

# SEARCHING FOR STREPSIPTERA: Museomics in the study of host-parasite relationships

RJ Millena



AMNH\_LZC\_004075



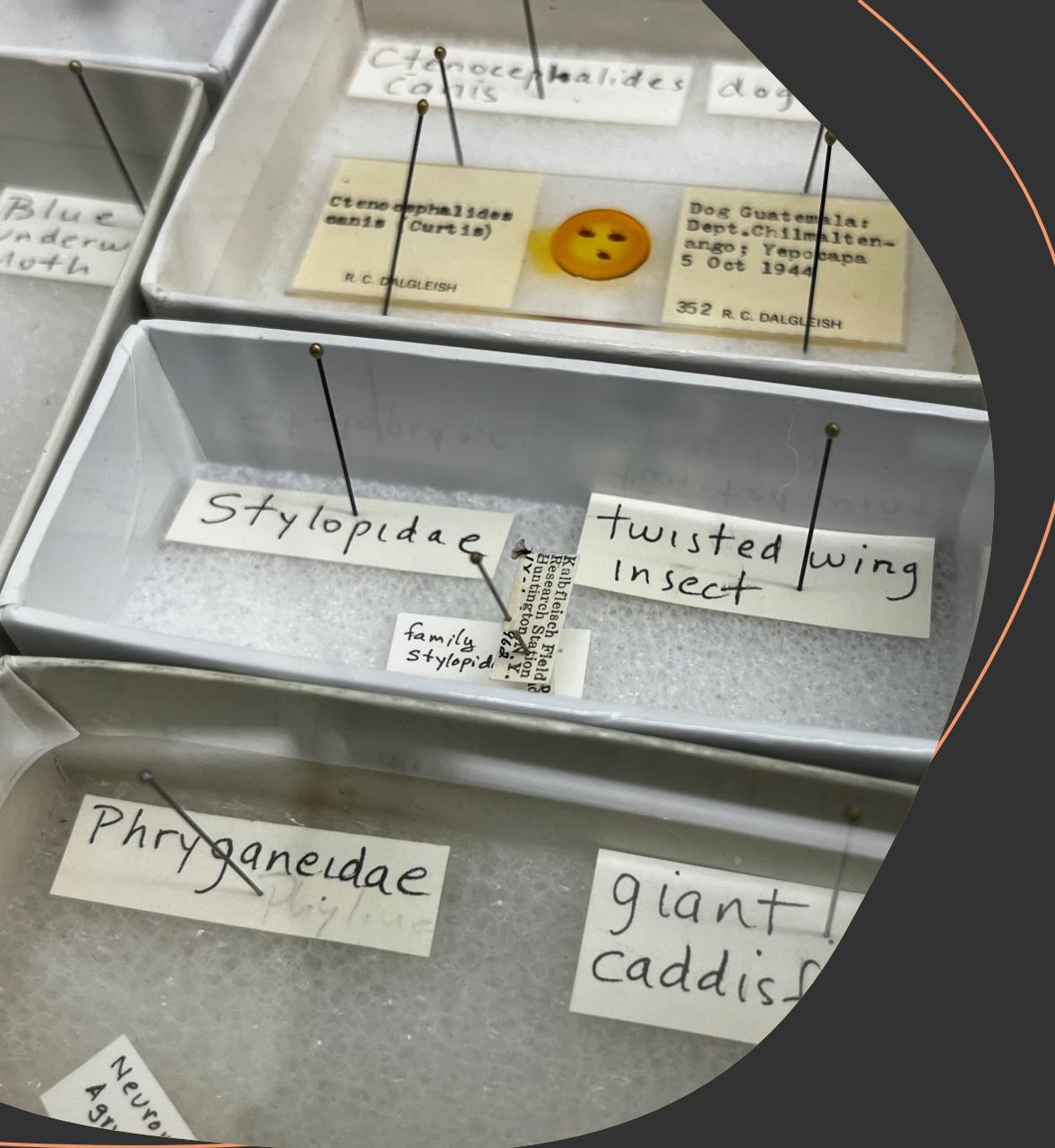
# Outline

Strepsiptera & Challenges

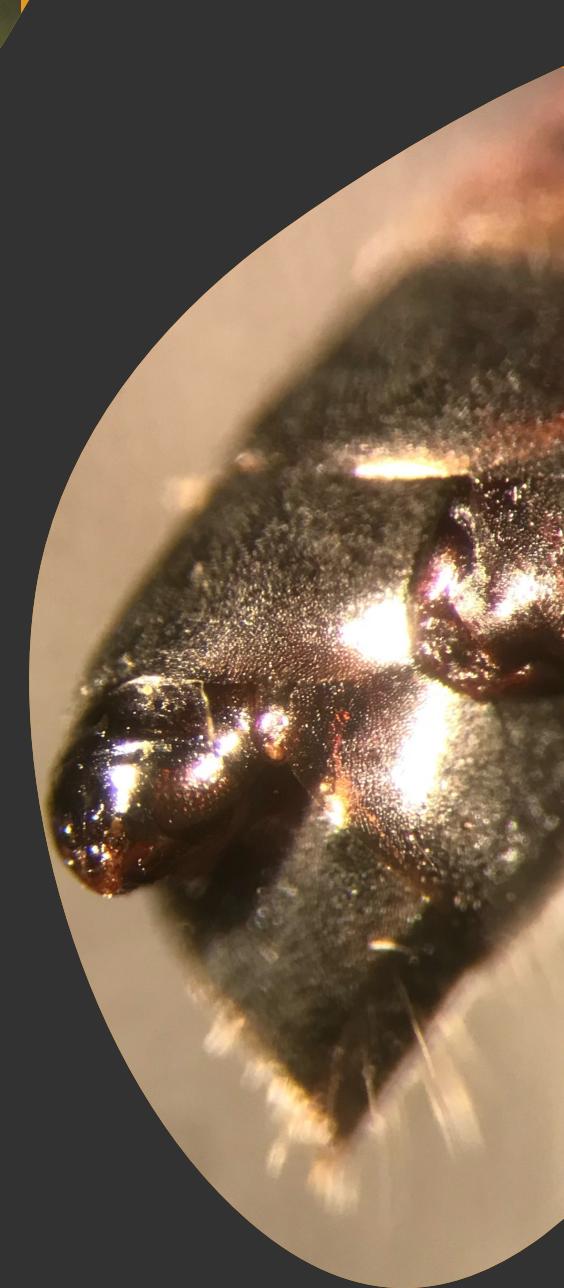
Ecological and Biological Questions

Museomics & Evolutionary Questions

Planned Work



# What is Strepsiptera?



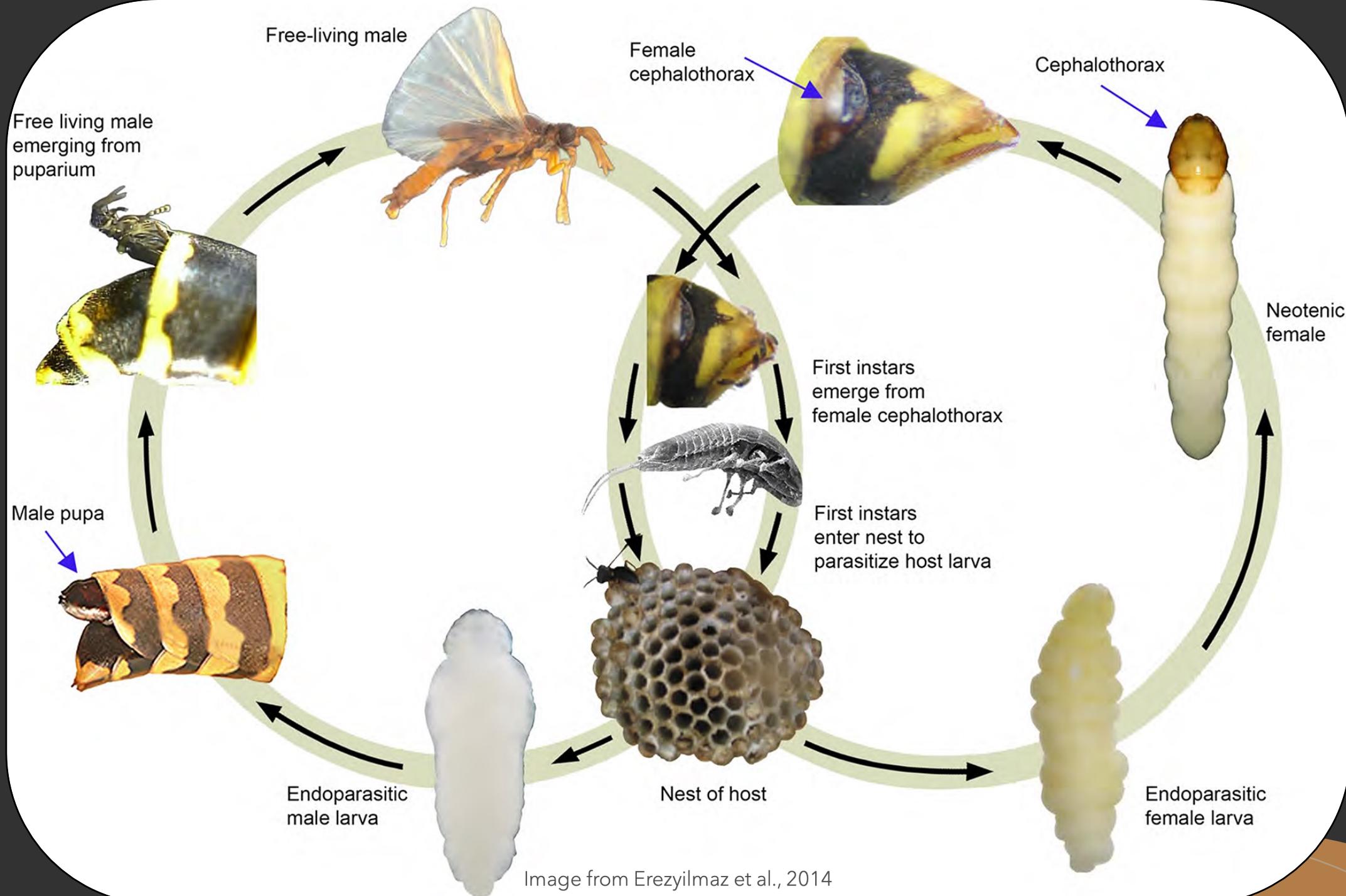
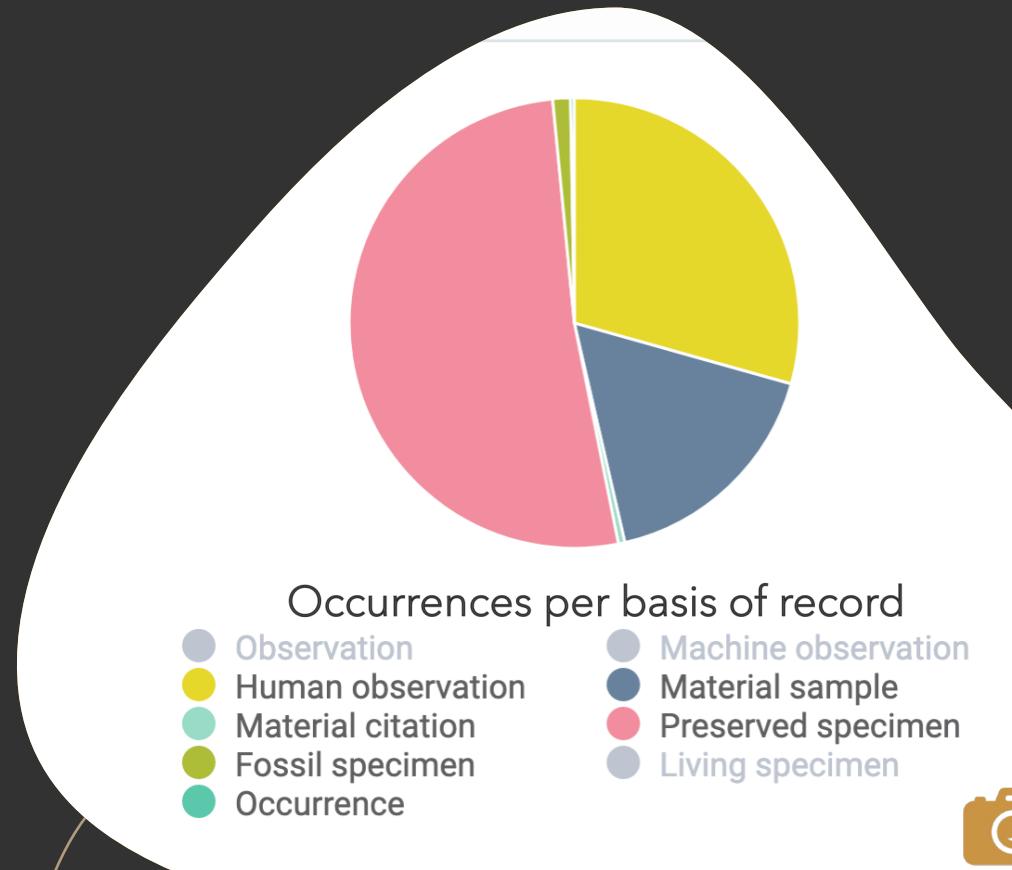


Image from Ereyilmaz et al., 2014



# Challenges in Study

- Extreme specialization
- Perceived rarity afield
- Distinctive genetic characteristics
  - High ribosomal DNA substitution rates
- Taxonomic confusion



# Benefits of Museum Collections

- Accessibility
- Wealth of data
- Temporal and spatial reach
- Education



# Ecological and Biological Questions

BIOLOGY  
LETTERS

royalsocietypublishing.org/journal/rsbl

Research



Check for updates

Cite this article: Millena RJA, Rosenheim JA. 2022 A double-edged sword: parental care increases risk of offspring infection by a maternally vectored virus.



BOHART  
Museum of Entomology  
University of California, Davis



UC DAVIS  
UNIVERSITY OF CALIFORNIA

Evolutionary biology

A double-edged sword: parental care increases risk of offspring infection by a maternally vectored virus

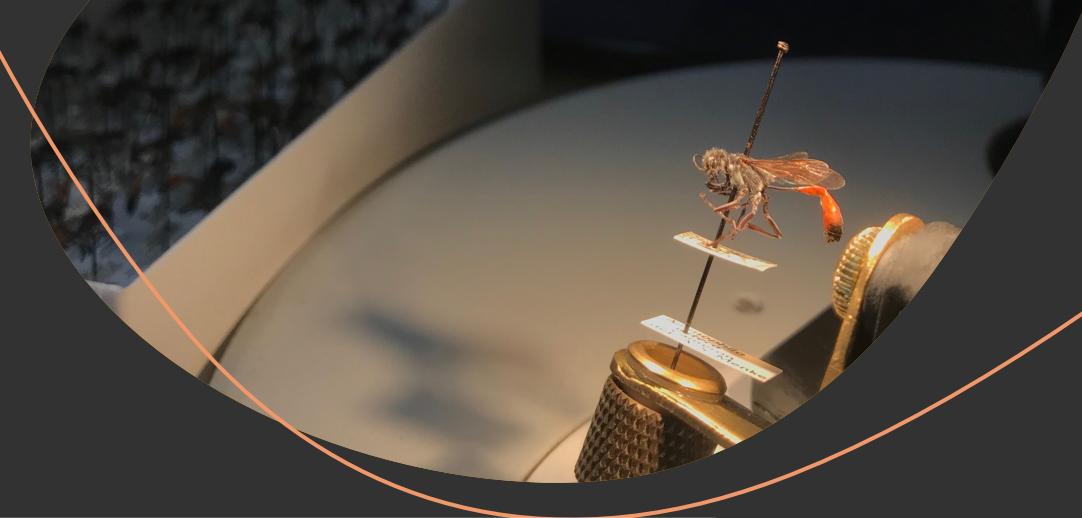
Rebecca Jean A. Millena<sup>1,2</sup> and Ja-

<sup>1</sup>RGGS, Invertebrate Zoology, American Museu

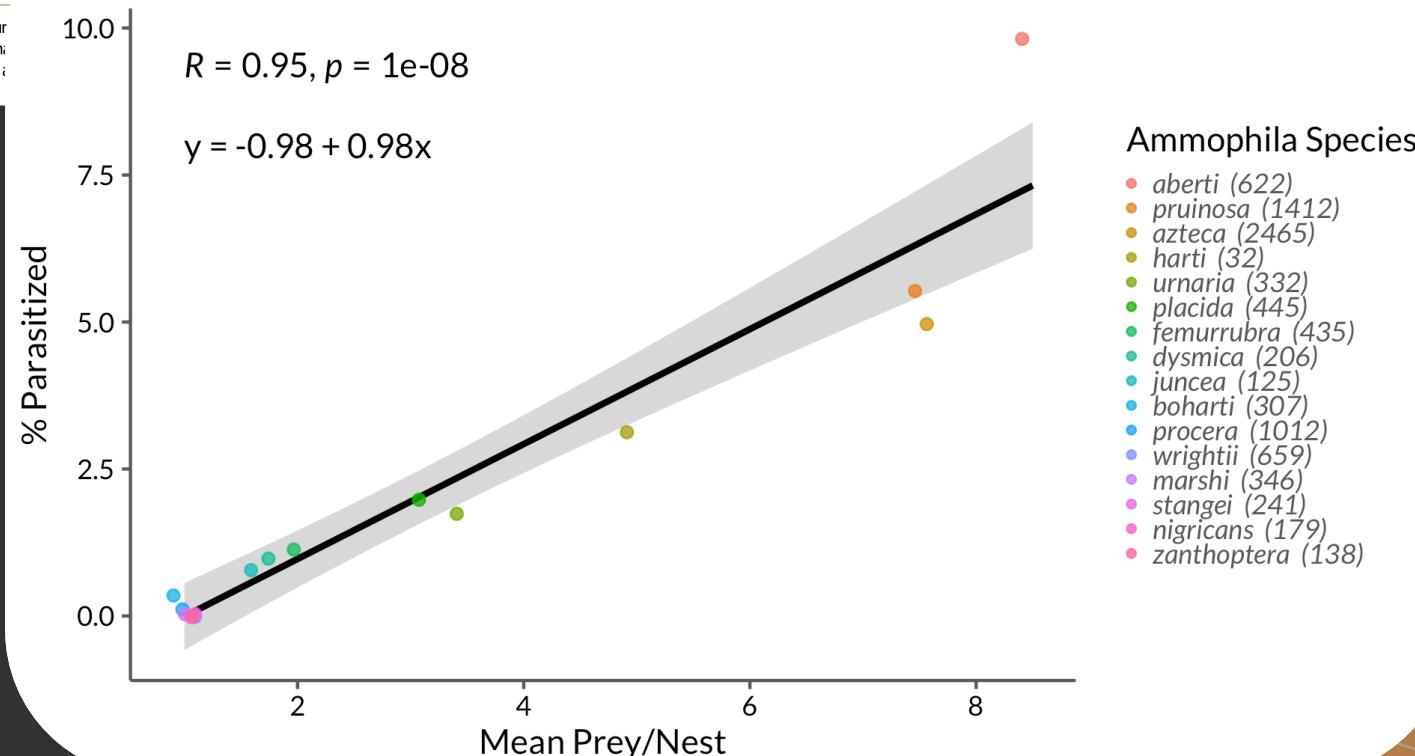
<sup>2</sup>Ecology and Evolution, Entomology and Nemi

<sup>3</sup>Department of Entomology and Nematology, i

Davis, CA, USA



Effect of Provisioning on Parasitism Rate Per Species

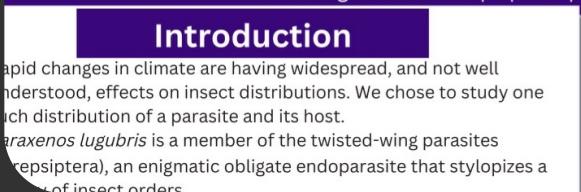


# Ecological and Biological Questions

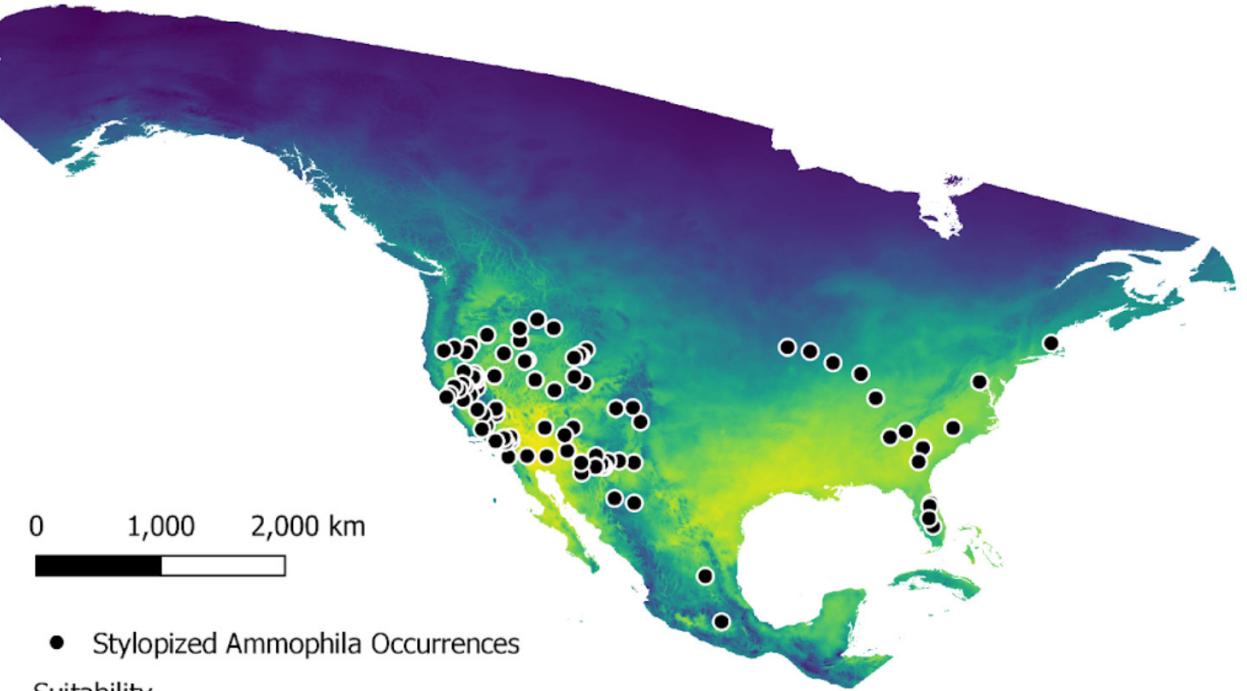
## Climate and collections: Modeling Strepsipteran host-parasite dynamics with museum collections

Wendy G. Frankel, Trinity Tobin, RJ Millena, Aaron Goodman

Title changed from: Strepsiptera parasitism rates across taxa in American museum collections



Map of Strepsiptera Ammophila Occurrences Against Ammophila Suitability



# Museomics

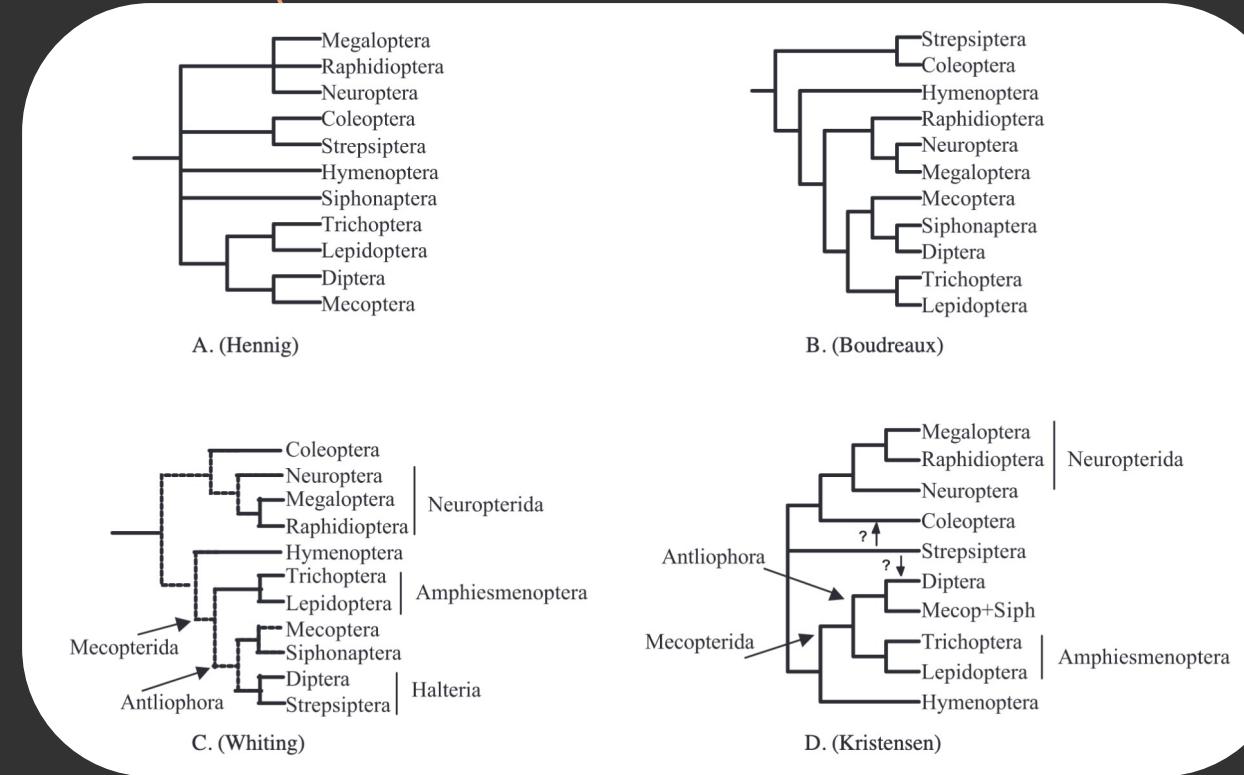
- The study of DNA sequences obtained from museum specimens
  - Archival storage vs natural degradation
- From short fragments of hDNA to whole genome
- Improving taxonomic coverage

hDNA target group	Specimen type	Sources for hDNA
Infectious diseases	Mammal dry skin Amphibian formalin/alcohol preserved skin	Skin or epithelial swabs
Microbial communities	Herbarium dry pressed	Leaf tissue
Plants	Herbarium dry pressed	Leaf tissue
Vertebrates	Anthropological material: Headdresses, clothing	Biological material
Mammals	Skins and hides	Hair, skin, muscle traces, claws
Mammals	Skeletal	Bone, teeth
Birds	Round and flat dry skins	Toe pads, skin
Birds; crabs	Eggs	Piece of egg or powder
Fishes Amphibians Reptiles	Fluid: formalin and/or ethanol preserved	Tissue including liver, muscle
Reptiles	Bone	Carapace, plastron, other bone elements
Insects Arachnids	Pinned, dried specimens	Whole limbs, whole specimens, subsamples
Bryozoa	Dried colonies	Partial specimens
Arachnids	Fluid: ethanol preserved	Whole or partial specimens



# Evolutionary Questions

- Enigmatic placement in the insect radiation
- Unresolved intraordinal relationships
- Ancestral characteristics
- Evolutionary trends

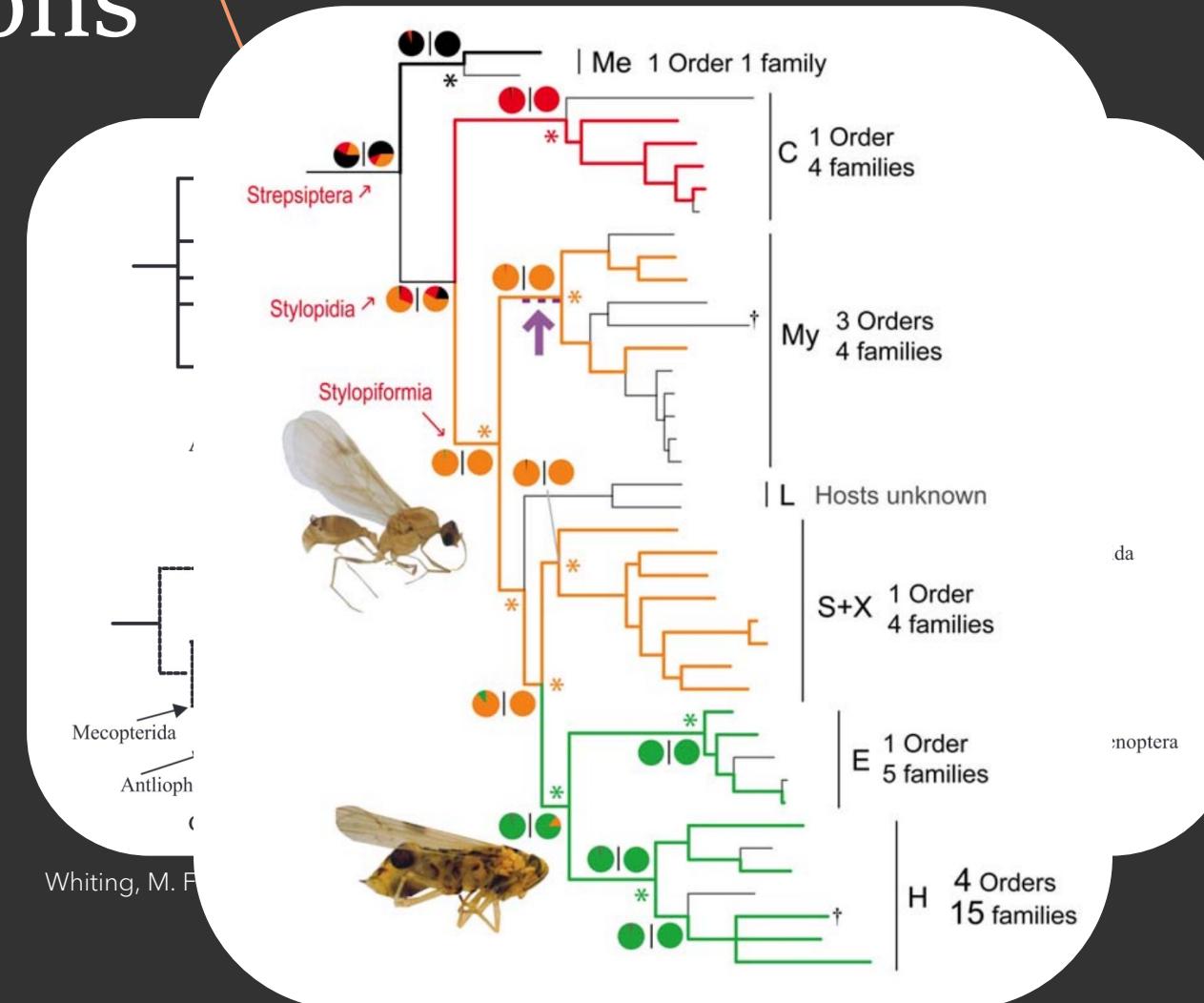


Whiting, M. F. (2002). *Zoologica Scripta*, 31, 3 - 15



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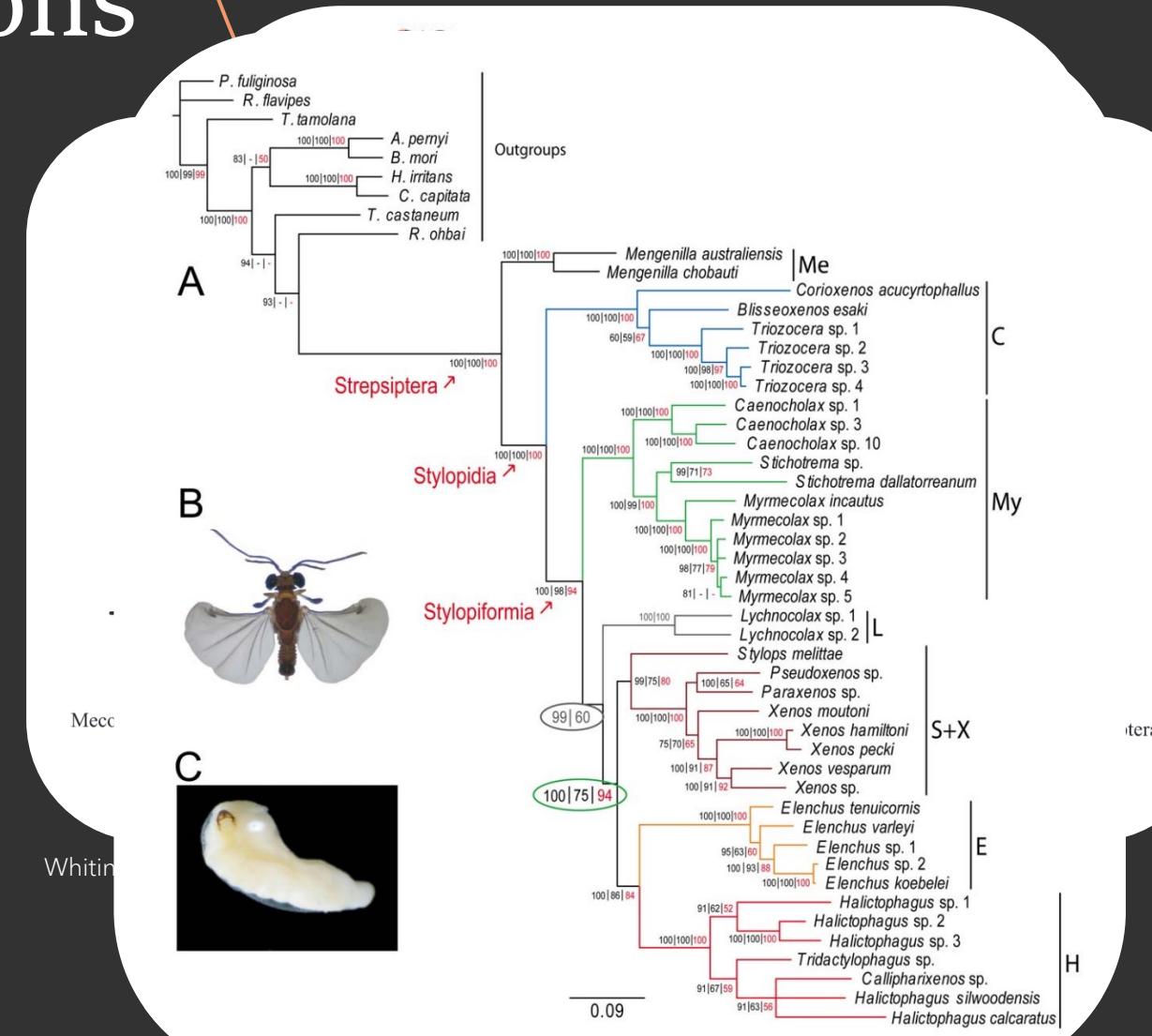


McMahon et al. (2011). PLoS ONE 6(6), e21206.



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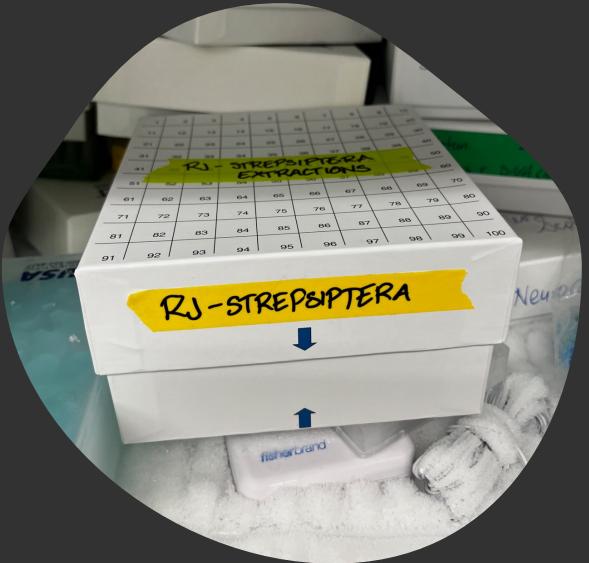
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# Goals and Methods



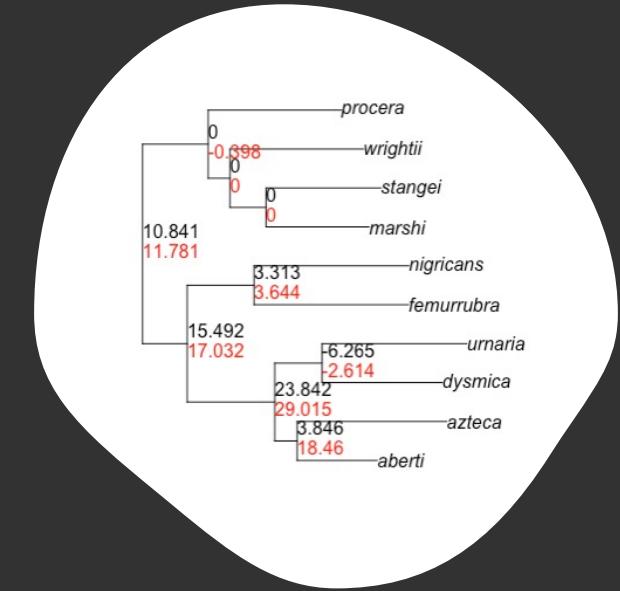
Find and dissect  
Strepsiptera from  
preserved hosts



Nondestructive  
protocol development  
& testing; extractions



Anchored hybrid  
enrichment (AHE)  
bait design

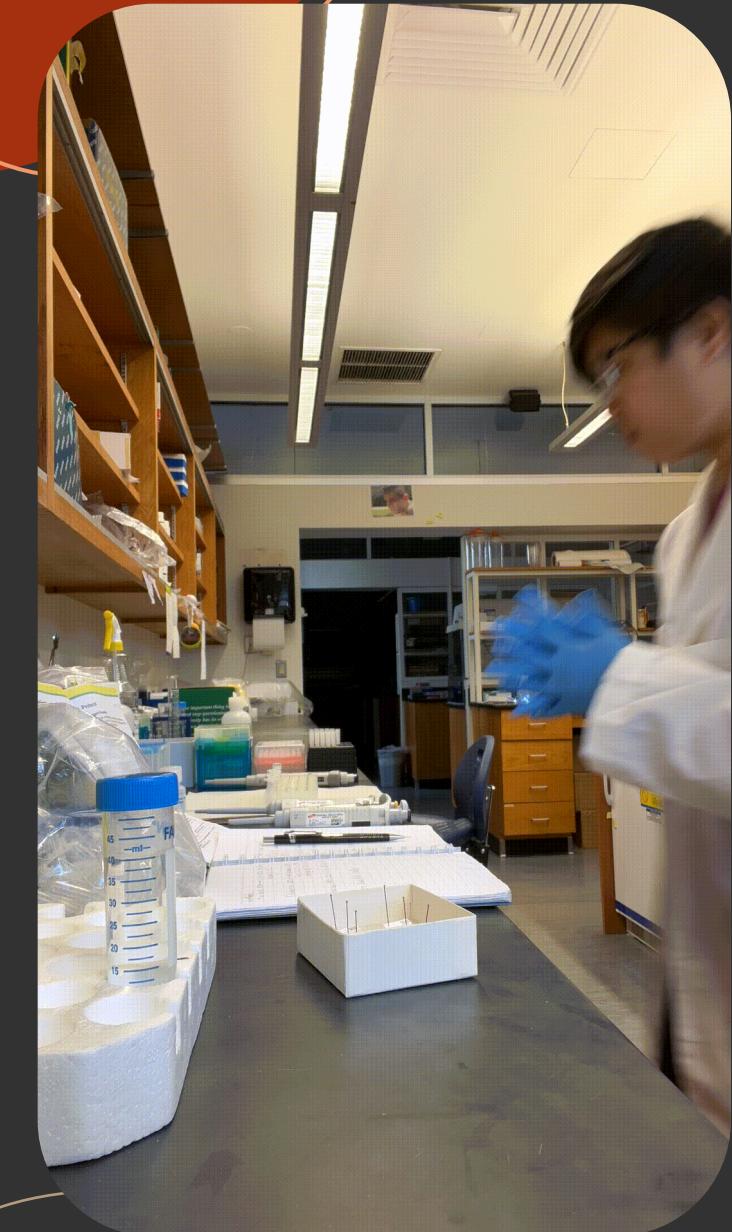
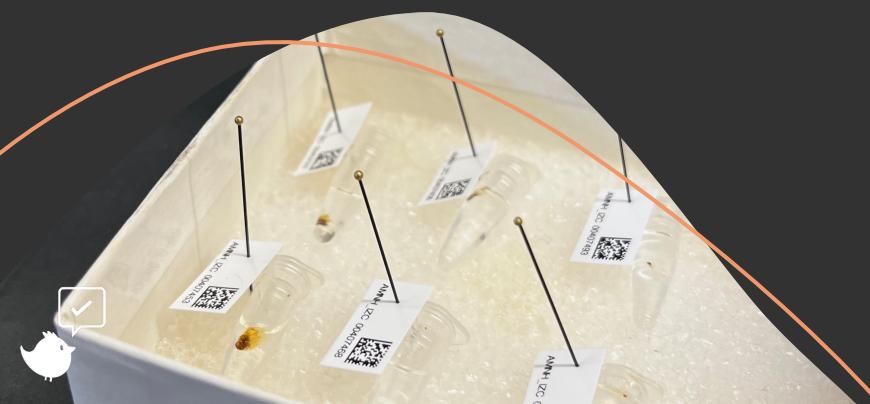


Phylogenetic  
analyses



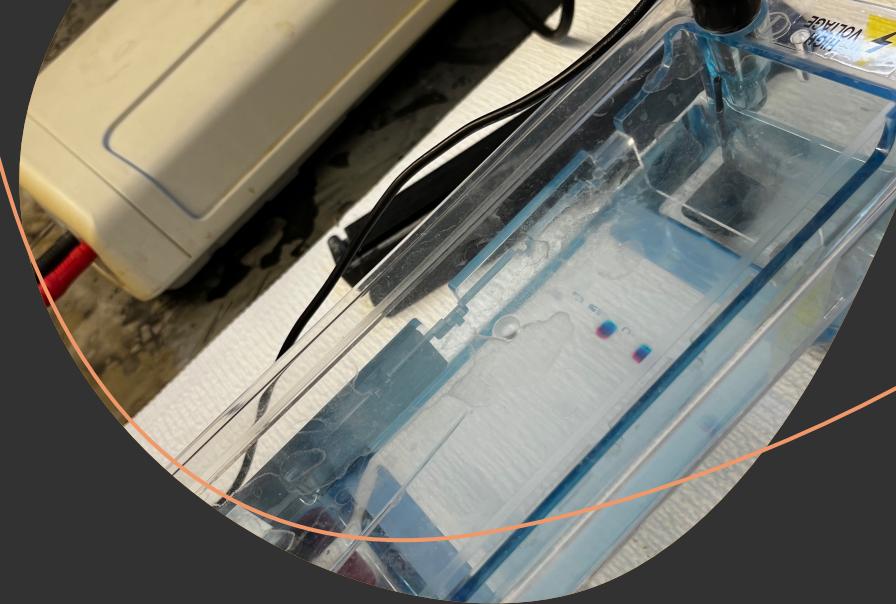
# Nondestructive Protocol

- Zymo Quick-DNA Miniprep Kit
- Whole body in proK and lysis buffer, 48 hr incubation
- Rinsed with ethanol, suspended in glycerin or slide mounted



# Testing the Protocol

- 6 specimens of *Paraxenos lugubris* from unidentified *Ammophila*
  - 2 USA, 4 ARG, 1974-2012
  - Malaise, dry
- Three tests
  - Preliminary: Nanodrop
  - PCR presence/absence test, Sanger
  - Oxford Nanopore MinION test



## Nanodrop Values

Sample #	Year Collected	ng/ $\mu$ L
1	1974	2.2
2	1974	2.2
3	2012	49.5
4	2012	1.6
5	1987	3.3
6	1992	2.7



HH HS 1H 1S 2H 2S 3H 3S

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4H 4S 5H 5S 6H 6S

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HH HS 1H 1S 2H 2S 3H 3S

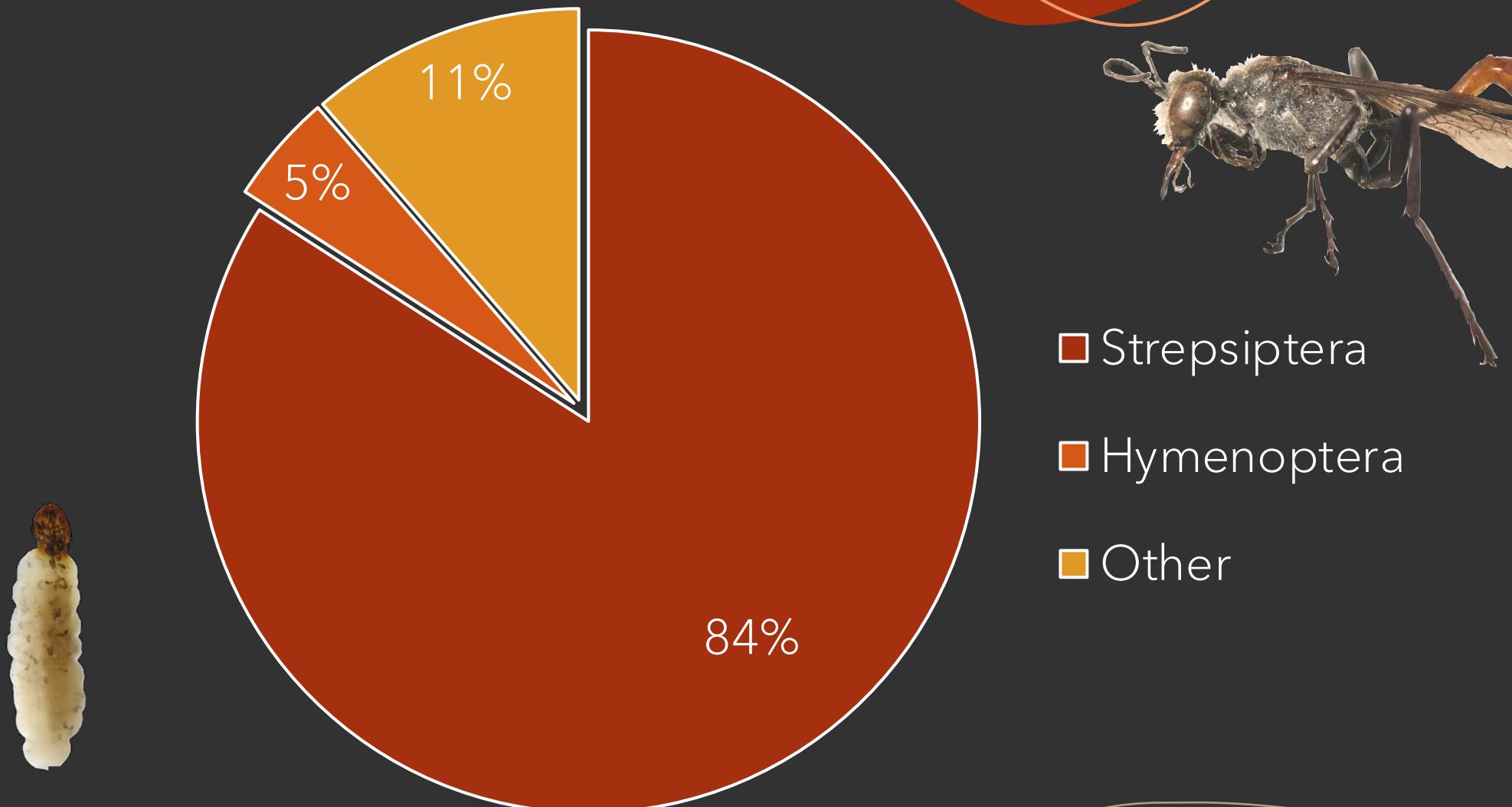
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4H 4S 5H 5S 6H 6S

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# First BLAST Result for Nanopore Reads



# Planned Work

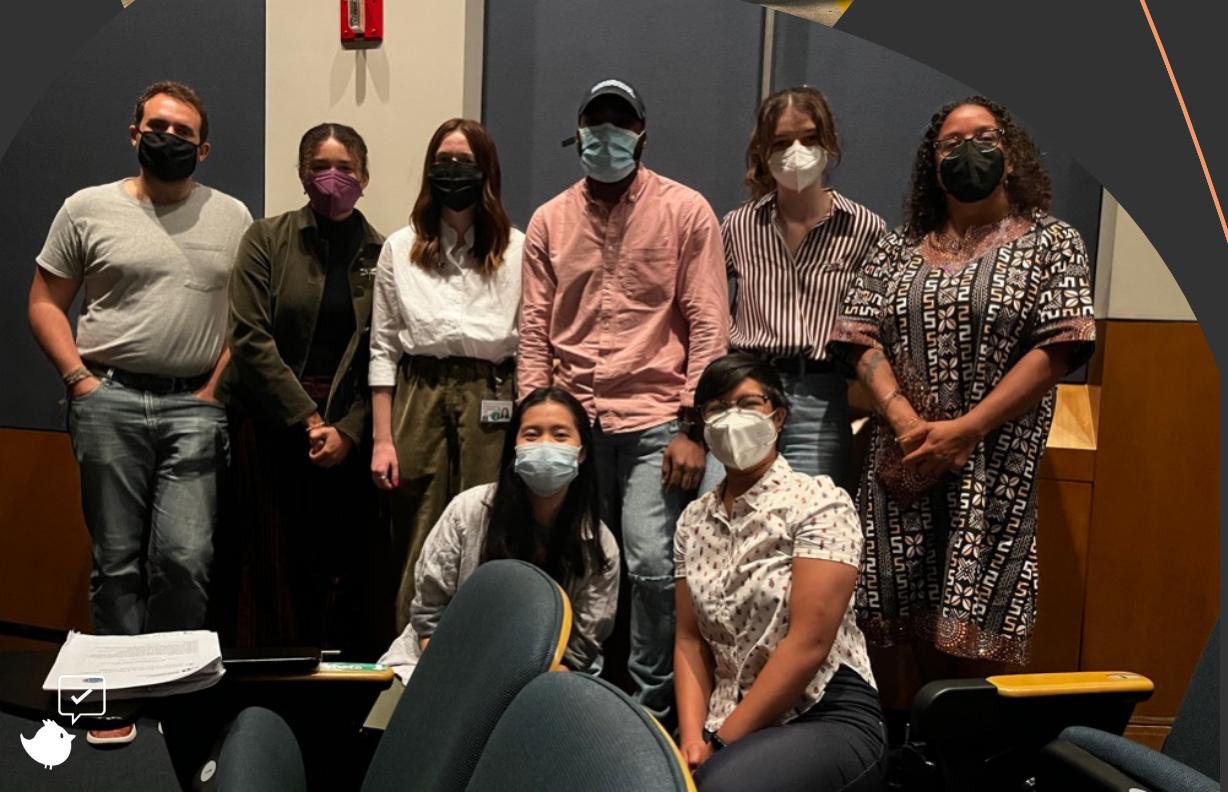
- Nanopore sequence of *Xenos peckii* whole genome, assembly & annotation
  - Reference genome
- Probe design with BaitFisher, AHE analyses, total evidence
- Host-parasite associations
  - Via collections and in phylogenetic context
- Species richness of Strepsiptera
- Optimizing museomics protocols



# Thank You!



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OF NATURAL HISTORY



**UCDAVIS**  
UNIVERSITY OF CALIFORNIA

Ware Lab  
Ruth Salas  
Christine Labeau  
Melody Doering



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