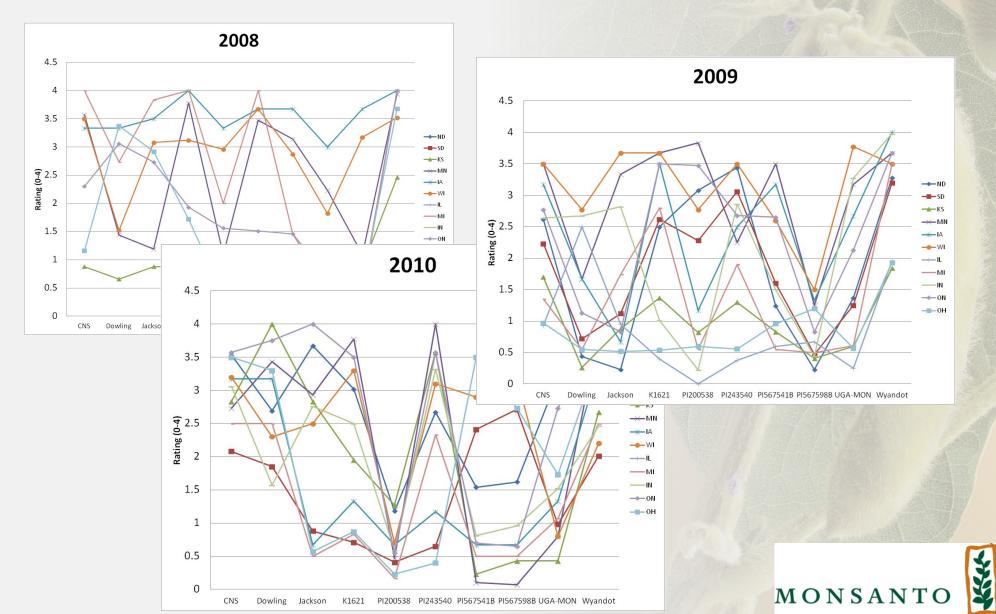
Locations



* Only 2 years of testing



Snapshot of Data



Data Transformation

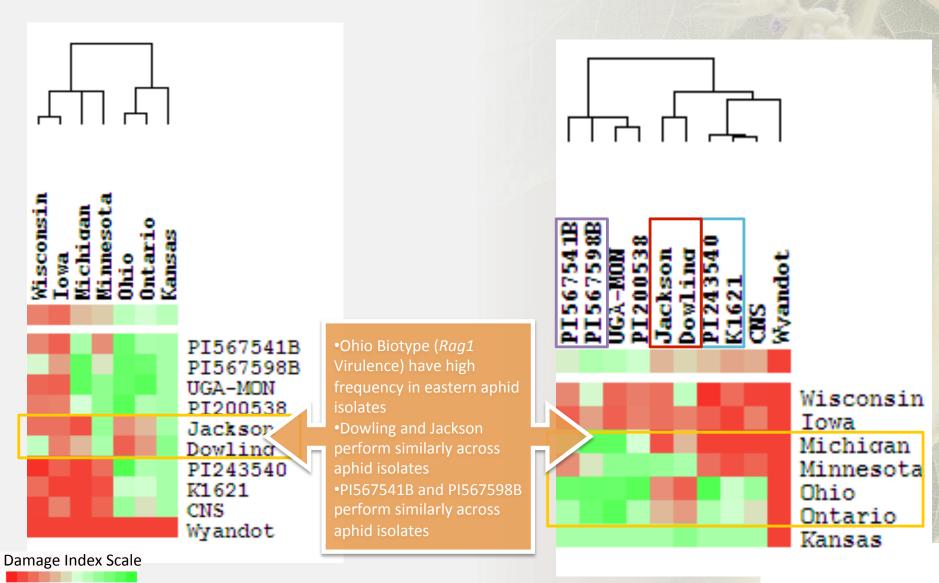
Rating Scale Converted to Damage Index (DI)

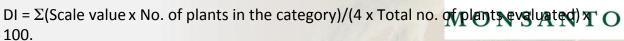
$$DI = \frac{\text{Average Scale Value of Entry}}{\text{Average Scale Value of Susceptible Check}} \times 100$$

Cluster Damage **Analysis** Index Value Color 50 10 40 30 20 20 10 40 -10 60 70 -20 -30 80 90 -40 -50 100 120 -60

- DI Scores ranged from 0-115%
- Cluster Analysis:
 - Gene Cluster© and TreeView© used to visualize trends
 - Transformed ratings by (50-DI)%

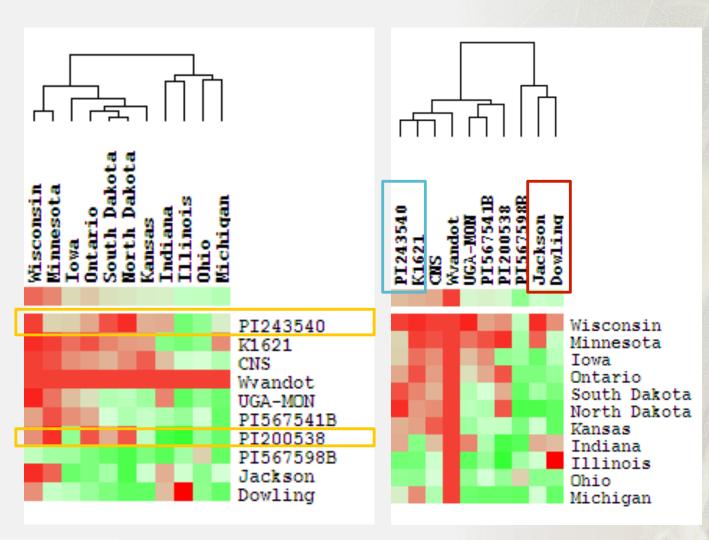








2009



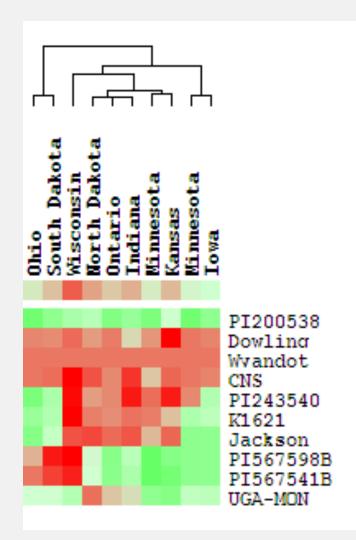
- •Rag1 virulent isolates were less frequent compared 2009
 •Dowling and Jackson perform
- Rag2 (PI243540 and PI200538) sources performed
- •Virulence to PI567598B appears to be rare

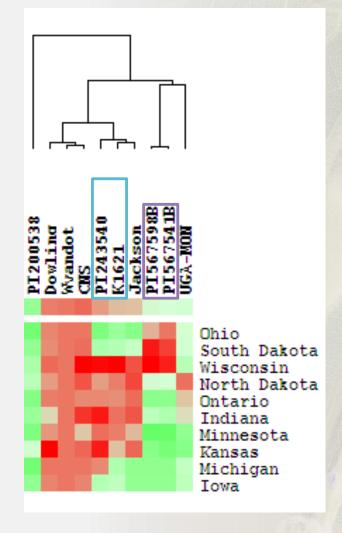
Damage Index Scale

DI = Σ (Scale value x No. of plants in the category)/(4 x Total no. of plants evaluated) x 100. MONSANTO



2010





- •Dowling and Jackson perform differently across aphid isolates
- •Rag1 virulent isolates were common and not limited to eastern sites
- Rag2 (PI243540 and PI200538) sources performed differently across aphid isolates
- •Virulence to PI567598 B appears to be rare





Trends

Line	Gene	Comment
K1621		Clustered together in
PI243540	Rag2	2008-2010
		Performed differently than PI243540
PI200538	Rag2	
PI567598B	rag1_b, rag3	Clustered together in 2008 and 2010; In
		general lowest aphid
PI567541B	rag1_c, rag4	ratings; both 2 recessive genes
Dowling	Rag1	
		Clustered together in
Jackson	Rag	2008-2009; single genes
		Did not consistently
		cluster with other
UGA-MON PI		entries
CNS	Variable Rxn	Often Susceptible

Learnings

- Prevalence of Ohio-biotype (Rag1 tolerance) populations varies year-to-year and geographyto-geography
- Aphid isolates often perform similarly on Dowling and Jackson across geographies
- Aphid isolates performed differently PI243540 and PI200538 across geographies
- Aphid isolates tolerant on PI567598B are also tolerant on PI567541B
- Isolates from different geographies did not consistently cluster together across years



Acknowledgements

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- Ag Stat (2009-10)
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- Marcy Green
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- Brian McCornack
- Rouf Mian
- Matt O'Neal
- Vaino Poysna
- Deirdre Prischmann-Voldseth
- David Ragsdale
- Dechun Wang

Seed

- Rouf Mian
- Dechun Wang
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Monsanto

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