

Identification of the Origins of *Coniatus* sp., a Non-Native Weevil in Western North America



Outline

- Intro
 - History in US
 - Identification
 - Range
- Questions
 - One/multiple introductions
 - Origin
 - Molecular phylogenetics
 - Lifecycle
 - Polymorphisms
 - Color
 - Pupal case
 - Impact on tamarisk
 - Shoots
 - Flowers



First sightings in US in 2006 in Arizona

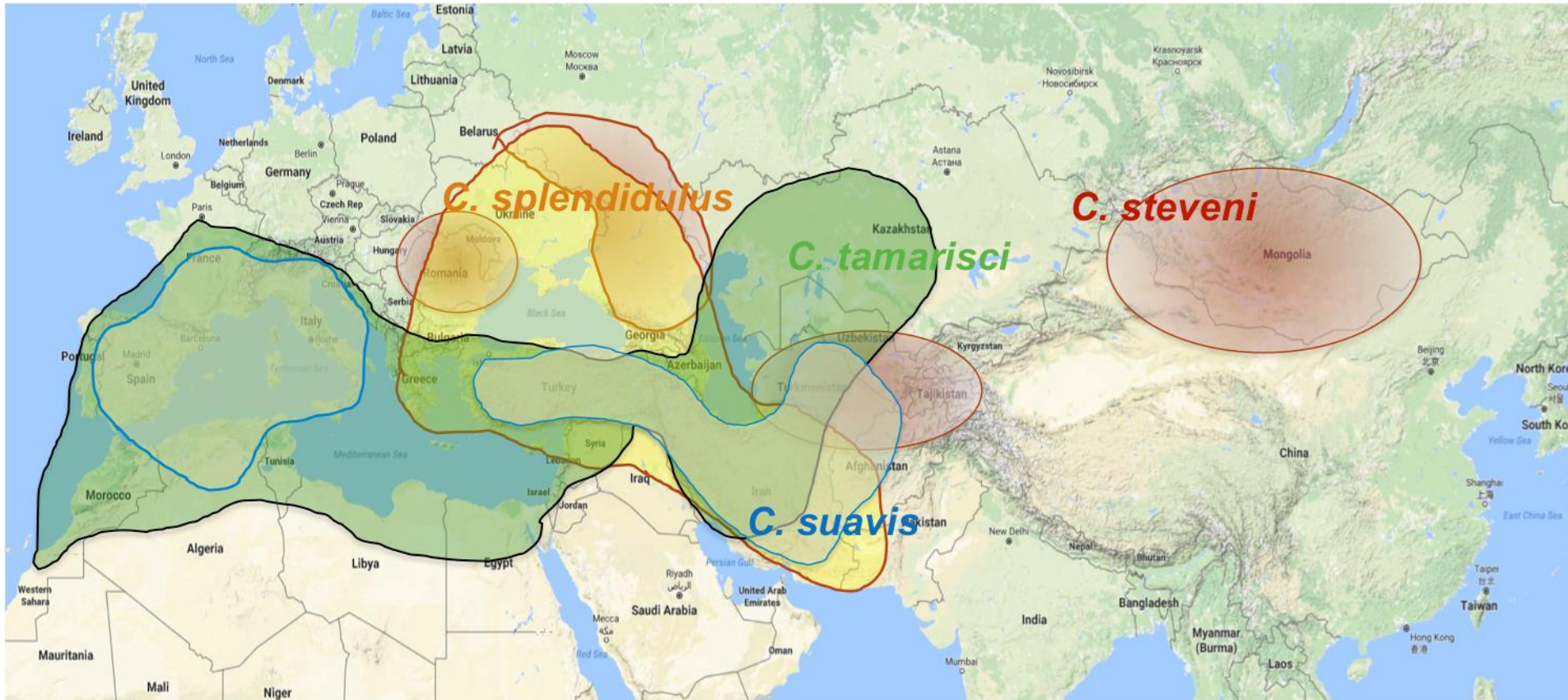
- Non-native to North America
- First published report in 2011
 - Jason R. Eckberg and Marissa E. Foster. 2011. First account of the splendid tamarisk weevil, *Coniatus splendidulus* Fabricius, 1781 (Coleoptera: Curculionidae) in Nevada. *The Pan-Pacific Entomologist* 87(1): 51-53.

Range of *Coniatus* sp. in North America



- 24 species in genus *Coniatus*
- Believed to be natural predators of *Tamarix* species
- Other *Coniatus* species have been studied and evaluated as possible biocontrol agents for tamarisk
 - *C. repandus*,
 - *C. tamarisci*
 - *C. stevini*
- *C. tamarisci* was approved by the Technical Advisory Group for Biological Control Agents of Weeds
 - no known intentional releases of any of *Coniatus* species have been made in US

Native Ranges of four *Coniatus* spp.



Based on Catalogue of Palaearctic Coleoptera

Questions

- One/multiple introductions?
- Origin?
 - Molecular phylogenetics

CO1 analysis

U.S. Collections –



Tom Dudley,
Karen Rosen,
Dan Bean,
Levi Jamison

Overseas collections



Massimo Cristofaro, Francesca Marini, René Sforza, Amanda Stahlke, Laibale Friedman

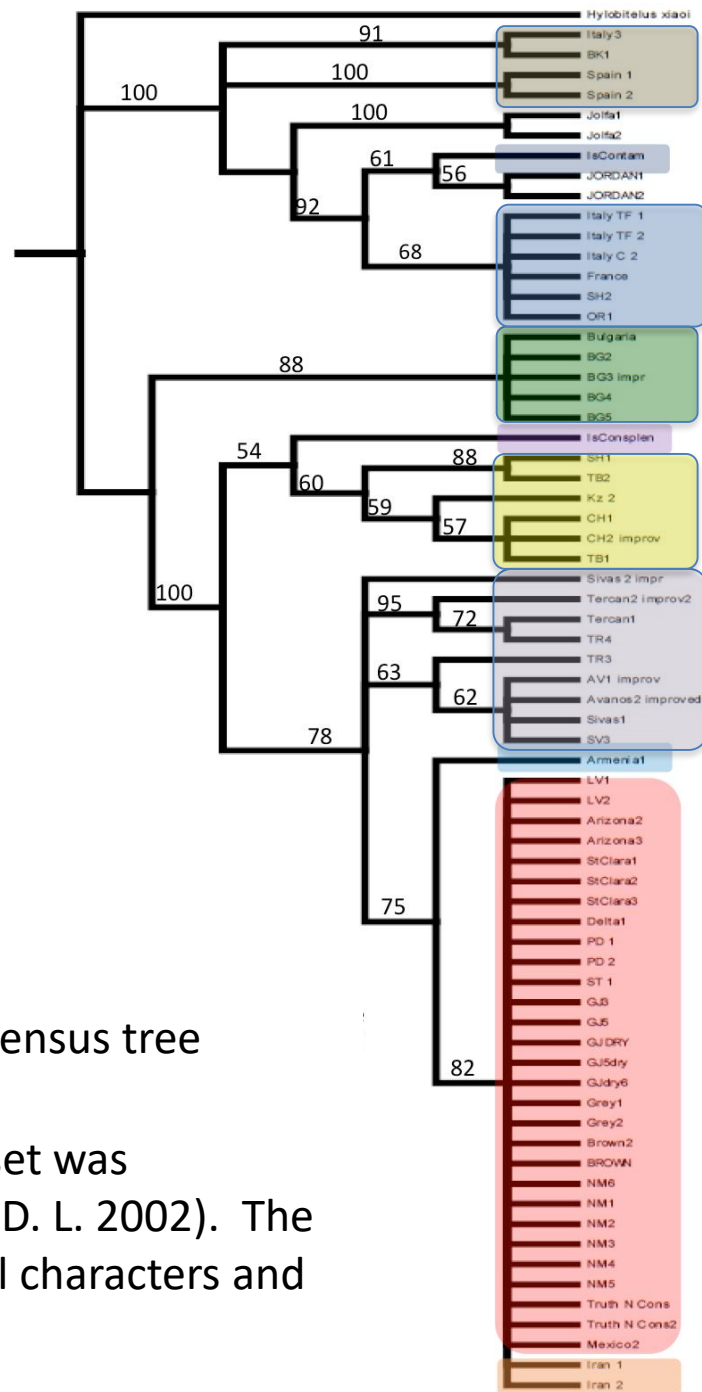
North American samples display little to none variation in CO1 (1296 bp)



Single introduction

Bootstrap 50% majority-rule Consensus tree

Maximum Parsimony analysis of the data set was performed using PAUP*v.4.b10 (Swofford, D. L. 2002). The bootstrap number was 1000 with 755 total characters and 217 informative characters.



N. America

[illegible]

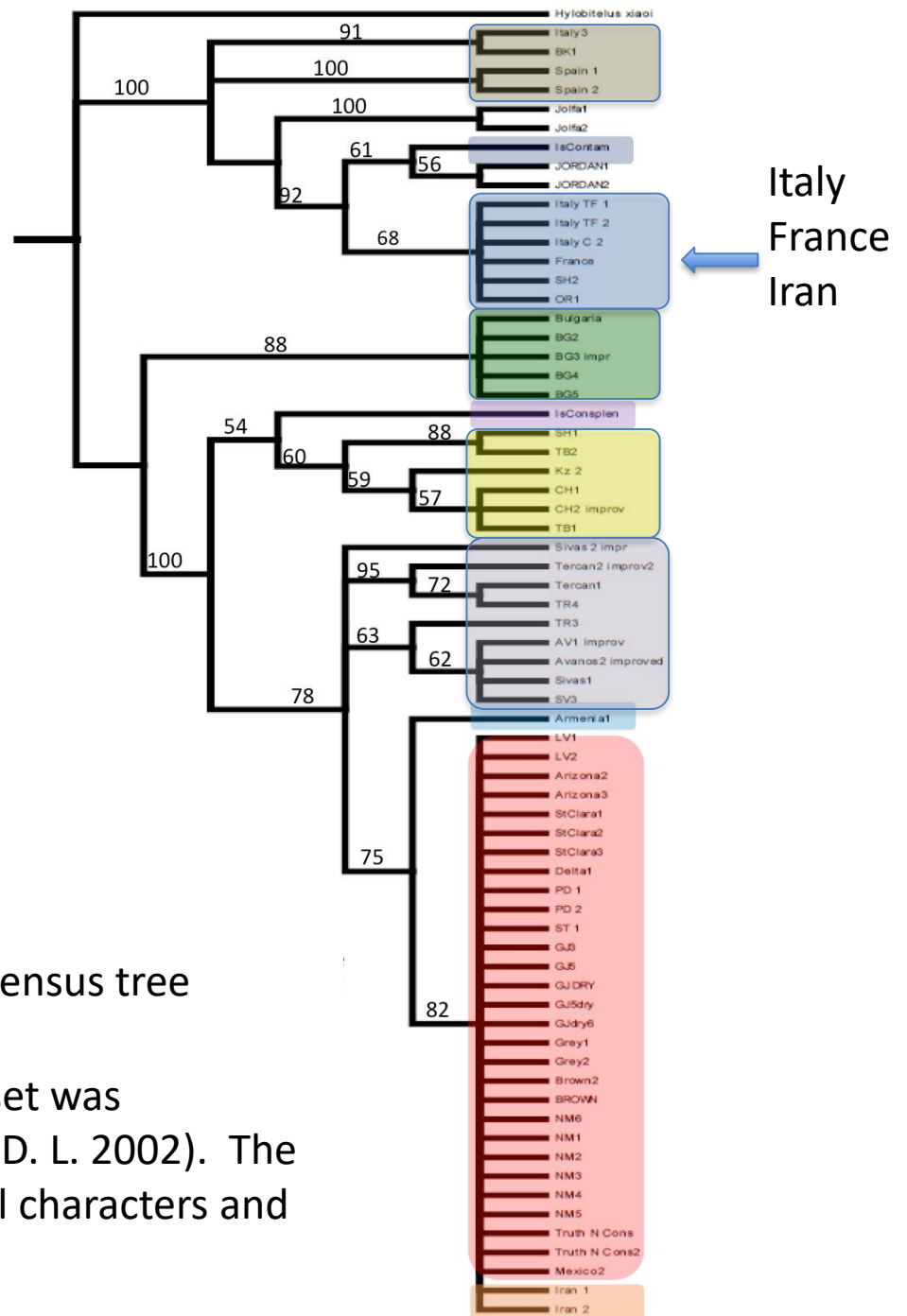
Some clades contain specimens collected from different sites



Some species present in several geographical areas

Bootstrap 50% majority-rule Consensus tree

Maximum Parsimony analysis of the data set was performed using PAUP*v.4.b10 (Swofford, D. L. 2002). The bootstrap number was 1000 with 755 total characters and 217 informative characters.



North American samples do NOT group with *Coniatus splendidulus*

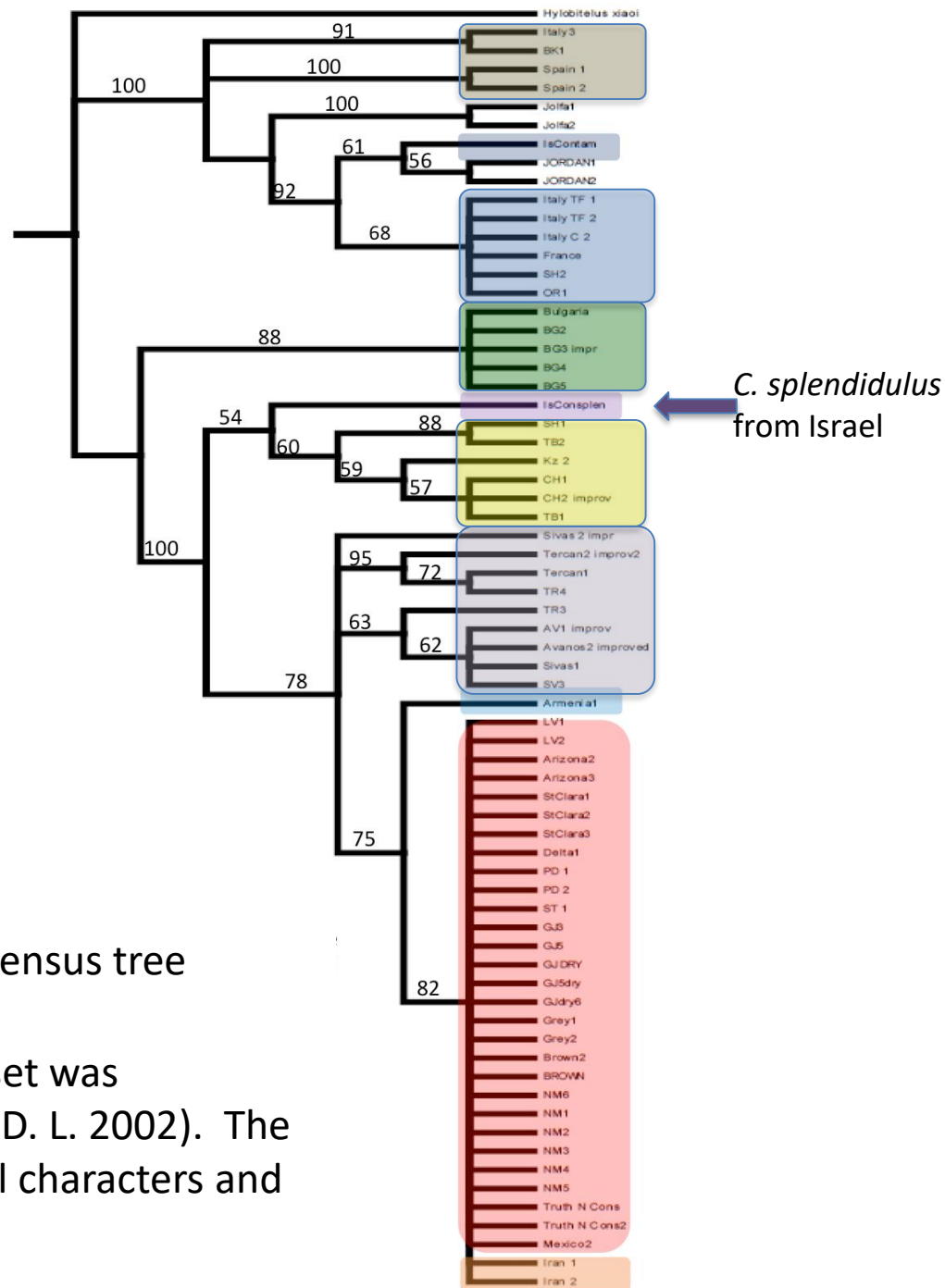


Species designations may need to be redone in the genus using DNA markers

Use of museum specimens which have already been taxonomically identified based on morphology

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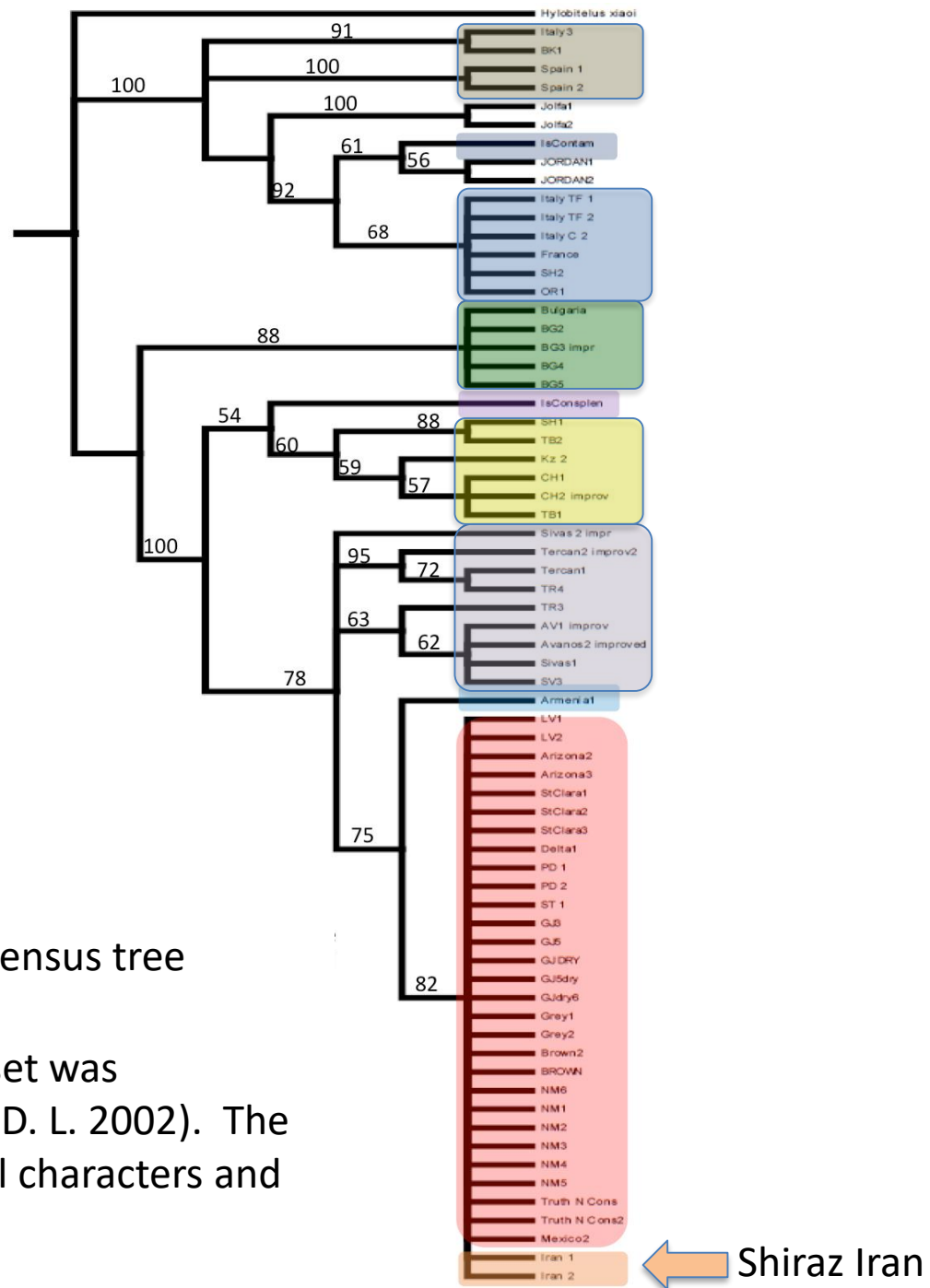
North American samples DO group with specimens collected near Shiraz, Iran



We may have an origin

Bootstrap 50% majority-rule Consensus tree

Maximum Parsimony analysis of the data set was performed using PAUP*v.4.b10 (Swofford, D. L. 2002). The bootstrap number was 1000 with 755 total characters and 217 informative characters.



Best Match: Shiraz, Iran

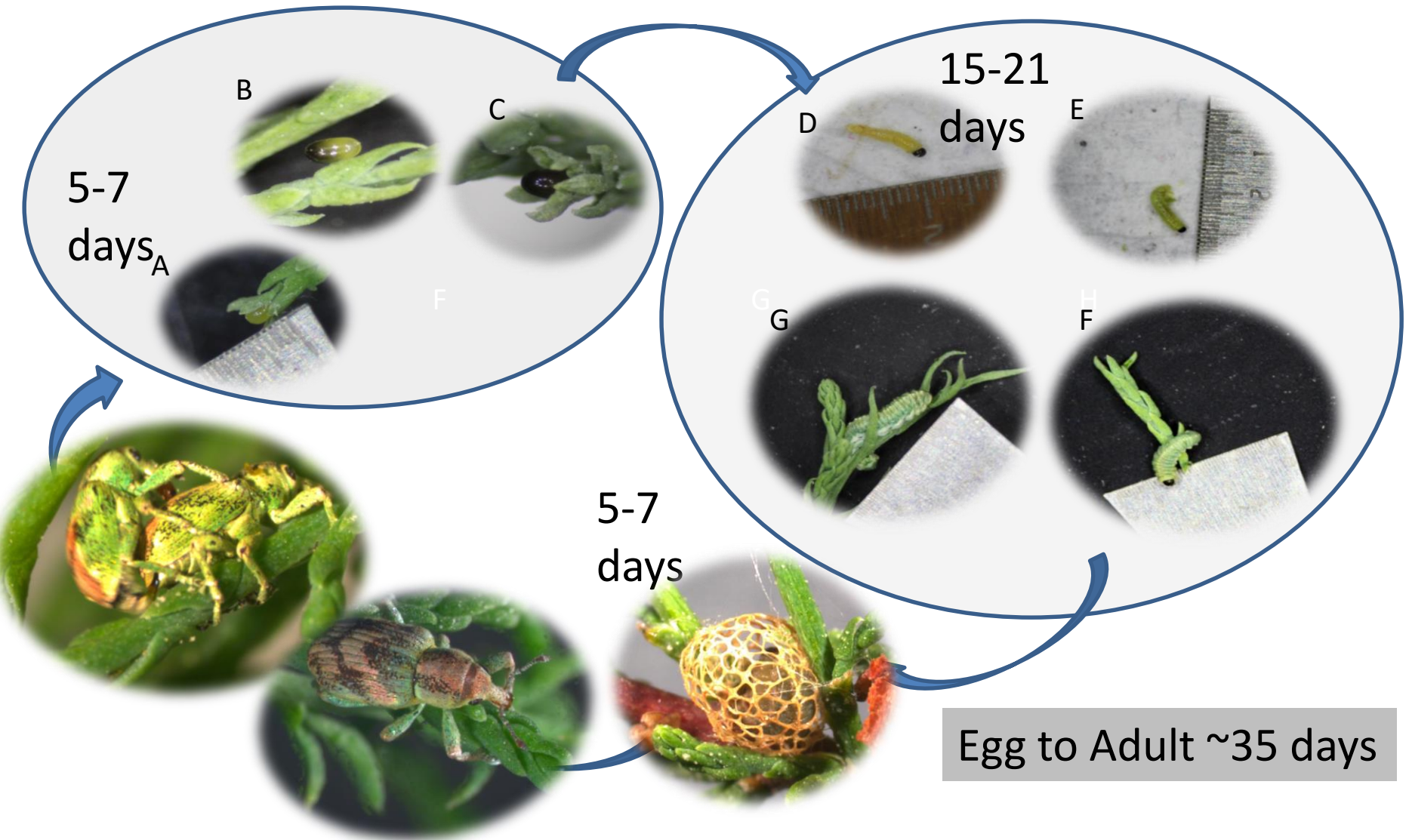


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Life cycle of the *Coniatus* sp. collected in GJ , CO

Insects were raised in an incubator at 25C at 14L:10D light cycle (similar to the day length observed in GJ late May/early June)



Color morphs



Color polymorphisms throughout the season

Pupal case polymorphisms



Questions- and some answers

– One/multiple introductions

– Origin

• Molecular phy

- 3-4 generations/season
- Long lifespan >12 months
- Found on tamarisk March-late November


– Lifecycle 

– Polymorphisms

• Color 

Seems to be dependent on the time of the season

• Pupal case

- 
- not due to sex of the larvae
 - May be an inherited trait

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Coniatus adults feed on the tip of the shoots



Oviposition at the shoot tips



Oviposition at the shoot tips
AND flower buds – lower seed production?



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Coniatus has the potential to stunt the plant growth and lower seed production

CMU Undergraduates



Austin Hadley



Rachel McLaughlin



Amanda Stahlke



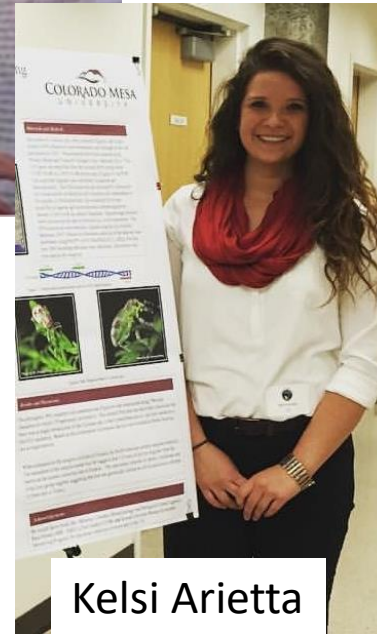
Ashly Pennington



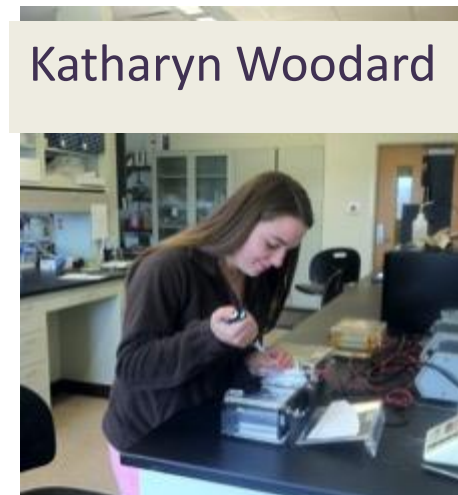
Greg Johnson



Madison Vincent



Kelsi Arietta



Katharyn Woodard

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