

# Update on the genetic characterization of West Coast populations of *Halyomorpha halys* and adventive populations of its egg parasitoid *Trissolcus japonicus*

Marie Claude BON and Vincent LESIEUR

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Brown Marmorated Stink Bug (BMSB) BioControl Identification Workshop  
BMSB Integrated Pest Management (IPM) Working Group Meeting

# *Update on the genetic characterization of West Coast populations of *Halyomorpha halys* and adventive populations of its egg parasitoid *Trissolcus japonicus**

## **Active collaboration**



- Elijah Talamas
- Matt Buffington

- Kim Hoelmer
- Christine Dieckhoff
- Kathleen Tatman

**AND many others...**

# Update on the genetic characterization of West Coast populations of *Halyomorpha halys* and adventive populations of its egg parasitoid *Trissolcus japonicus*

- Phylogeography of *Halyomorpha halis* in the U.S.
- Genetic diversity of *Trissolcus japonicus*
- Phylogeny of the Asian *Trissolcus*

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## What do we know so far?



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Biol Invasions  
DOI: 10.1007/s10530-013-0510-3

ORIGINAL PAPER

Tracing the origin of US brown marmorated stink bugs,  
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Jiawu Xu · Dina M. Fonseca · George C. Hamilton ·  
Kim A. Hoelmer · Anne L. Nielsen



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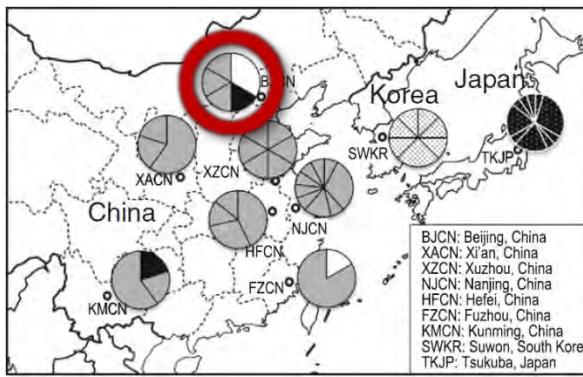
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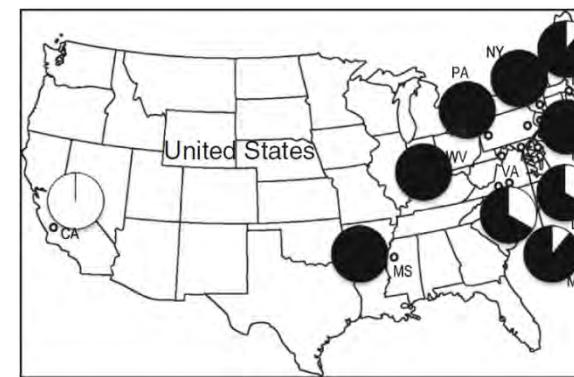
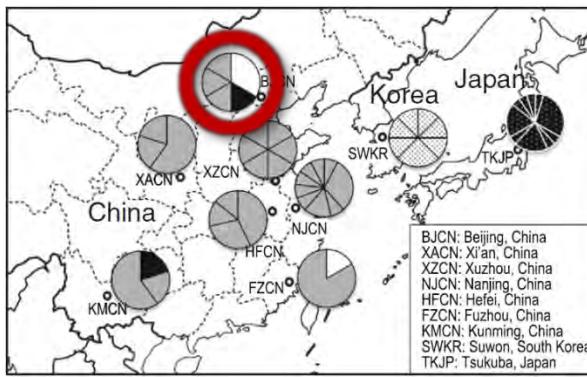
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J Pest Sci (2014) 87:17–28  
DOI 10.1007/s10340-013-0529-3

ORIGINAL PAPER

Occurrence, genetic diversity, and potential pathways of entry  
of *Halyomorpha halys* in newly invaded areas of Canada  
and Switzerland

T. D. Gariepy · T. Haye · H. Fraser ·  
J. Zhang



- A single introduction of small propagules size matches the invasion history in Eastern US
- Beijing area in China being the likely source of introduction
- Similar scenario Gariepy et al (2014)
  - US populations acted as bridgeheads for Canadian populations

## What about Western US?



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Phylogeographic approach

Same methods as Xu et al. (2013)

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3 mtDNA markers

- 
- COII
  - 12S rRNA
  - 12S CR



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Same methods as Xu et al. (2013)

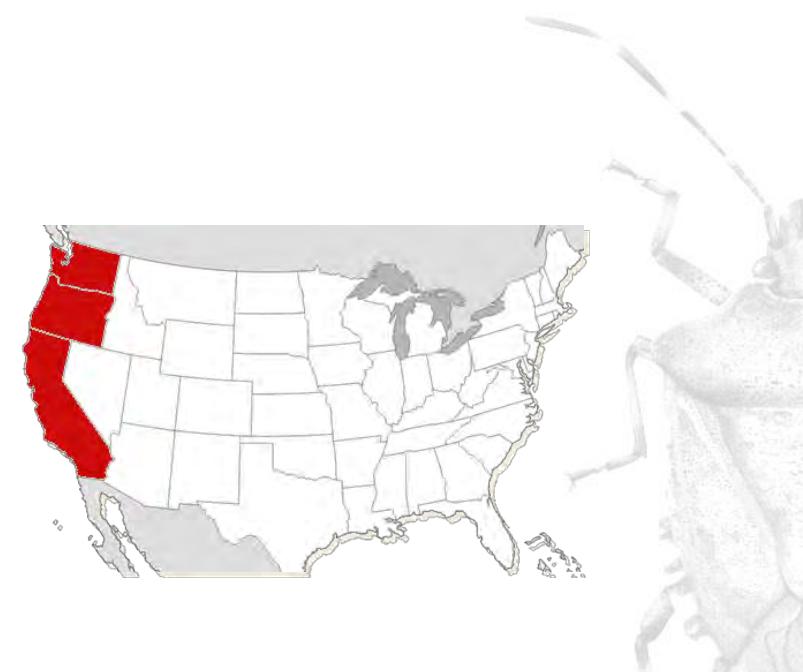


### 3 mtDNA markers

- COII
- 12S rRNA
- 12S CR

### Sampling

- 119 specimens sampled
- 18 sampling locations  
→ Oregon, Washington and California



# Genetic characterization of BMSB and its egg parasitoid

1.North American BMSB

2.North American *Trissolcus*

3.Phylogeny of *Trissolcus*

3

## Genetic diversity measures of BMSB

Geographic area	n	Number of haplotypes	Unique haplotype	Haplotype diversity	Nucleotide diversity
Western US (CA, OR, WA)	119	6	1	0.72	0.00168
Eastern US	55	2	0	0 - 0.67	0 – 0.0006
China	18	26 (2-5 per location)	24	0.81-1	0.0011- 0.0038
Korea (1 site)	8	7	7	0.96	0.0023
Japan (1 site)	21	10	10	0.87	0.0027

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- High genetic diversity in the native range

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● High genetic diversity in the native range

● Significantly lower in eastern US

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- High genetic diversity in the native range
- Higher diversity in western US
- Significantly lower in eastern US

# Genetic characterization of BMSB and its egg parasitoid

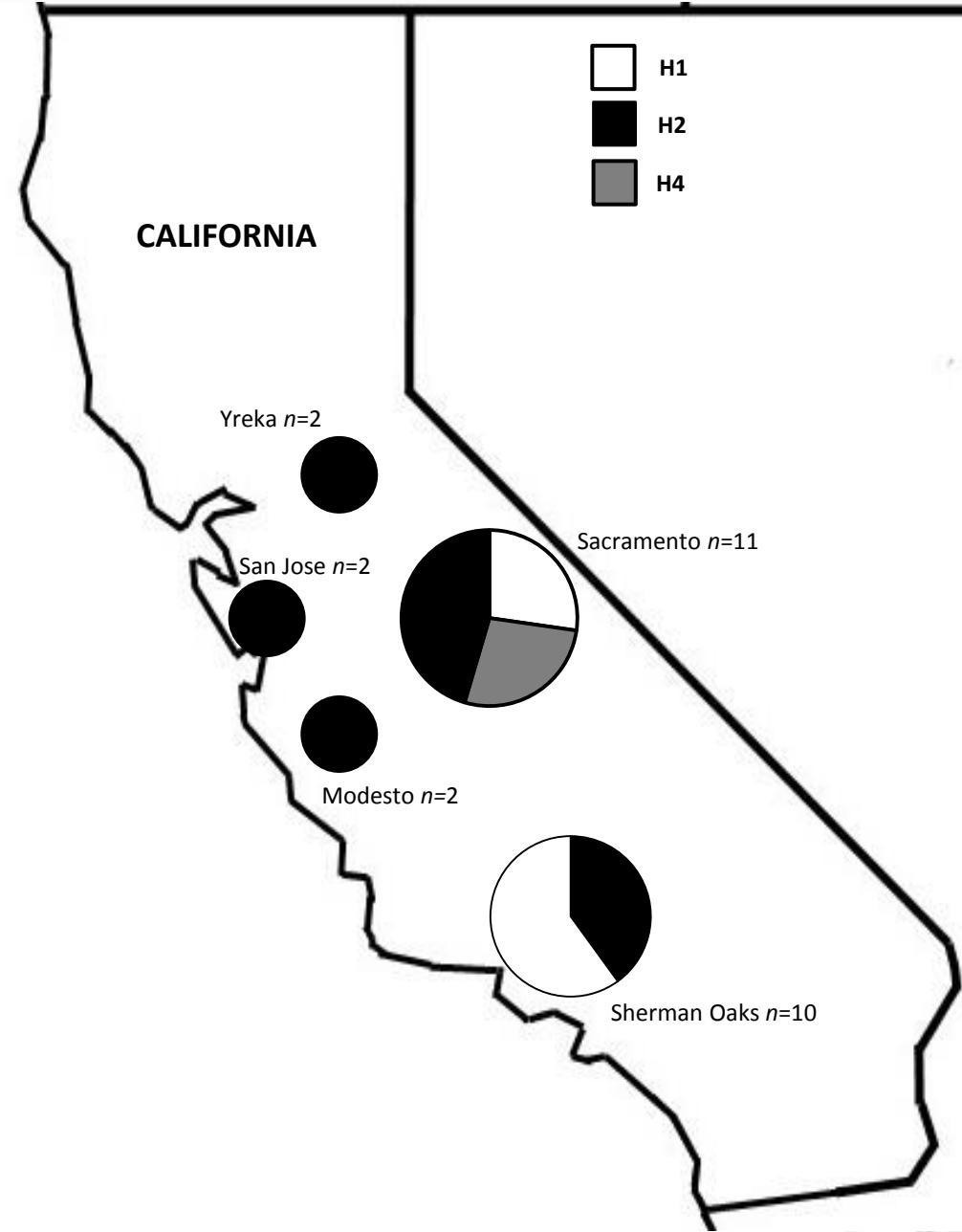
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## California



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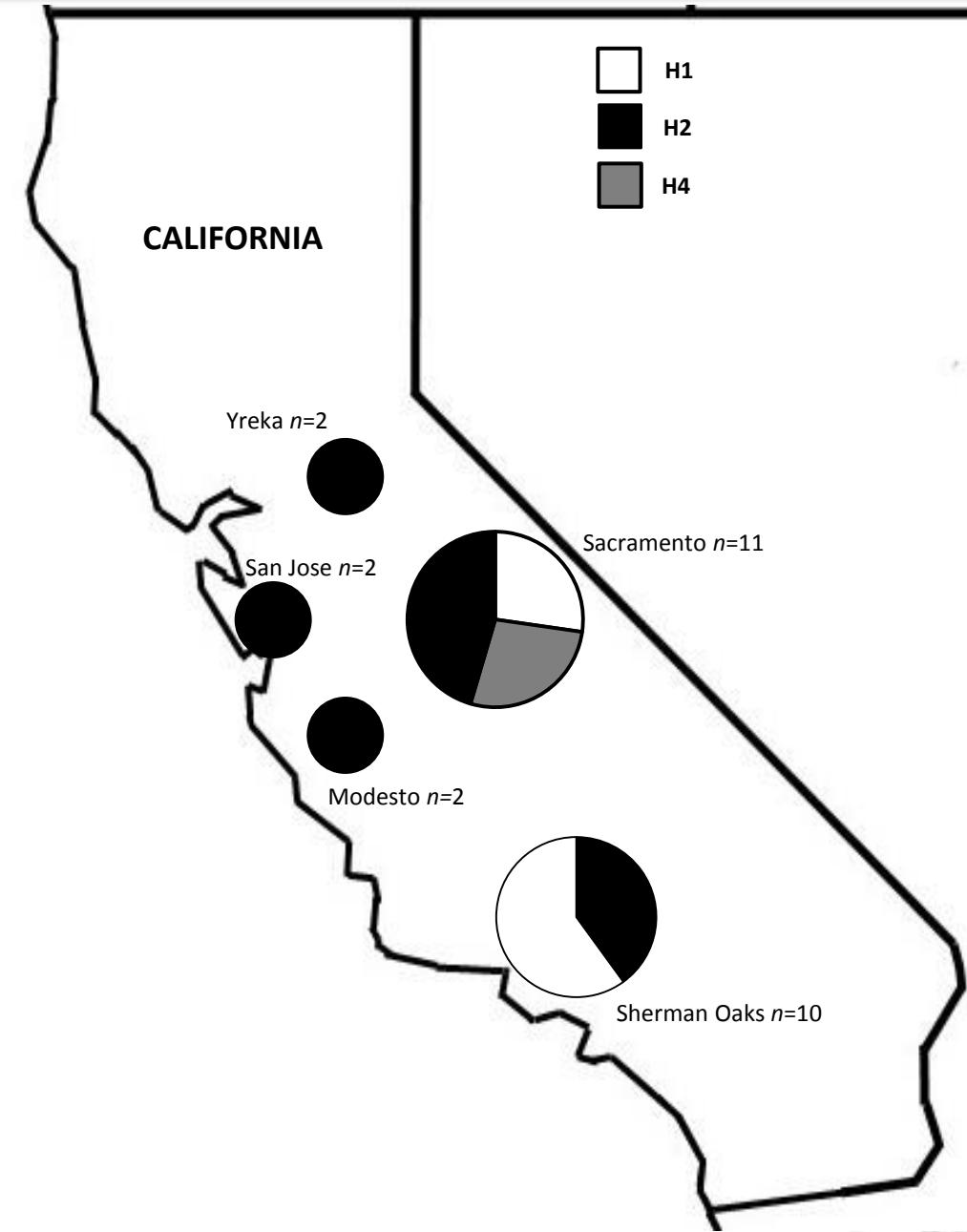
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### California

- 3 mt DNA haplotypes (combined dataset) recovered from 31 specimens
- Prevalence of H1 and H2 (similarly to the pattern observed in East Coast)



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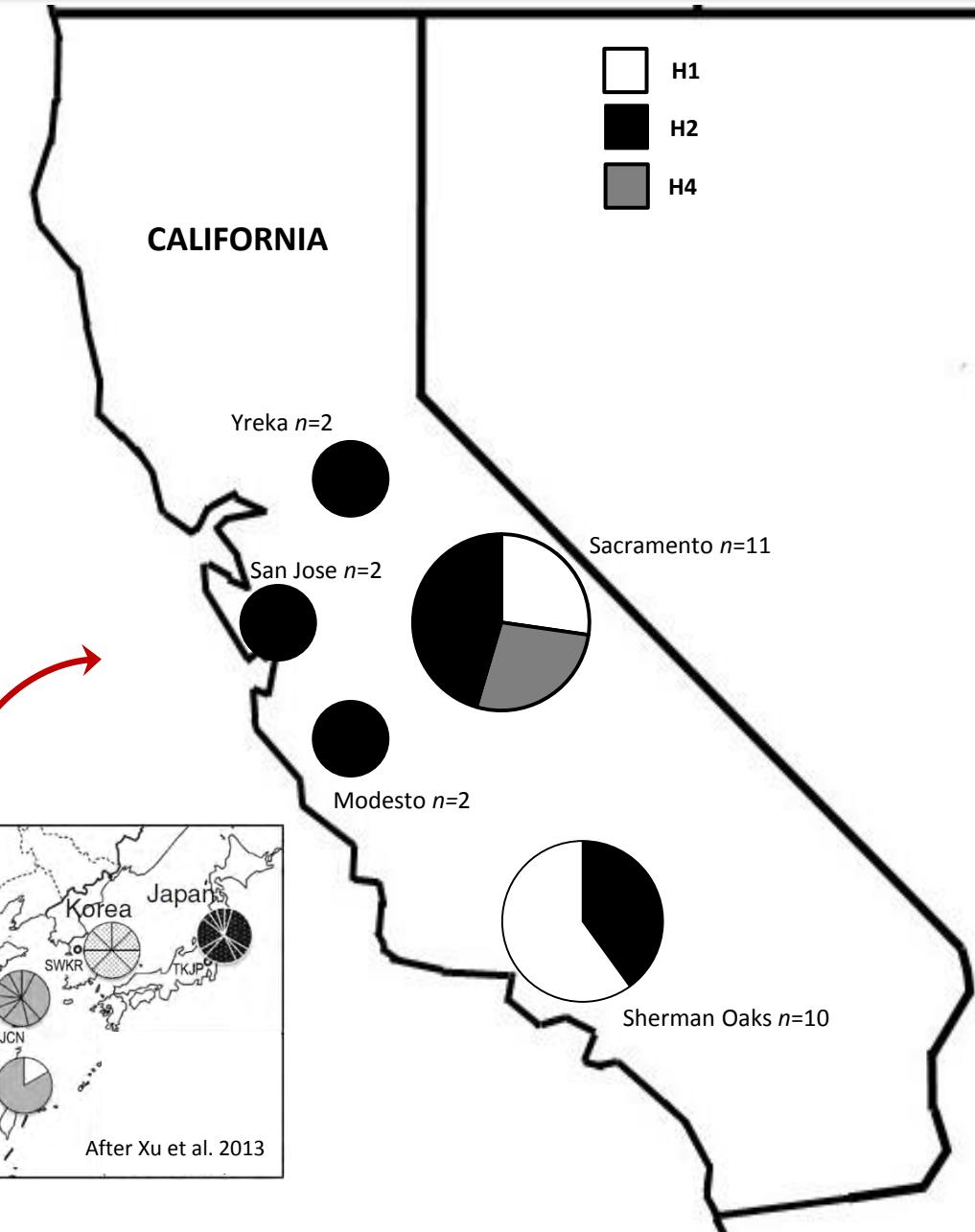
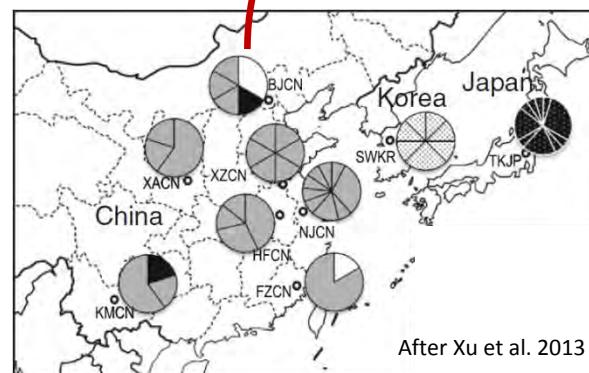
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Beijing area in China:  
the likely source region for California

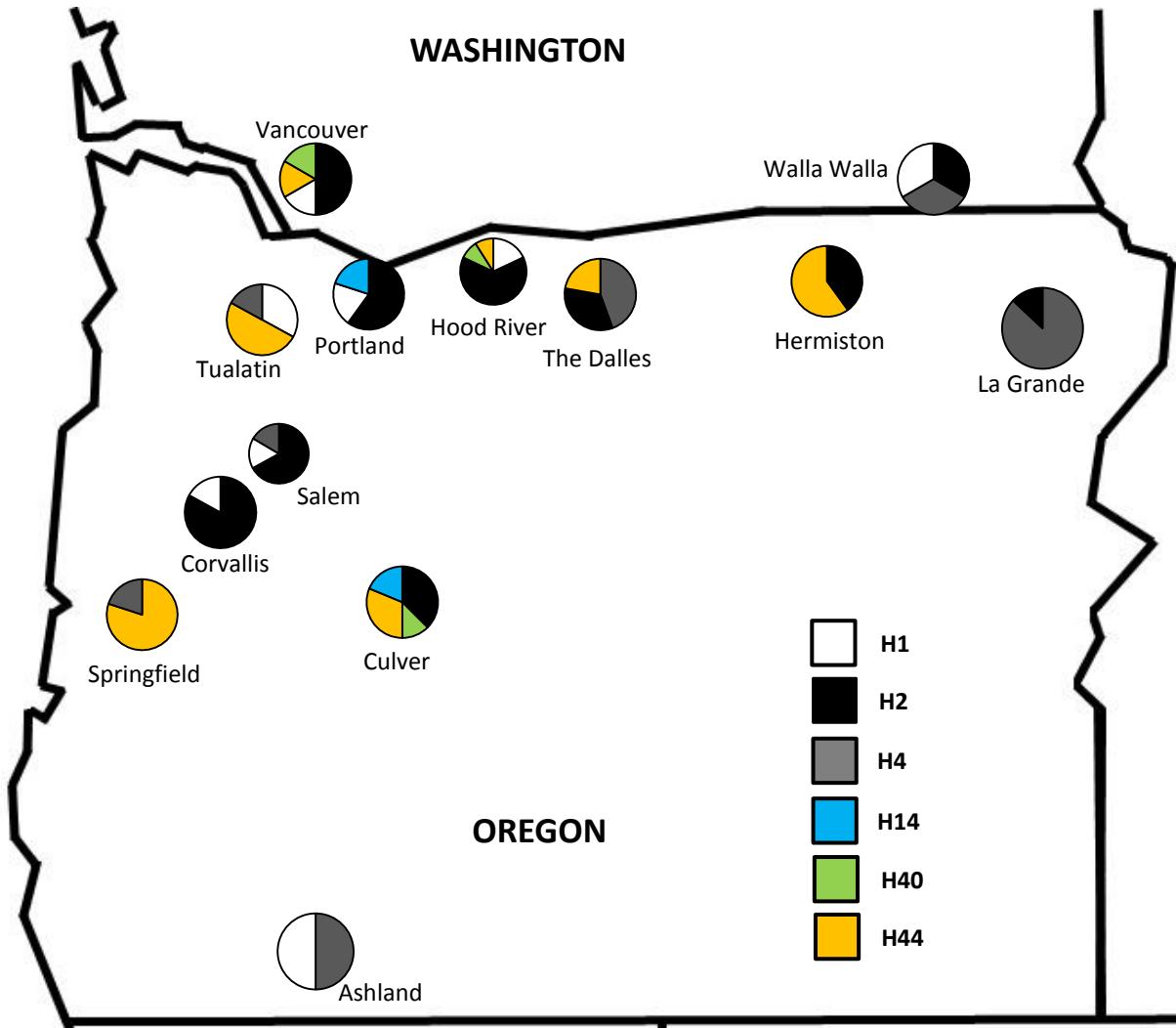


## Washington and Oregon



## Washington and Oregon

- 6 haplotypes recovered from 88 specimens



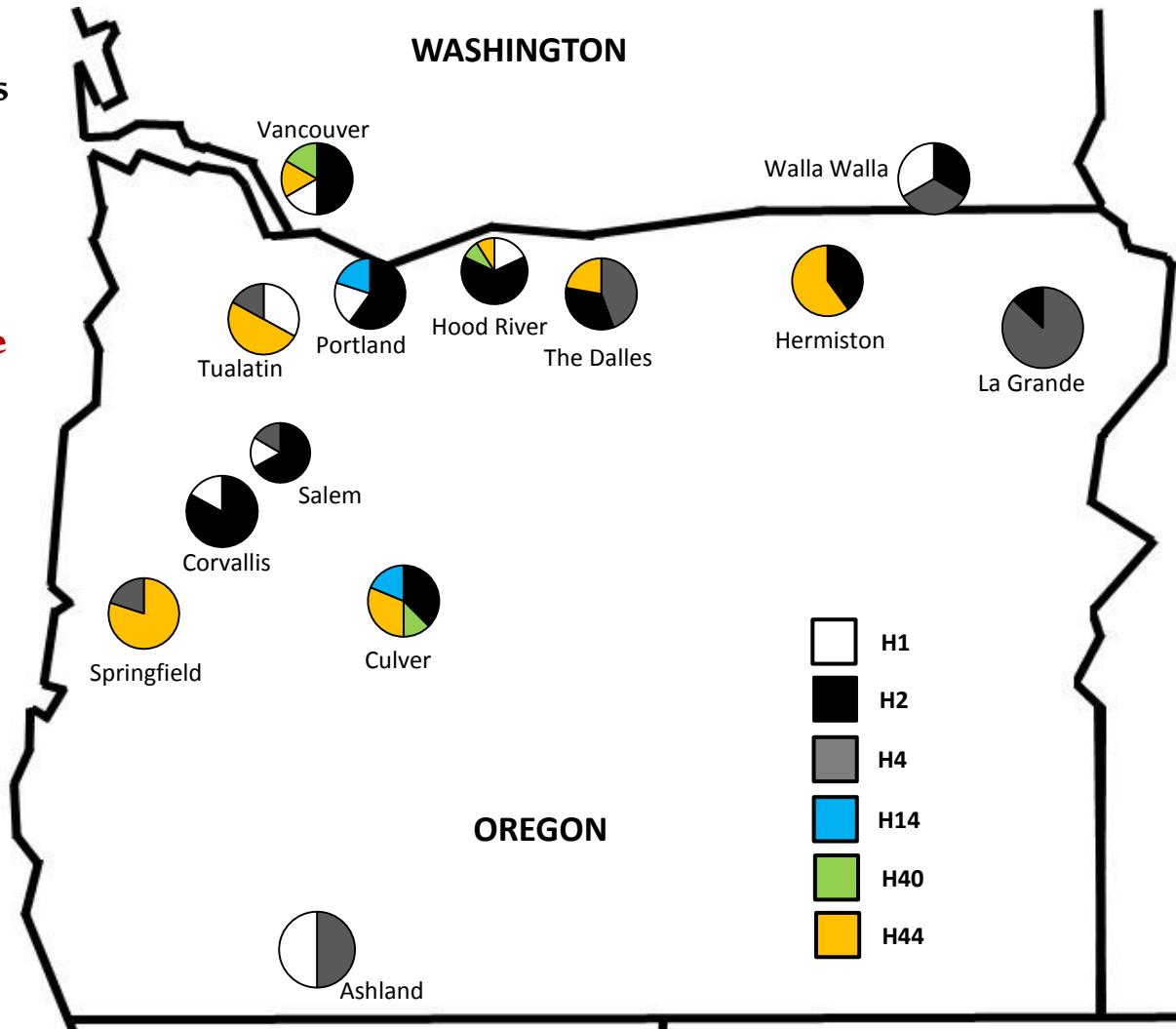
## Washington and Oregon

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- Prevalence of H2 (39%) and to a less extent H44 (21%)



H44 is evidenced for the first time



## Washington and Oregon

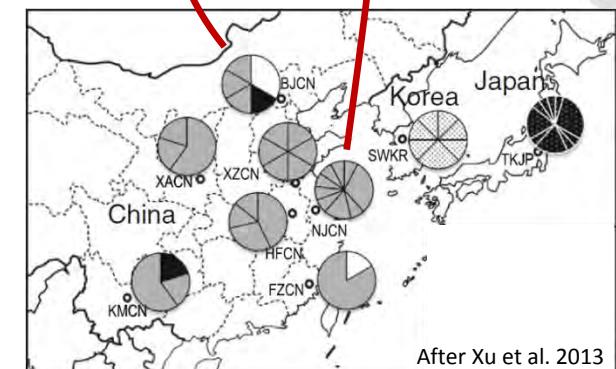
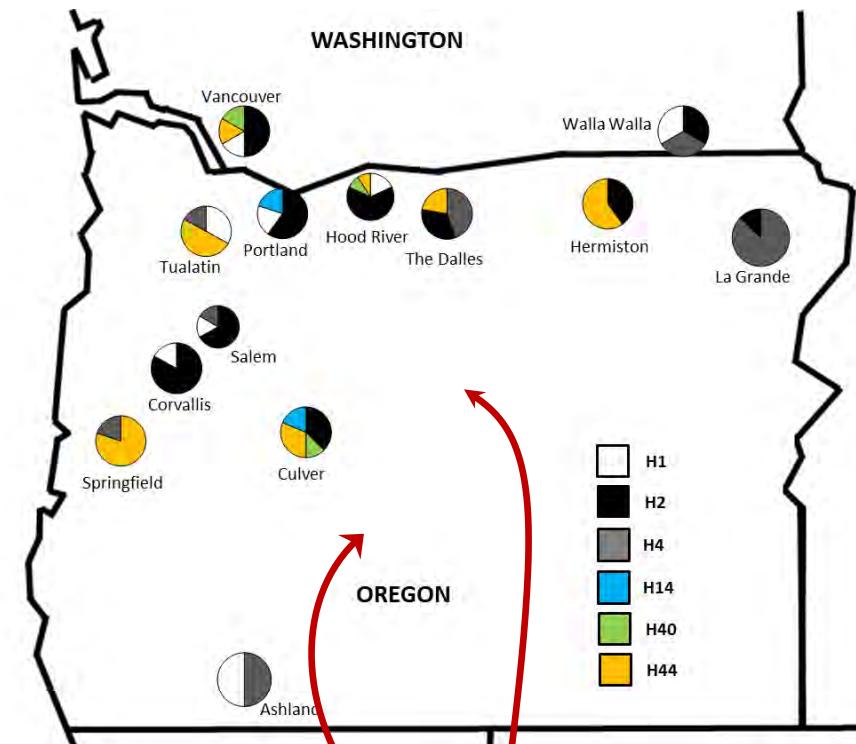
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H1, H2, H4 and H14

Beijing and Nanjing areas in China:  
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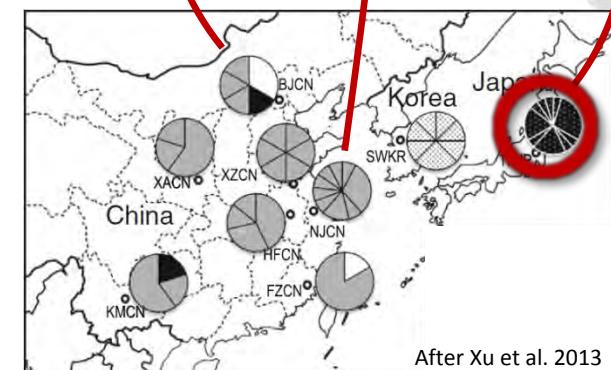
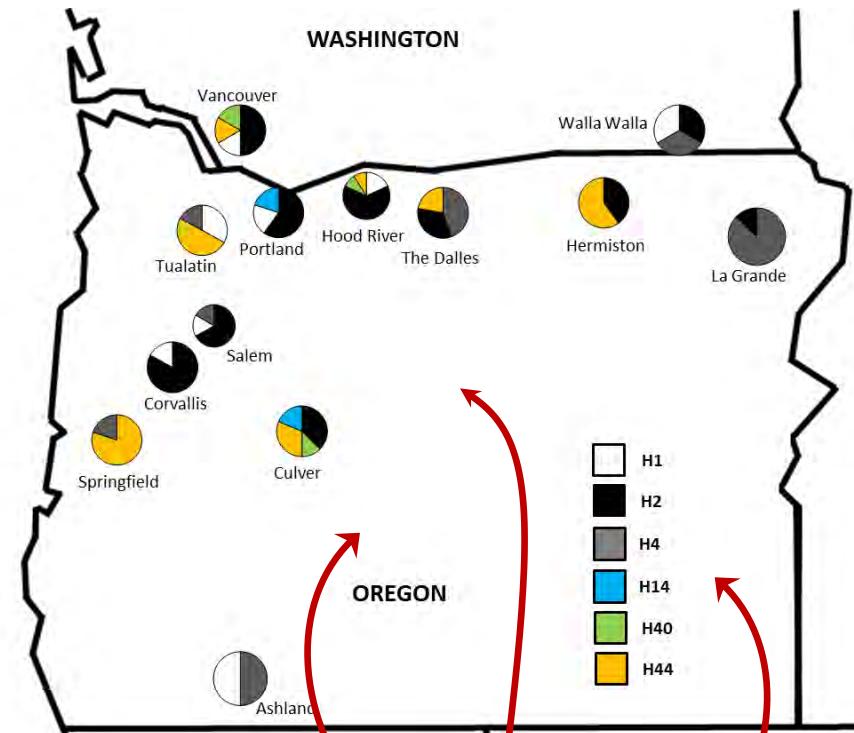
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Beijing and Nanjing areas in China:  
the likely source regions for WA and OR

But not only... Japan

Historical interception from Japan in  
British Columbia

After Cariepy et al. 2014



## Most likely scenario evidenced by the molecular data



Invasion in Western US resulting from multiple sources in Asia

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After Xu et al. 2013



Manuscript in preparation

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JHR 33: 113–117 (2013)  
doi: 10.3897/jhr.33.5627  
[www.pensoft.net/journals/jhr](http://www.pensoft.net/journals/jhr)

RESEARCH ARTICLE



## New synonymy of *Trissolcus halyomorphae* Yang

Elijah J. Talamas<sup>1</sup>, Matthew Buffington<sup>1</sup>, Kim Hoelmer<sup>2</sup>

- Specialist egg parasitoid of BMSB
- Different strains from China, Japan and South Korea in culture
  - Evaluation of its host range and efficacy



Talamas et al. 2015



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Talamas et al. 2015

JHR 43: 119–128 (2015)  
doi: 10.3897/JHR.43.4661  
<http://jhr.pensoft.net>

SHORT COMMUNICATION



## *Trissolcus japonicus* (Ashmead) (Hymenoptera, Scelionidae) emerges in North America

Elijah J. Talamas<sup>1</sup>, Megan V. Herlihy<sup>2</sup>, Christine Dieckhoff<sup>3,4</sup>, Kim A. Hoelmer<sup>4</sup>,  
Matthew L. Buffington<sup>1</sup>, Marie-Claude Bon<sup>5</sup>, Donald C. Weber<sup>2</sup>

A survey conducted in 2014 revealed that *T. japonicus* was already present in the wild

- How did *Trissolcus japonicus* come to this site?
- Accidental quarantine escape?

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## **Comparison of the genetic diversity**



**« Recovered population »**

**Quarantine colonies**



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## Comparison of the genetic diversity



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Quarantine colonies

## Development of microsatellites

Population genetics requires successful development of microsatellites

→ These tools have been developed at the onset of the BMSB project

24 microsatellites

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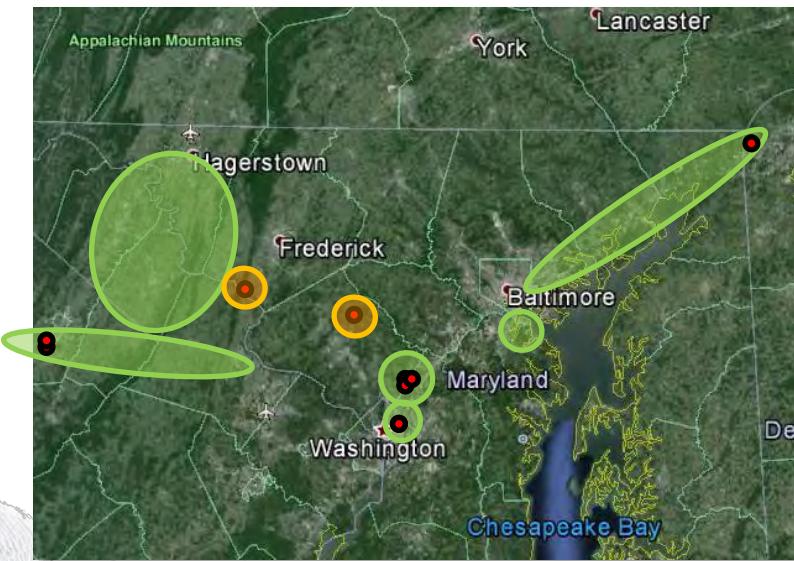
Sentinel egg masses



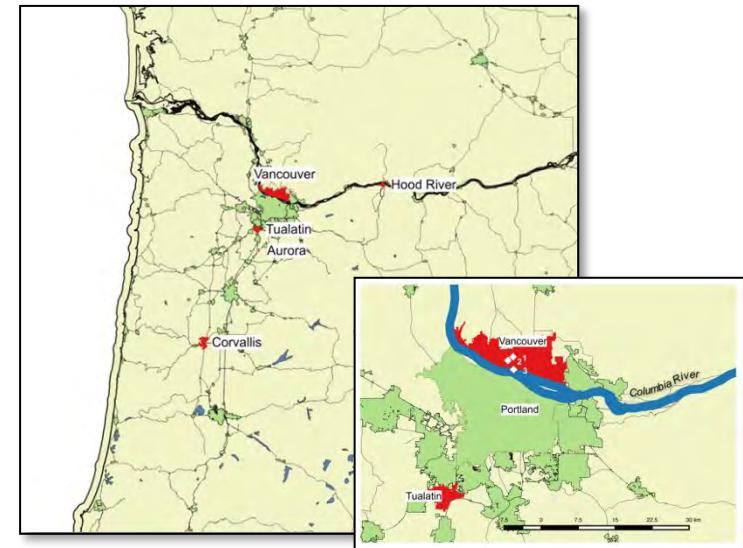
67 individuals

- 2014       $n=38$
- 2015       $n=29$

East Coast Survey



West Coast Survey



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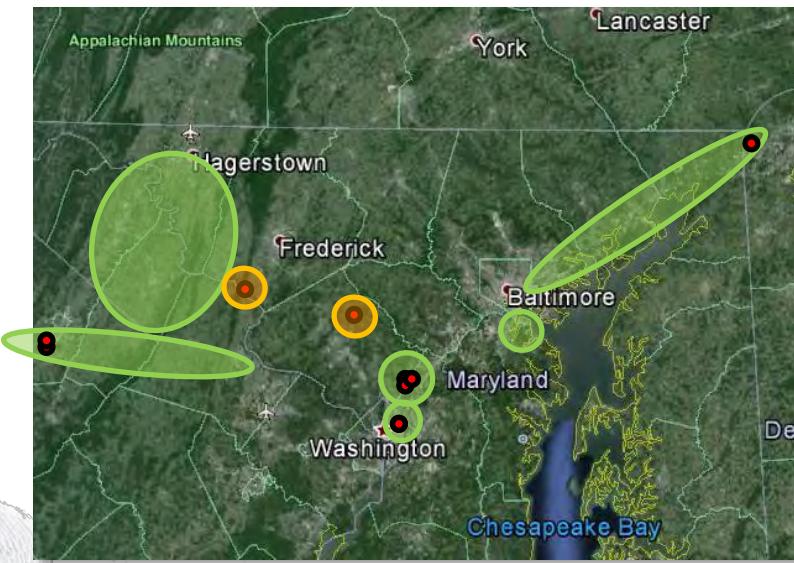
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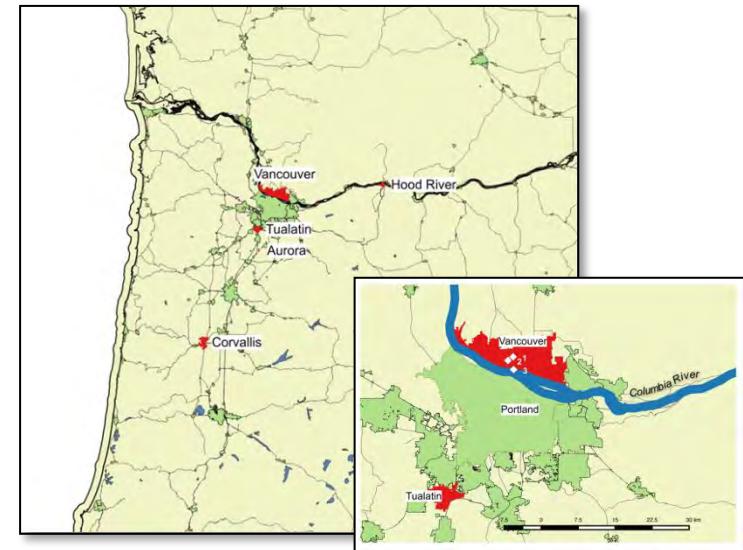
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East Coast Survey



West Coast Survey



Taxonomy

Multilocus genotyping approach (24 microsatellites)

Barcode sequencing (COI)



Wild specimens  
can be identified

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Genus	Species		EBCL id	Sex	Cuid	Label1
Trissolcus	euschistii	fragment	Tj221	?	destructive method	dissected from wild Euschistus EM#O-15-15
Trissolcus	japonicus	diss out	Tj222	f?	destructive method	BIIR sentinel ex Thyanta on Ash#NS-13-15
Trissolcus	japonicus		Tj223	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj224	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
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Trissolcus	japonicus		Tj226	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
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Trissolcus	japonicus	partial	Tj230	?	destructive method	WC 1783 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj231	?	destructive method	WC 11810 BARC, M. Herlihy
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Trissolcus	japonicus	partial	Tj233	?	destructive method	WC10720 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj234	?	destructive method	WC11(1)8 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj235	?	destructive method	WC 3831
Trissolcus	japonicus	partial	Tj236	?	destructive method	OE283
Trissolcus	japonicus	partial	Tj237	?	destructive method	ON5817
Trissolcus	japonicus	partial	Tj238	?	destructive method	WC10810
Trissolcus	japonicus		Tj239	f	non invasive method	WC10727 BARC, M. Herlihy
Trissolcus	japonicus		Tj240	f	non invasive method	OE5720 BARC, M. Herlihy
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Trissolcus	sp closely related		Tj249	f	non invasive method	BARC 15 Aug 2015
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Trissolcus	japonicus		Tj251	f	non invasive method	BARC EM#6, 17 June 2015
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## ● Prevalence of *T. japonicus* in the populations recovered

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# Genetic characterization of BMSB and its egg parasitoid

1.North American BMSB

2.North American *Trissolcus*

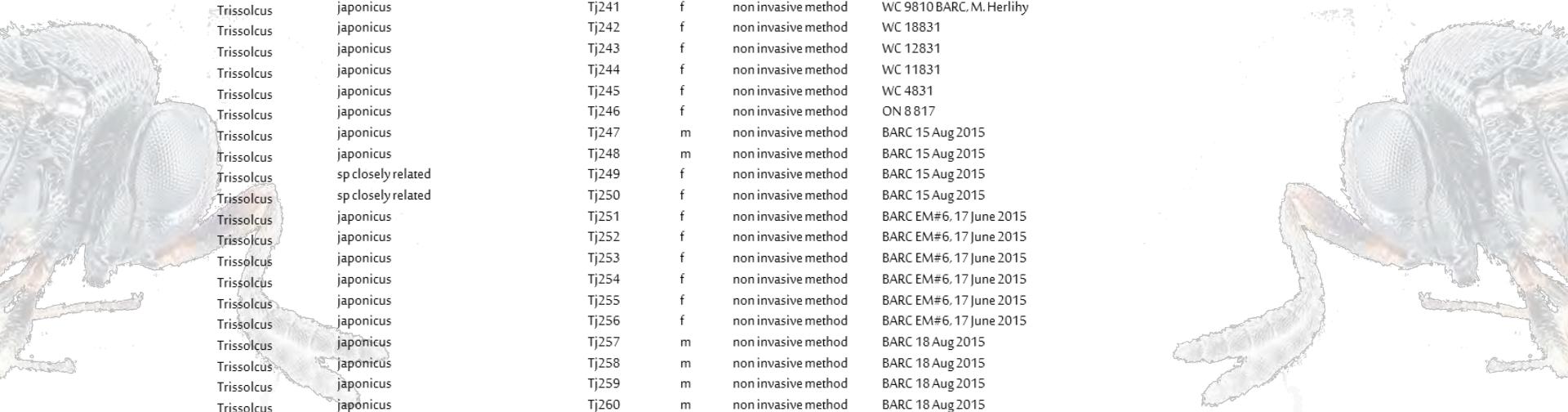
3.Phylogeny of *Trissolcus*

## ● Prevalence of *T. japonicus* in the populations recovered

## ● *Trissolcus euphorbiae*= North American species



Genus	Species		EBCL id	Sex	Cuid	Label1
Trissolcus	euphorbiae	fragment	Tj221	?	destructive method	dissected from wild Euphorbia EM#O-15-15
Trissolcus	japonicus	diss out	Tj222	?	destructive method	BIIR sentinel ex Thyanta on Ash#NS-13-15
Trissolcus	japonicus		Tj223	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj224	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj225	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj226	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj227	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj228	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus		Tj229	m	destructive method	BIIR ex P. mac sentinel EM on ash tree August 2015 possible sentinel lab contaminant
Trissolcus	japonicus	partial	Tj230	?	destructive method	WC 1783 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj231	?	destructive method	WC 11810 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj232	?	destructive method	DE? 783 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj233	?	destructive method	WC10720 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj234	?	destructive method	WC11(1)8 BARC, M. Herlihy
Trissolcus	japonicus	partial	Tj235	?	destructive method	WC 3831
Trissolcus	japonicus	partial	Tj236	?	destructive method	OE283
Trissolcus	japonicus	partial	Tj237	?	destructive method	ON5817
Trissolcus	japonicus	partial	Tj238	?	destructive method	WC10810
Trissolcus	japonicus		Tj239	f	non invasive method	WC10727 BARC, M. Herlihy
Trissolcus	japonicus		Tj240	f	non invasive method	OE5720 BARC, M. Herlihy
Trissolcus	japonicus		Tj241	f	non invasive method	WC 9810 BARC, M. Herlihy
Trissolcus	japonicus		Tj242	f	non invasive method	WC 18831
Trissolcus	japonicus		Tj243	f	non invasive method	WC 12831
Trissolcus	japonicus		Tj244	f	non invasive method	WC 11831
Trissolcus	japonicus		Tj245	f	non invasive method	WC 4831
Trissolcus	japonicus		Tj246	f	non invasive method	ON 8817
Trissolcus	japonicus		Tj247	m	non invasive method	BARC 15 Aug 2015
Trissolcus	japonicus		Tj248	m	non invasive method	BARC 15 Aug 2015
Trissolcus	sp closely related		Tj249	f	non invasive method	BARC 15 Aug 2015
Trissolcus	sp closely related		Tj250	f	non invasive method	BARC 15 Aug 2015
Trissolcus	japonicus		Tj251	f	non invasive method	BARC EM#6, 17 June 2015
Trissolcus	japonicus		Tj252	f	non invasive method	BARC EM#6, 17 June 2015
Trissolcus	japonicus		Tj253	f	non invasive method	BARC EM#6, 17 June 2015
Trissolcus	japonicus		Tj254	f	non invasive method	BARC EM#6, 17 June 2015
Trissolcus	japonicus		Tj255	f	non invasive method	BARC EM#6, 17 June 2015
Trissolcus	japonicus		Tj256	f	non invasive method	BARC EM#6, 17 June 2015
Trissolcus	japonicus		Tj257	m	non invasive method	BARC 18 Aug 2015
Trissolcus	japonicus		Tj258	m	non invasive method	BARC 18 Aug 2015
Trissolcus	japonicus		Tj259	m	non invasive method	BARC 18 Aug 2015
Trissolcus	japonicus		Tj260	m	non invasive method	BARC 18 Aug 2015



# Genetic characterization of BMSB and its egg parasitoid

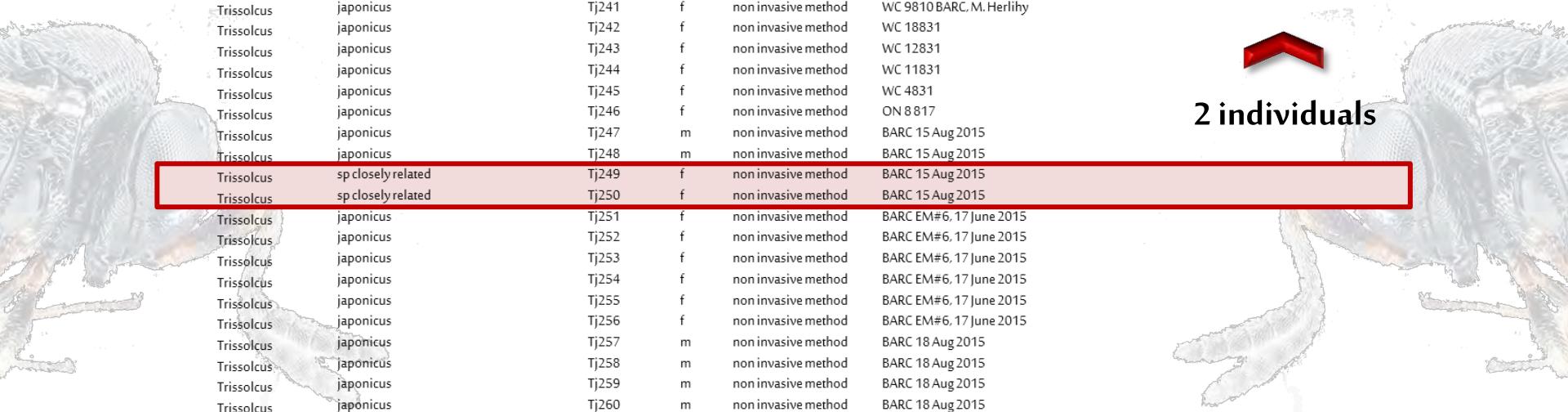
1.North American BMSB

2.North American *Trissolcus*

3.Phylogeny of *Trissolcus*

## Prevalence of *T. japonicus* in the populations recovered

## *Trissolcus euphorchis*= North American species



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Trissolcus	japonicus		Tj259	m	non invasive method	BARC 18 Aug 2015
Trissolcus	japonicus		Tj260	m	non invasive method	BARC 18 Aug 2015

## Evidence of a second lineage of egg parasitoid ?

2 individuals

# Genetic characterization of BMSB and its egg parasitoid

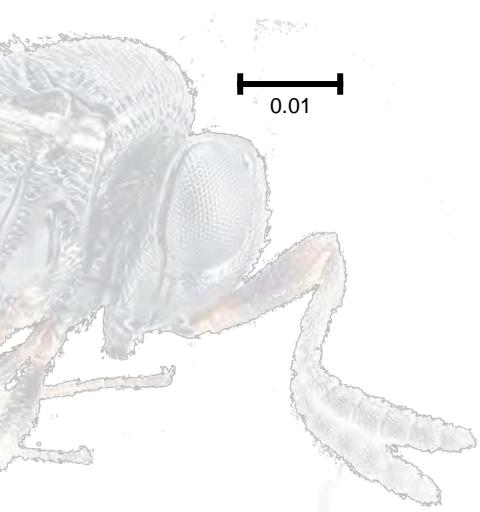
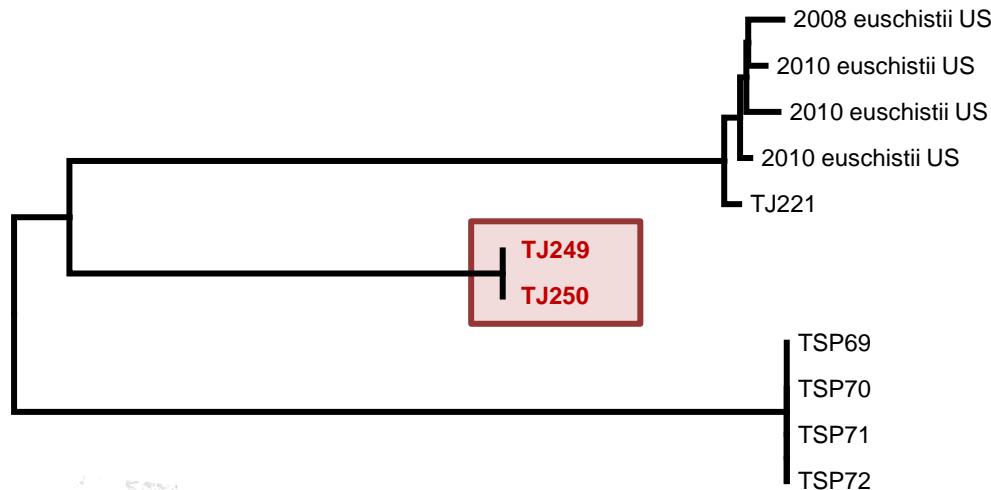
1.North American BMSB

2.North American *Trissolcus*

3.Phylogeny of *Trissolcus*

11

## COI (barcode) results



# Genetic characterization of BMSB and its egg parasitoid

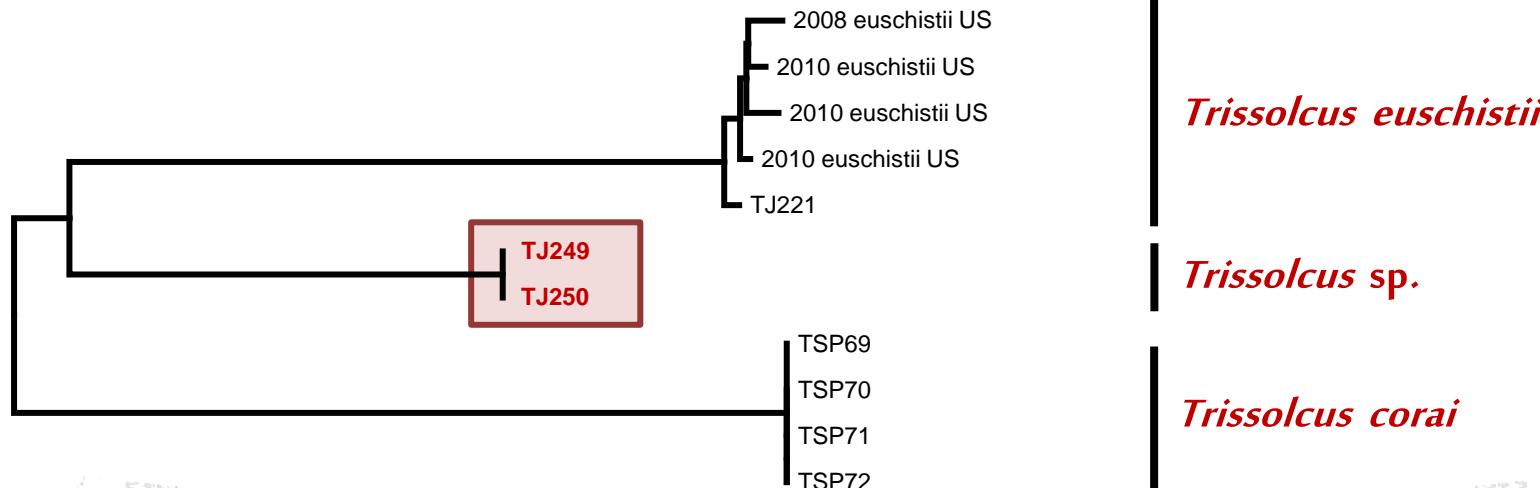
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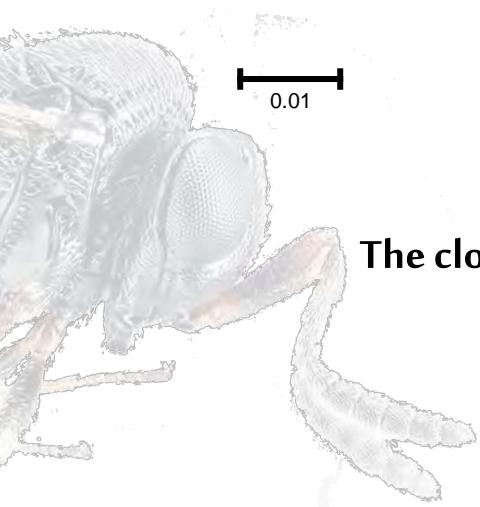
11

## COI (barcode) results



The closest taxa

*Trissolcus euschistii*  
*Trissolcus corai*



# Genetic characterization of BMSB and its egg parasitoid

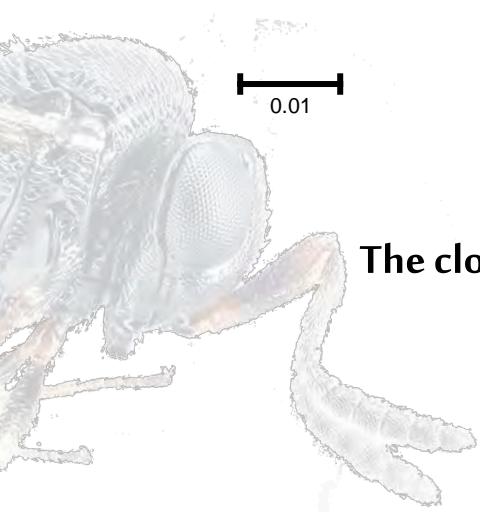
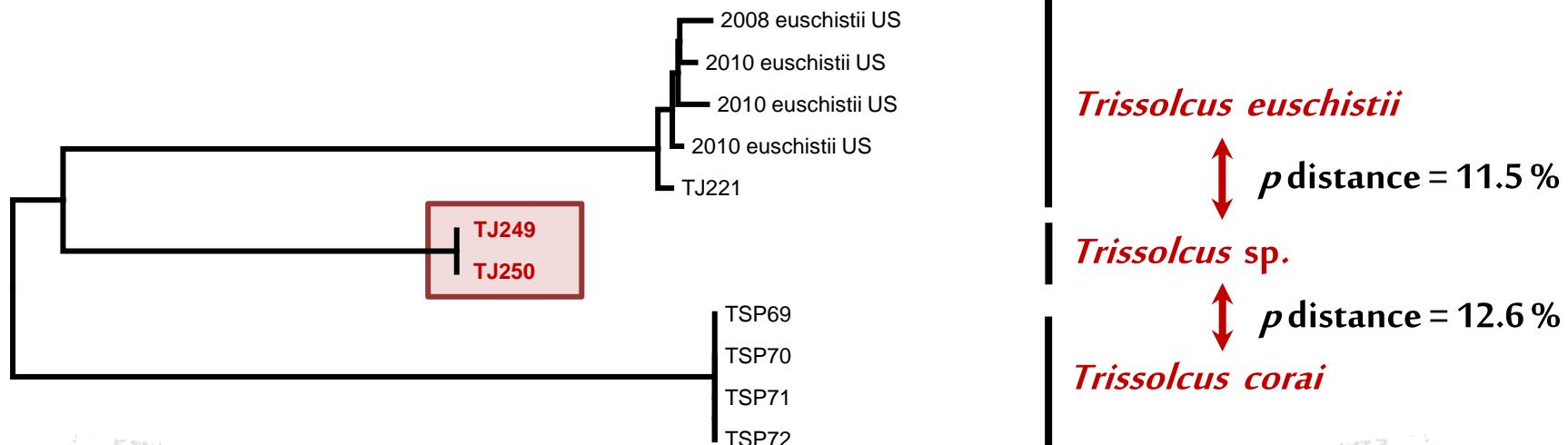
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## COI (barcode) results



The closest taxa

→ *Trissolcus euschistii*  
→ *Trissolcus corai*

Morphology  
*T. euschistii*  
More genes !



# Genetic characterization of BMSB and its egg parasitoid

1.North American BMSB

2.North American *Trissolcus*

3.Phylogeny of *Trissolcus*

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## Bayesian clustering approach

Structure software



# Genetic characterization of BMSB and its egg parasitoid

1.North American BMSB

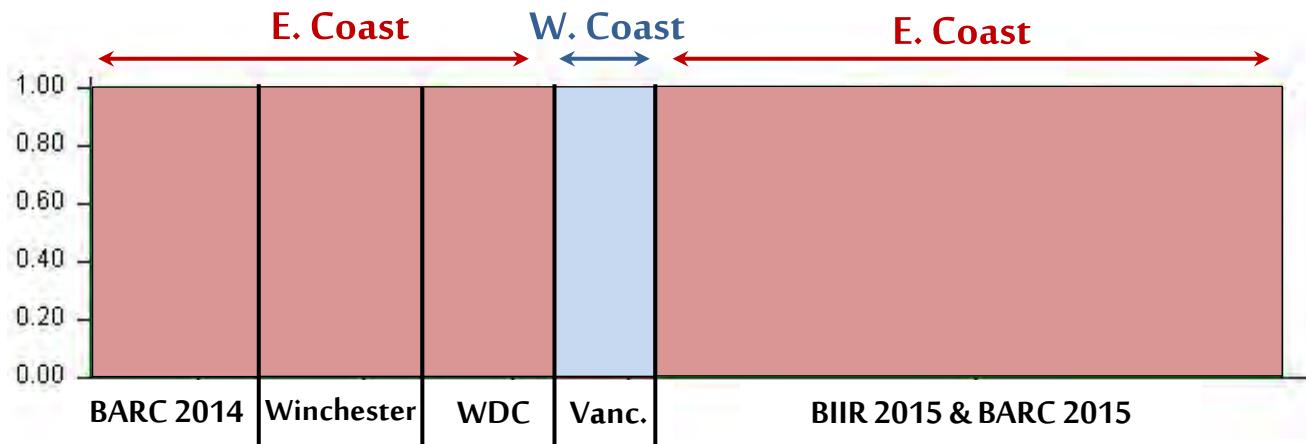
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## Bayesian clustering approach

Structure software



# Genetic characterization of BMSB and its egg parasitoid

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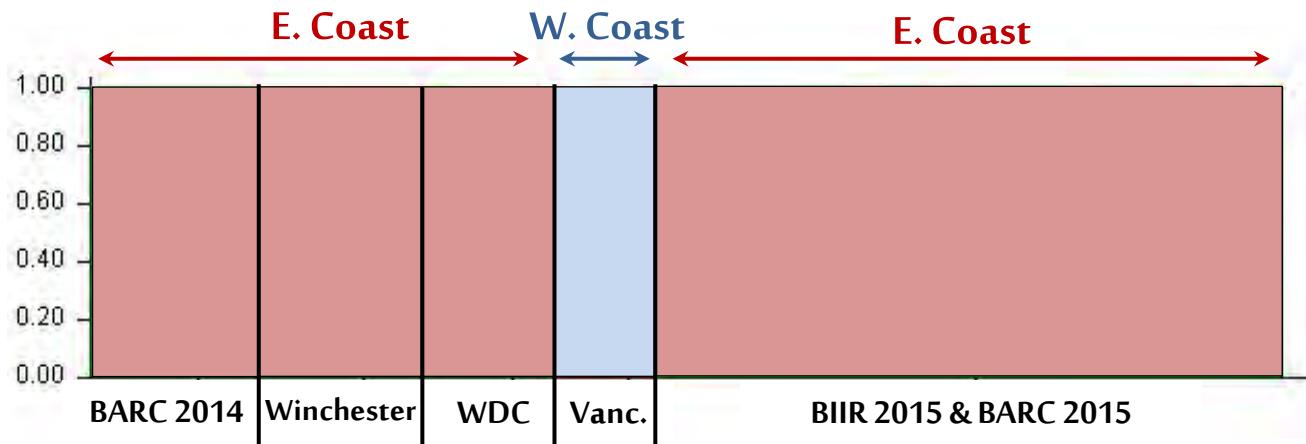
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## Bayesian clustering approach

Structure software

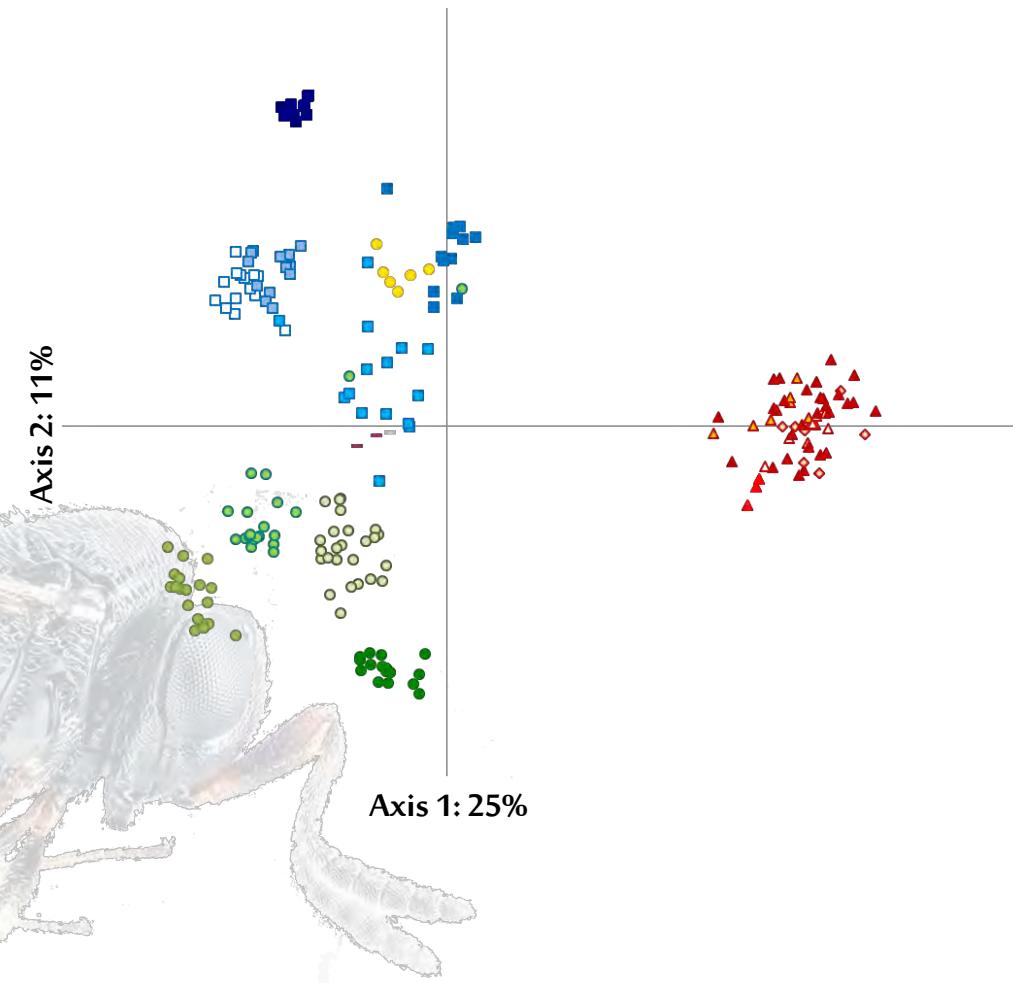


- The most likely value:  $K = 2$
- Different clustering pattern → West Coast population *versus* East Coast populations

Different sources for East Coast and West Coast populations



## Principale Coordinate Analysis (PCoA)

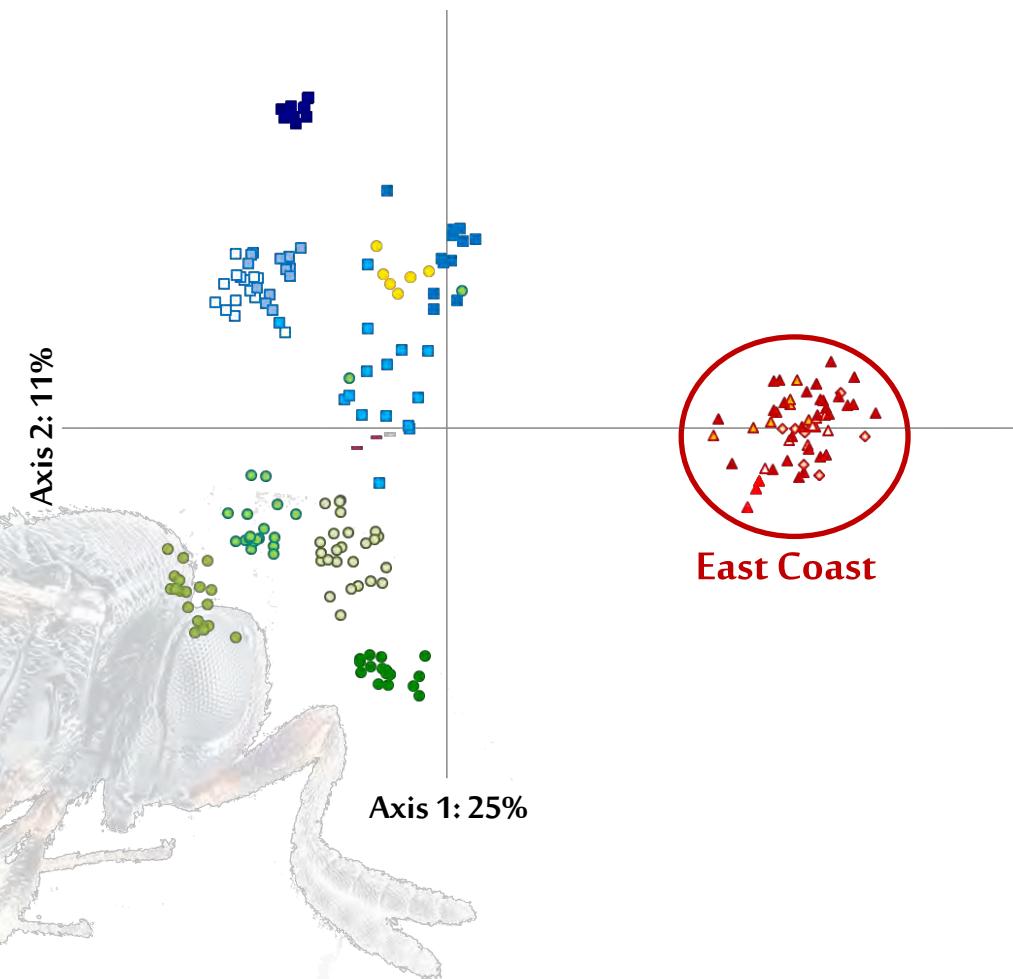


## Principale Coordinate Analysis (PCoA)

- Axis 1: Split between East Coast populations and all other populations



Populations recovered in East Coast do not correspond to an accidental quarantine escape

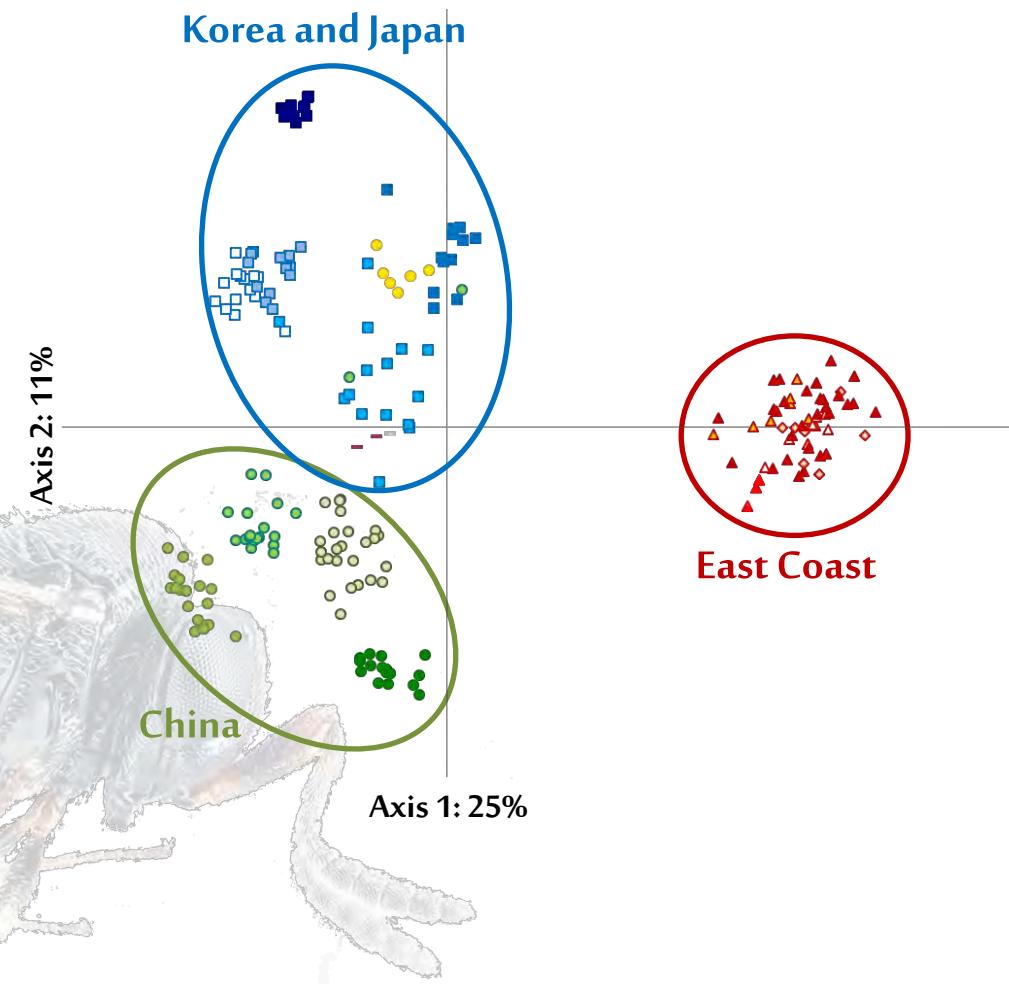


## Principale Coordinate Analysis (PCoA)

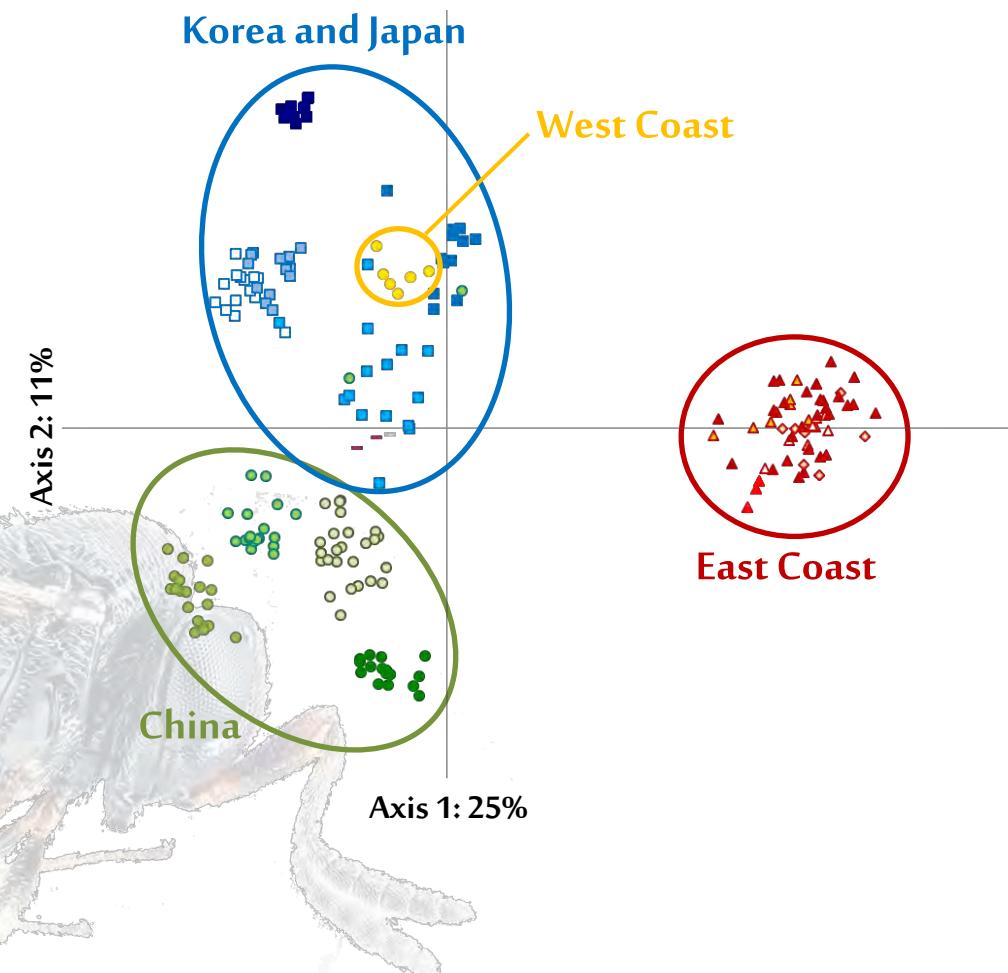
- Axis 1: Split between East Coast populations and all other populations



Populations recovered in East Coast do not correspond to an accidental quarantine escape



## Principale Coordinate Analysis (PCoA)



- Axis 1: Split between East Coast populations and all other populations



Populations recovered in East Coast do not correspond to an accidental quarantine escape

- West Coast individuals

→ Cluster with Korean group



No reason that those individuals correspond to an accidental quarantine escape

Korean origin following an introduction of parasitized eggs or hitchiking adults

## What's next?

- Manuscript in preparation



- A master degree student

→ To type all potential endosymbionts present in the quarantine colonies (using a multilocus approach)

# Update on the genetic characterization of West Coast populations of *Halyomorpha halys* and adventive populations of its egg parasitoid *Trissolcus japonicus*

Phylogeography of *Halyomorpha halys* in the U.S.

Genetic diversity of *Trissolcus japonicus*

Phylogeny of the Asian *Trissolcus*

# Genetic characterization of BMSB and its egg parasitoid

## 1. North American BMSB

## 2. North American Trissolcus

## 3. Phylogeny of *Trissolcus*

14



● Comprehensive Phylogenetic reconstruction based on concatenated analysis (1616bp) of mitochondrial and nuclear genes

● 289 individuals used



# Genetic characterization of BMSB and its egg parasitoid

## 1.North American BMSB

## 2.North American Trissolcus

14

## 3.Phylogeny of *Trissolcus*



● Comprehensive Phylogenetic reconstruction based on concatenated analysis (1616bp) of mitochondrial and nuclear genes

● 289 individuals used



Congruence between molecular delineation and morphological characters for most lineages

Evidence of 19 *Trissolcus* species  
Some are new to science

Real tool for diagnostic

# Genetic characterization of BMSB and its egg parasitoid

1.North American BMSB

2.North American *Trissolcus*

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## What's next?

- 2 more genes are sequenced to get a robust phylogeny
- Final results for datation are expected before ICE2016
- Preliminary results and the philosophy of the project



→ Poster and oral presentation of the project

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*Thank you for your attention....*



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*Thank you for your attention....*



# Genetic characterization of BMSB and its egg parasitoid

1.North American BMSB

2.North American *Trissolcus*

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## COI (barcode) results

