

# Bridging gaps in the study of parasitoid wasps: a perspective from the global south.

Eduardo Mitio Shimbori - 29 février 2024 - CBGP



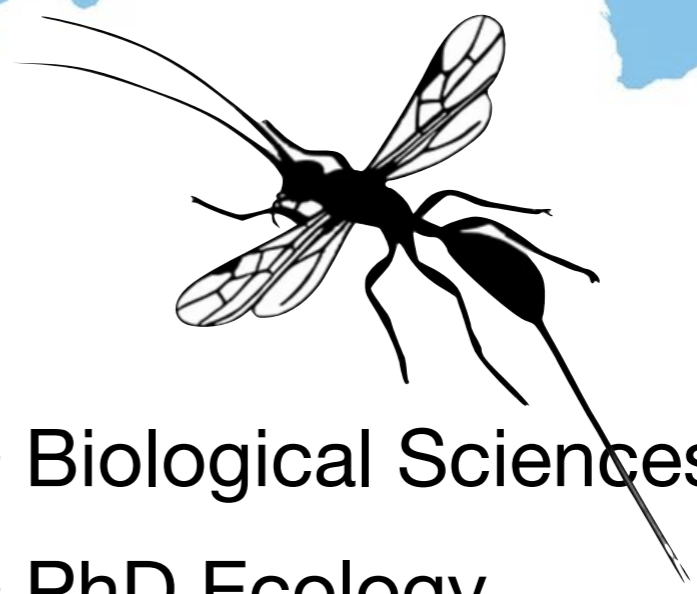
UNIVERSIDAD NACIONAL  
AUTÓNOMA DE MÉXICO



• Current: postdoc



**Universidade Federal  
de São Carlos**



• Biological Sciences  
• PhD Ecology

# Topics

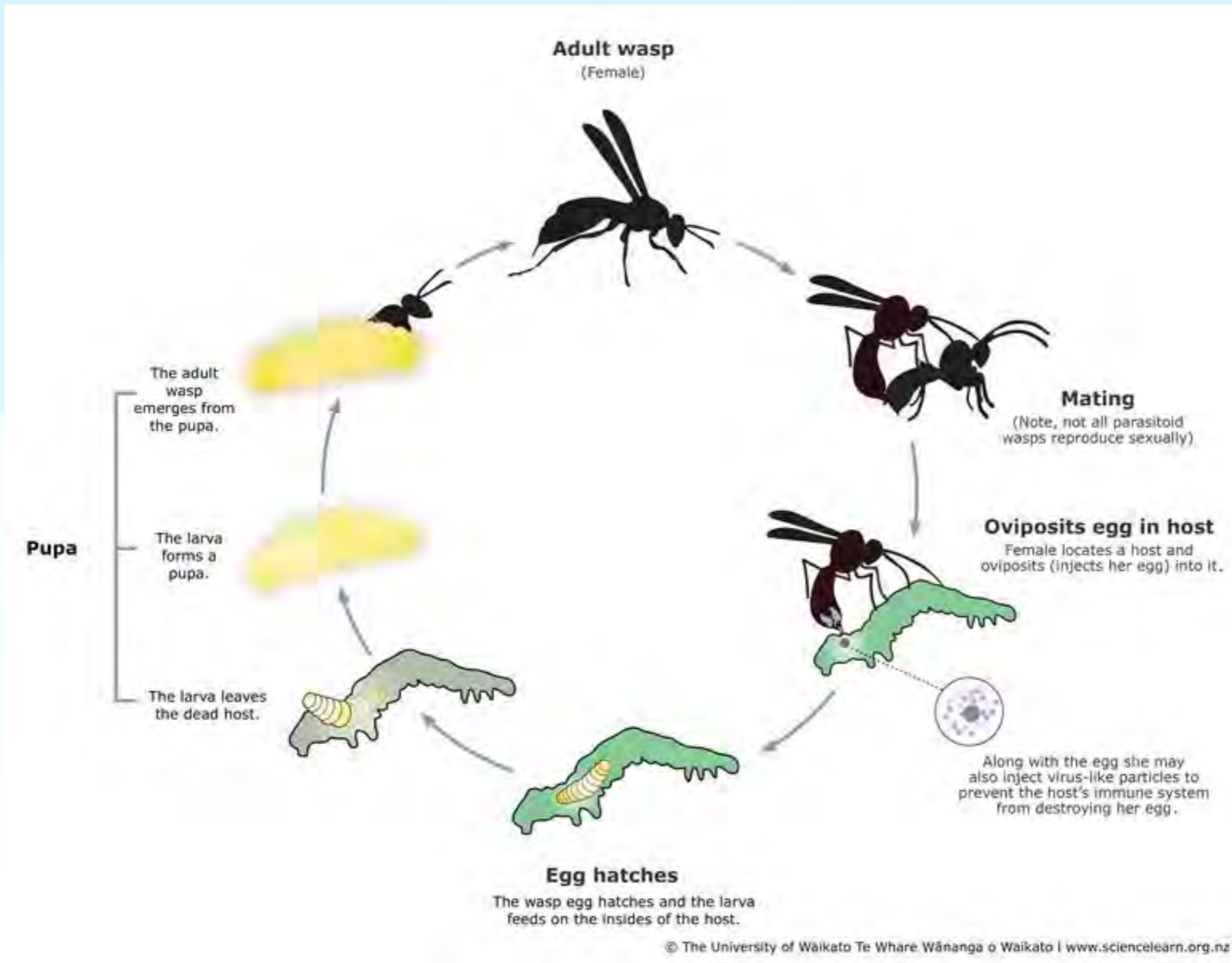
## research centred on parasitoid wasps

1. Parasitoid basics - biology, ecology and diversity
2. Braconidae basics - systematics, biology and diversity
3. The taxonomic gap and the taxonomic impediment
4. How deep is the gap?
5. Can we bridge the gaps?
6. Biological control and taxonomy of parasitoids
7. Current projects



# 1. Parasitoids basics

## Biology, ecology and diversity





# 1. Parasitoids basics

## Biology, ecology and diversity

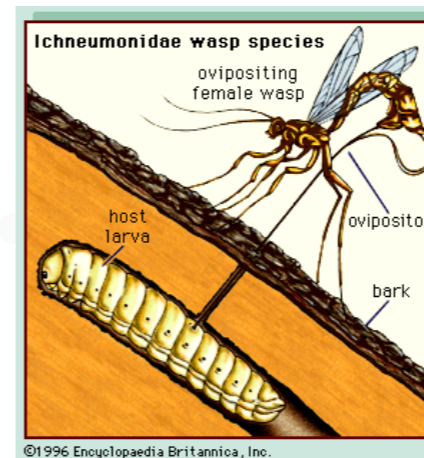
Hymenoptera: venom glands and ovipositor



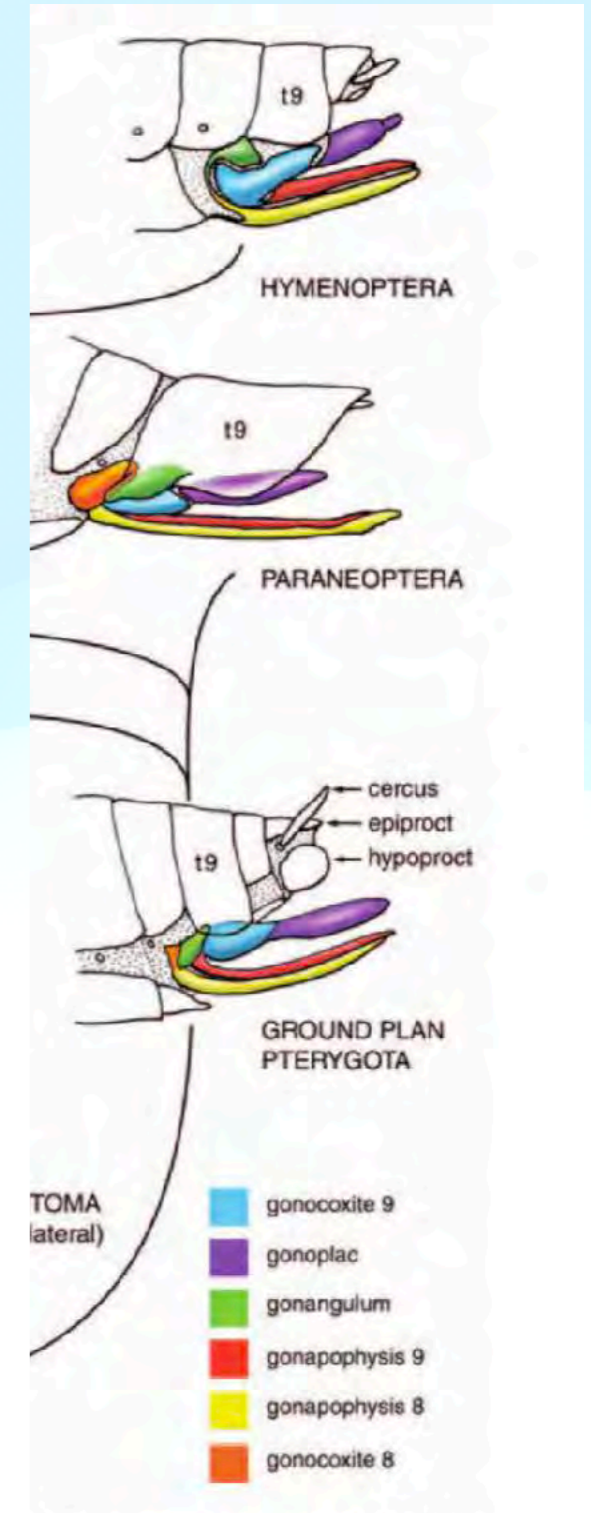
*Leptopilina*



Venom gland (VG) and reservoir (R)



©1996 Encyclopaedia Britannica, Inc.

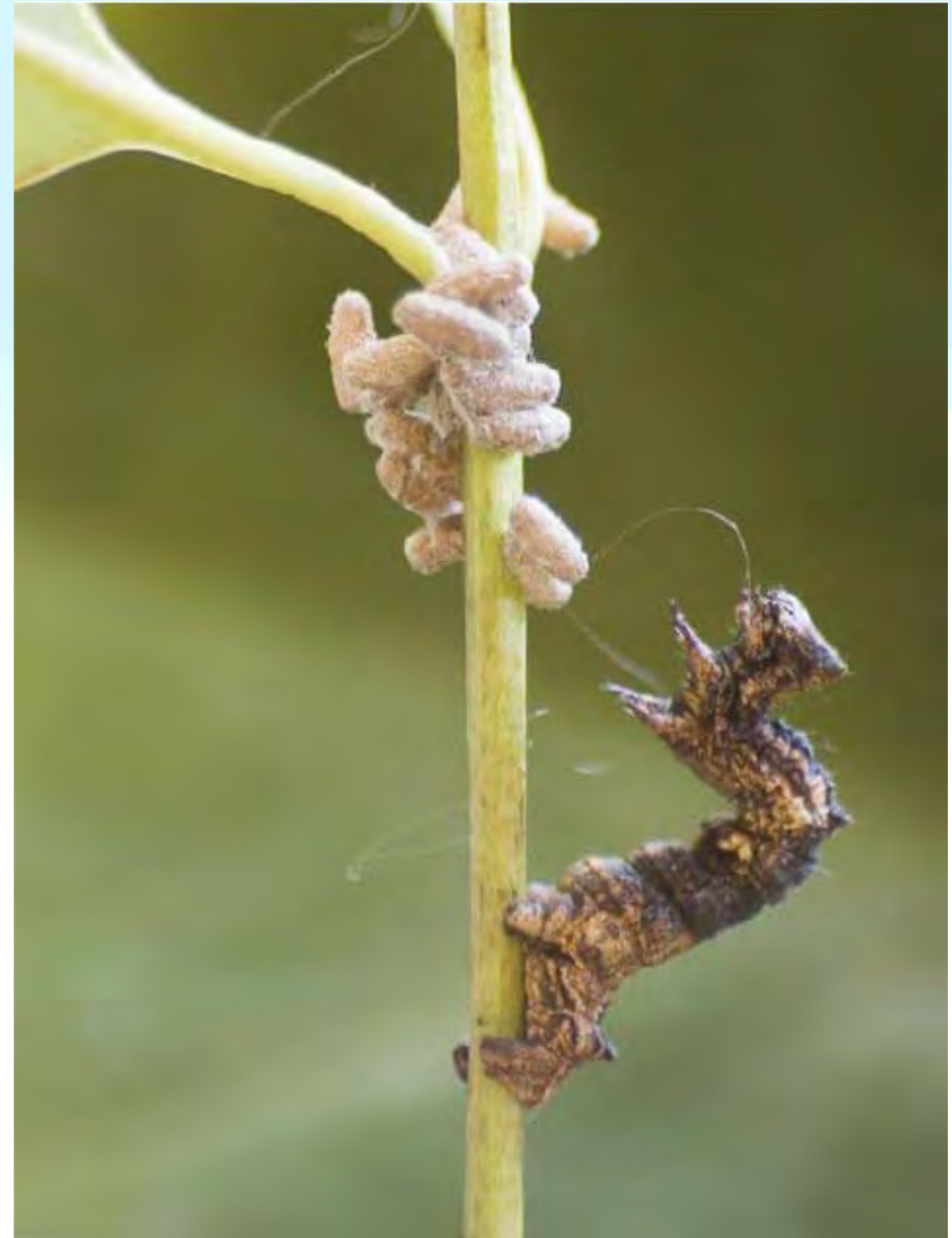


# 1. Parasitoids basics

## Biology, ecology and diversity

*Glyptapanteles* sp.

*Thyrinteina leucocera*



Grosman et al 2008, PLoS One 3(6)



# 1. Parasitoids basics

## Biology, ecology and diversity



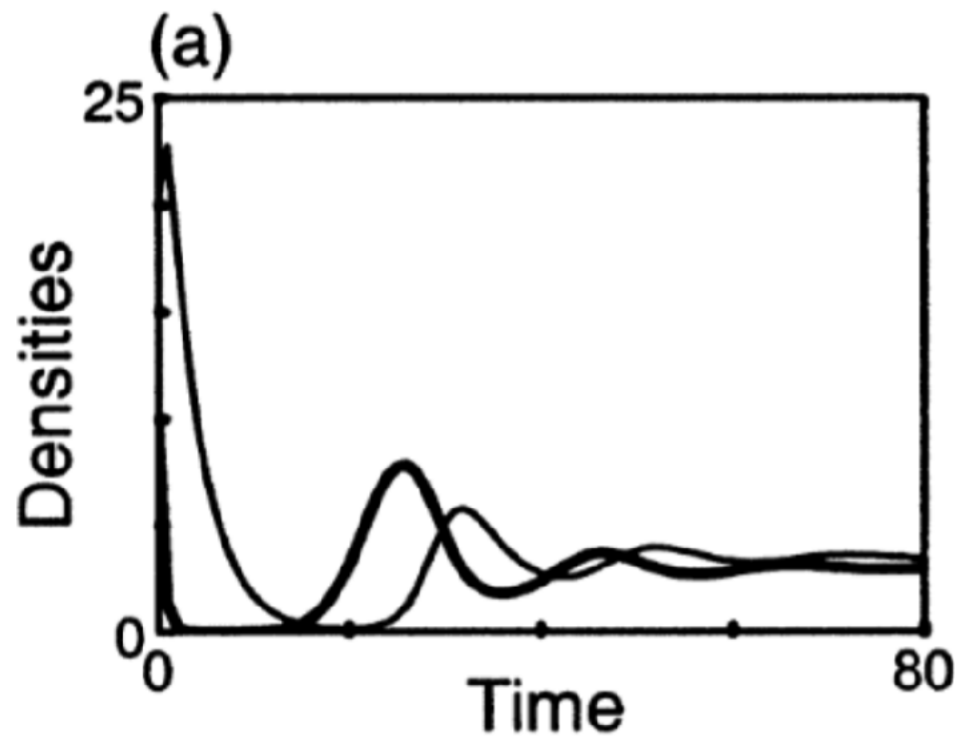
Pontoppidan et al 2009, PLoS One 4(3)

# 1. Parasitoids basics

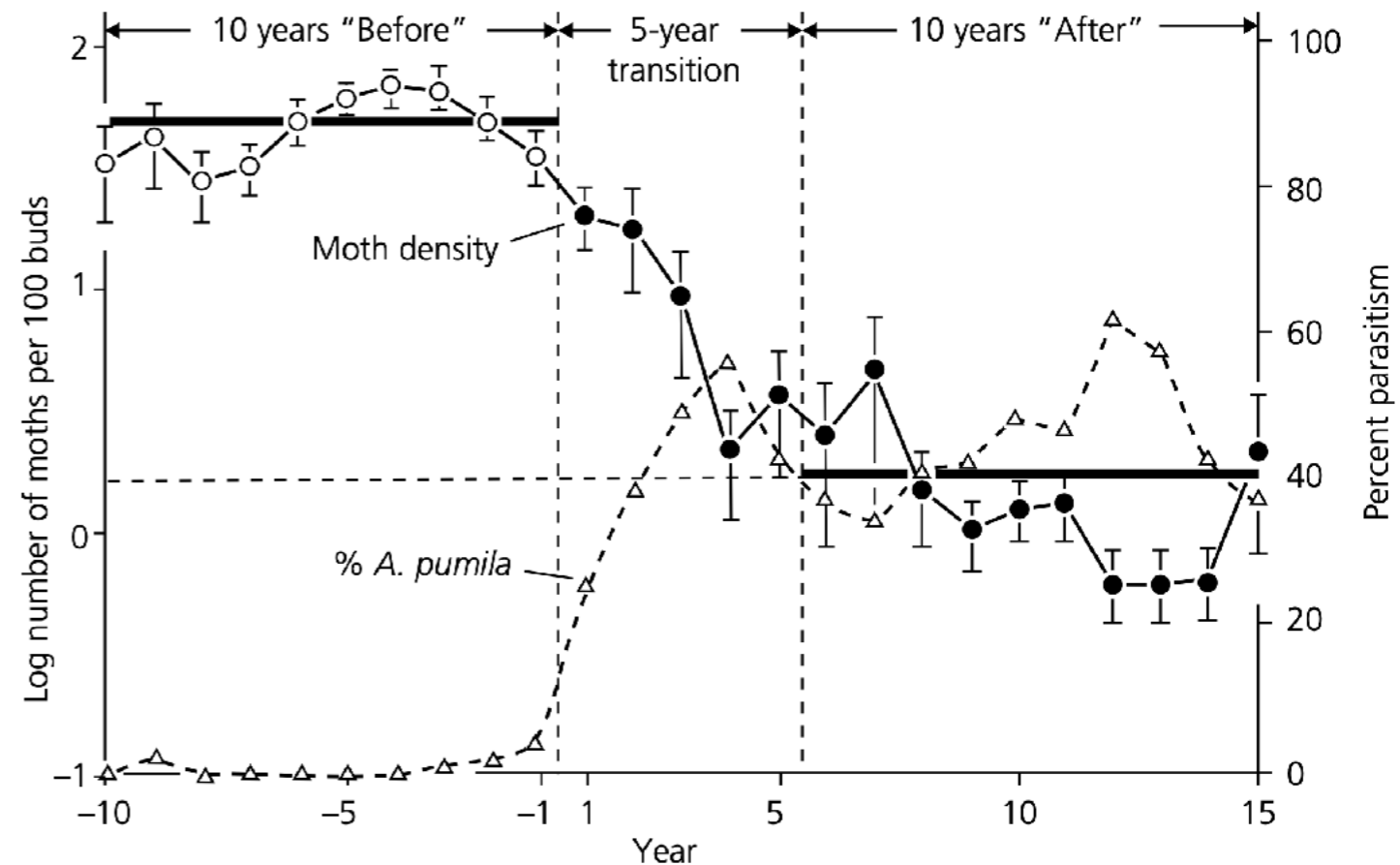
## Biology, ecology and diversity



*Coleophora laricella* density variation in Oregon (USA) for 25 yr relative to the introduction of *Agathis pumila* (Ryan 1997)



Simulation of the Lotcka-Volterra equation model plus logistic prey,  $K=10$  (Murdoch et al 2003; Consumer-Resource Dynamics)





# 1. Parasitoids basics

## Biology, ecology and diversity

- Ecosystem services are valuable
- Natural control of pest in USA = US\$ 4.5 billion/year\*
- Africa: *Anagyrus lopezi* against cassava mealybug = US\$8–20 billion /40 years\*\*
- food security

\* Losey & Vaughan 2006, *BioScience* 56(4)

\*\* Zeddies et al. 2005, *Agr Econ* 24(2)



# 1. Parasitoids basics

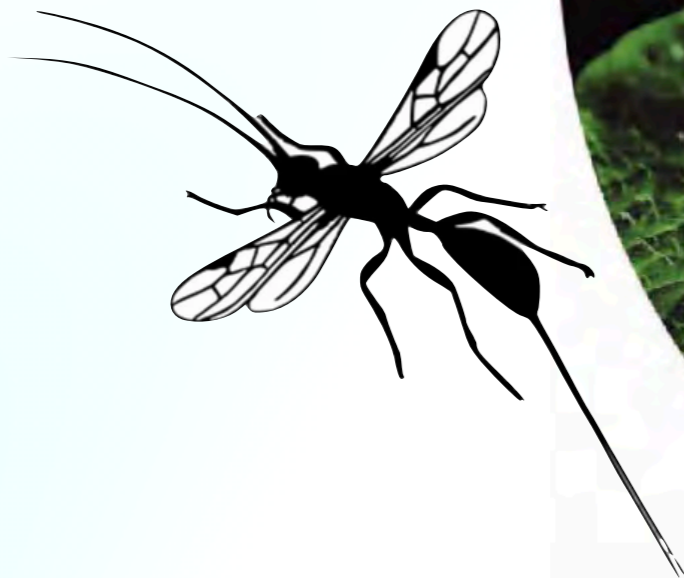
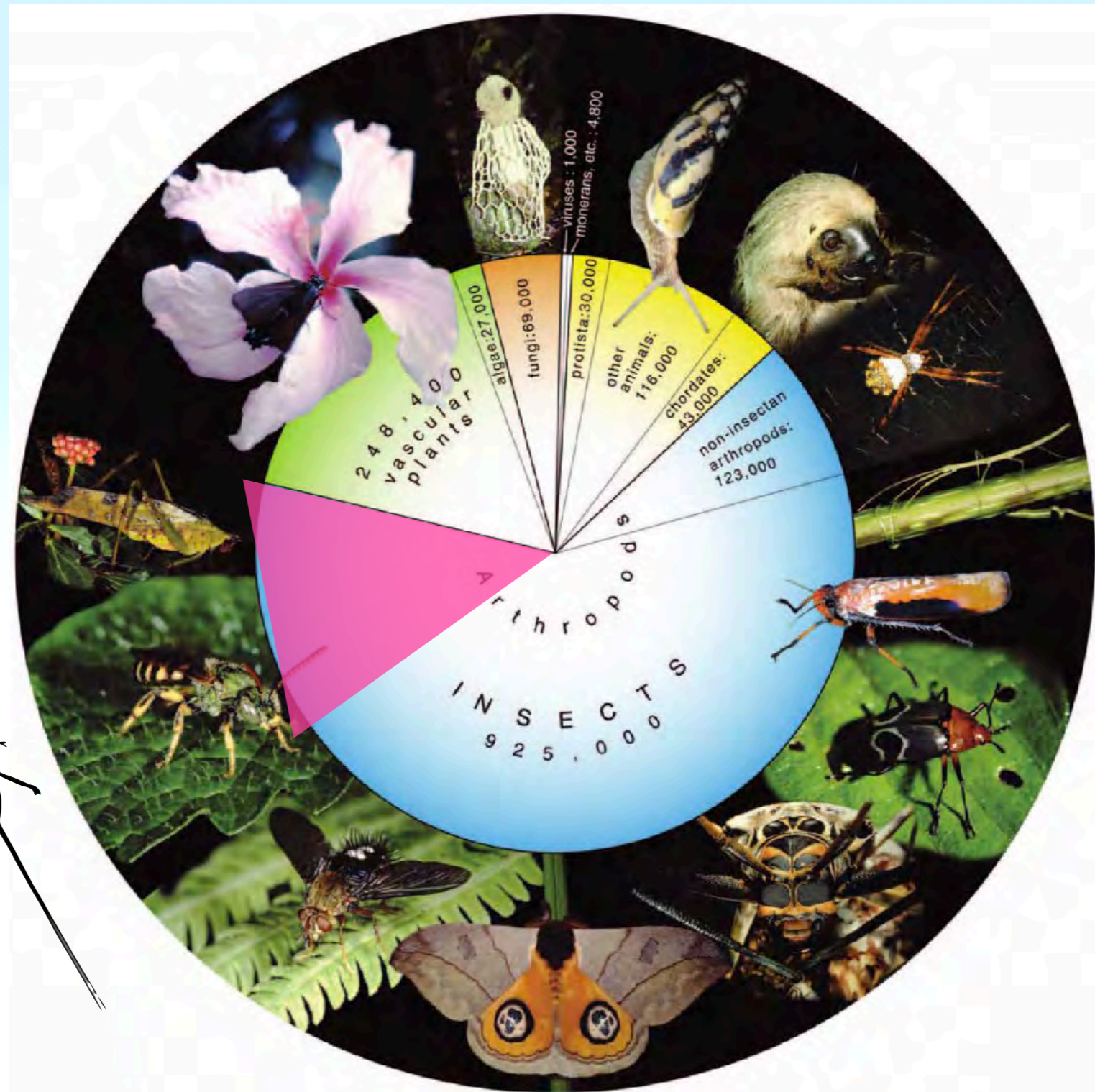
## Biology, ecology and diversity





# 1. Parasitoids basics

## Biology, ecology and diversity

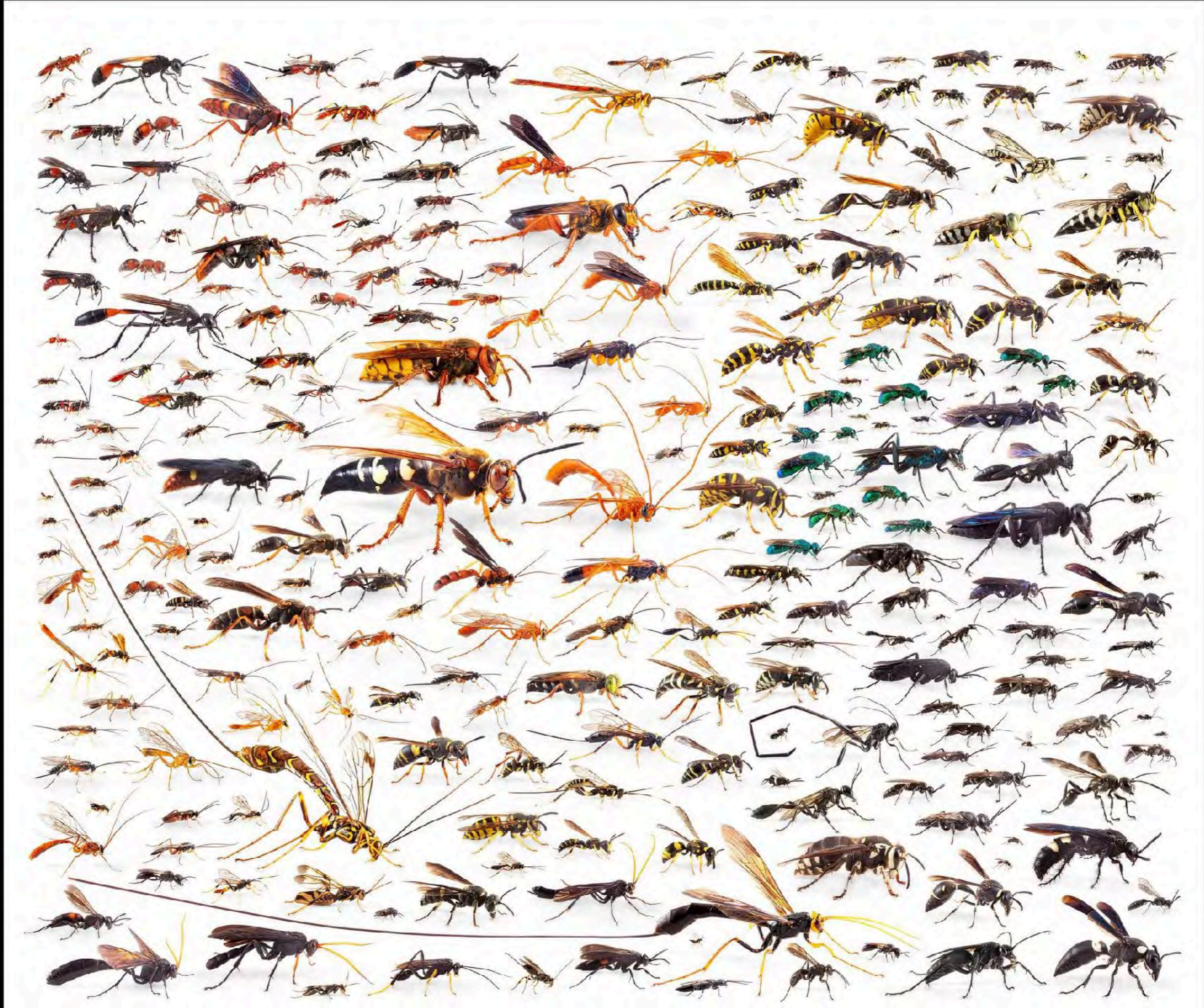




Sloan Tomlinson @thatwaspguy







Sloan Tomlinson @thatwaspguy



# 1. Parasitoids basics

## Reasons to chose them

- Very cool life histories;
- A huge part of the biodiversity in any terrestrial ecosystem;
- Ability to control populations/ model insects/ biological control;
- Extreme adaptations to parasitic life, including manipulation of host behaviour;
- Having a grad student friend needing help with field-work and sample sorting





# 2. Braconidae

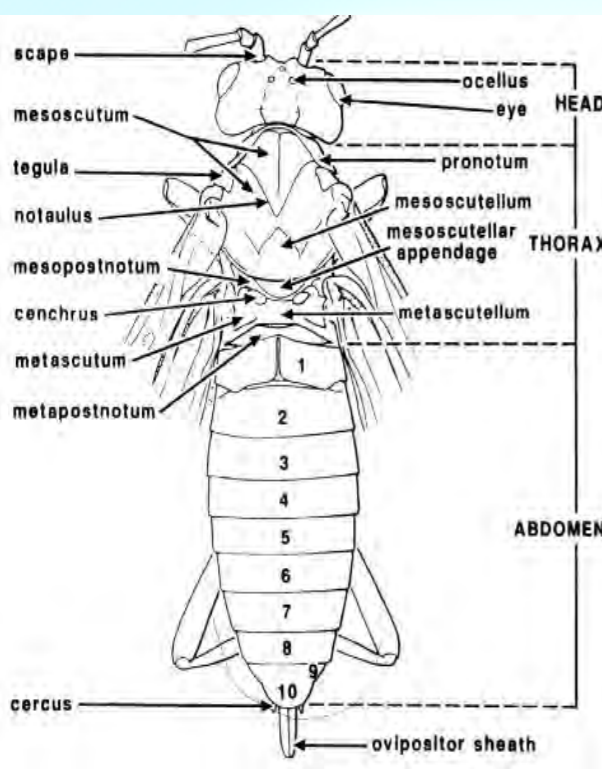
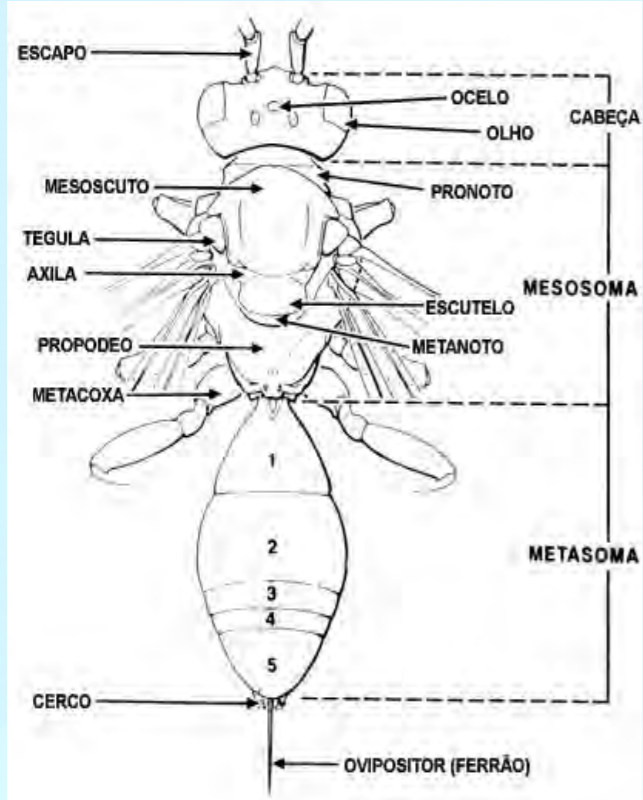
## Phylogeny, morphology and biology

- Braconidae + Ichneumonidae = Ichneumonoidea
- The most diverse families within Hymenoptera
- Braconidae - 1,100 genera in 42 subfamilies; ~22,000 species

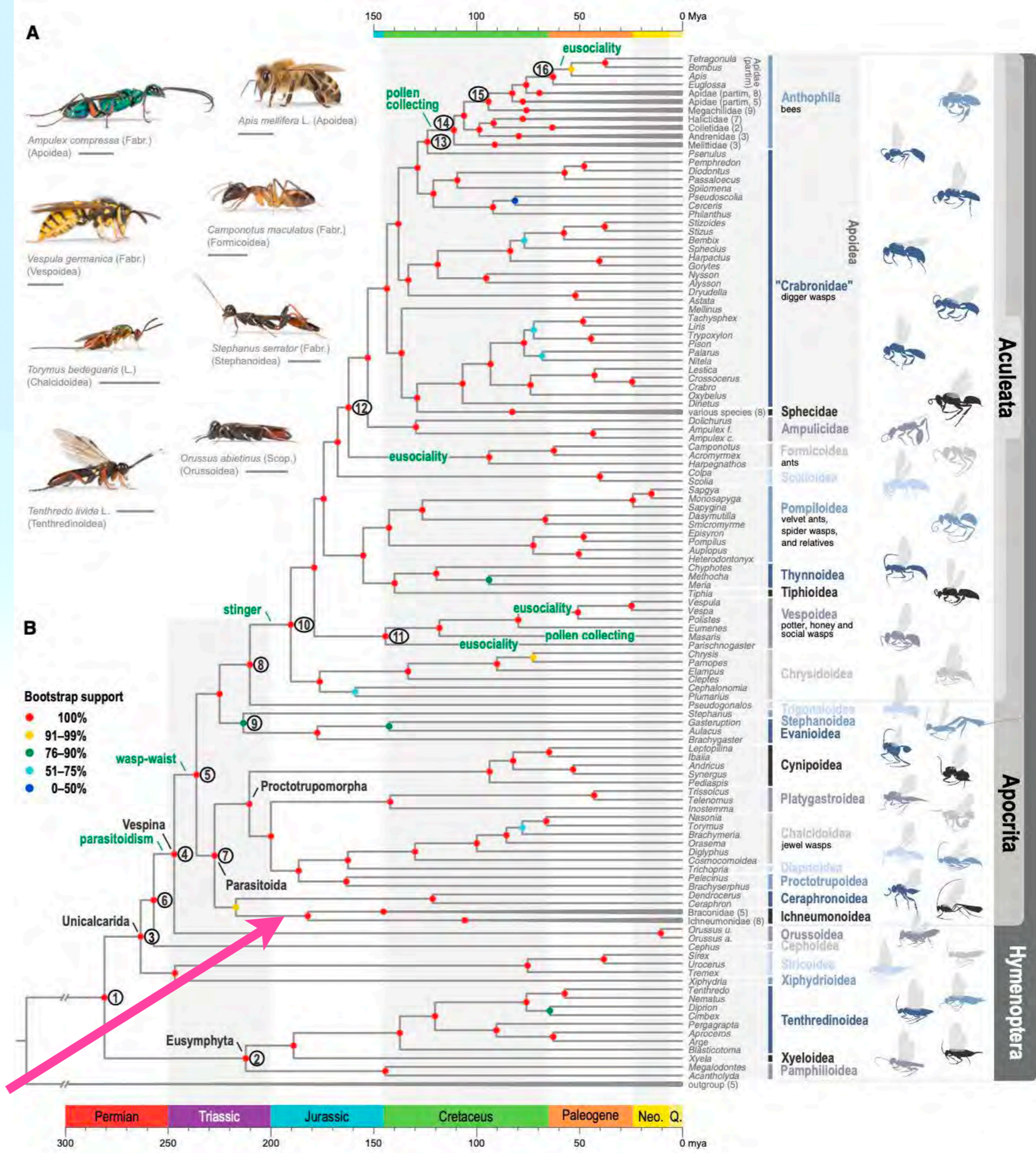




# Phylogeny

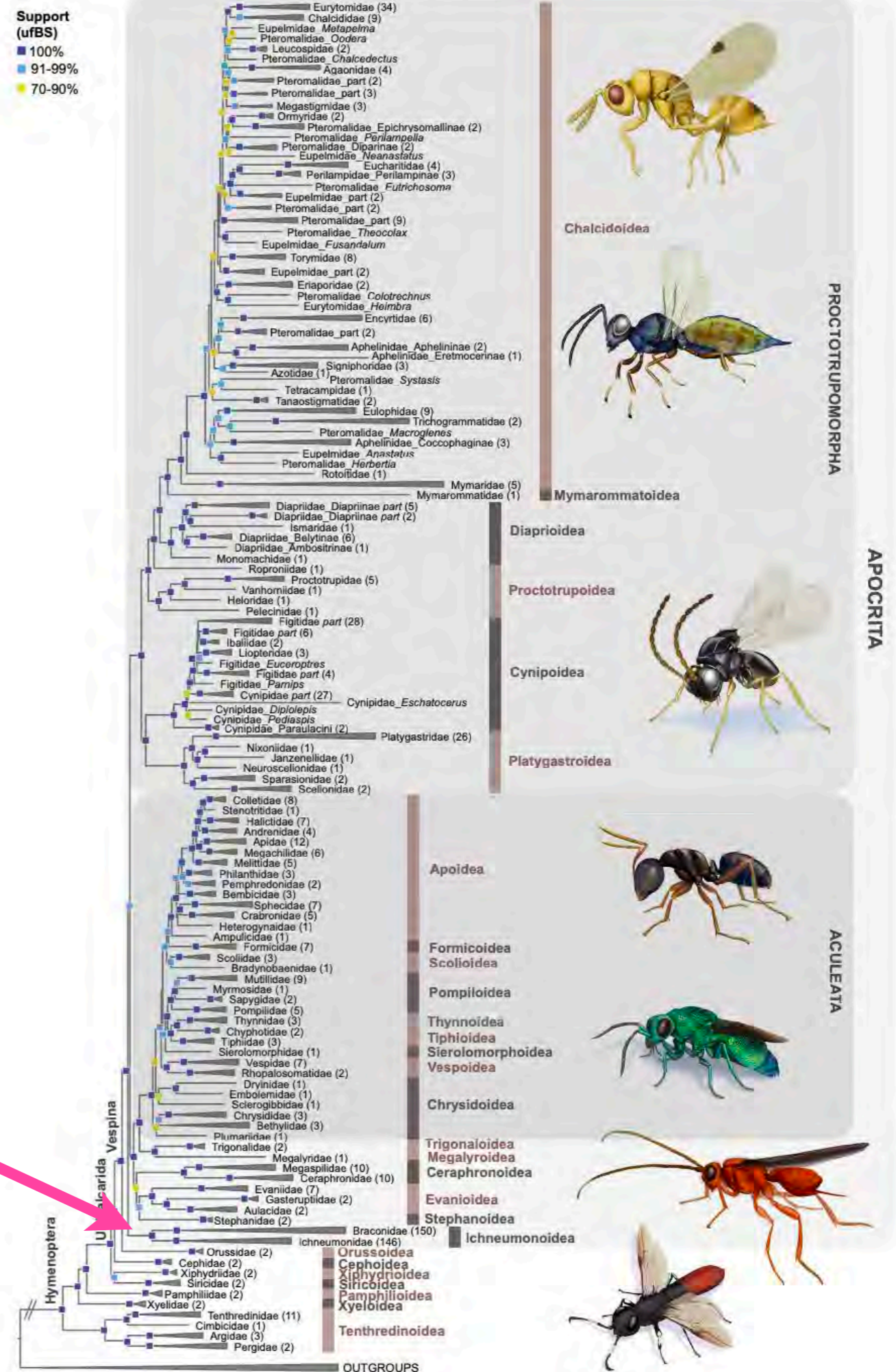


Peters et al. 2017





# Phylogeny



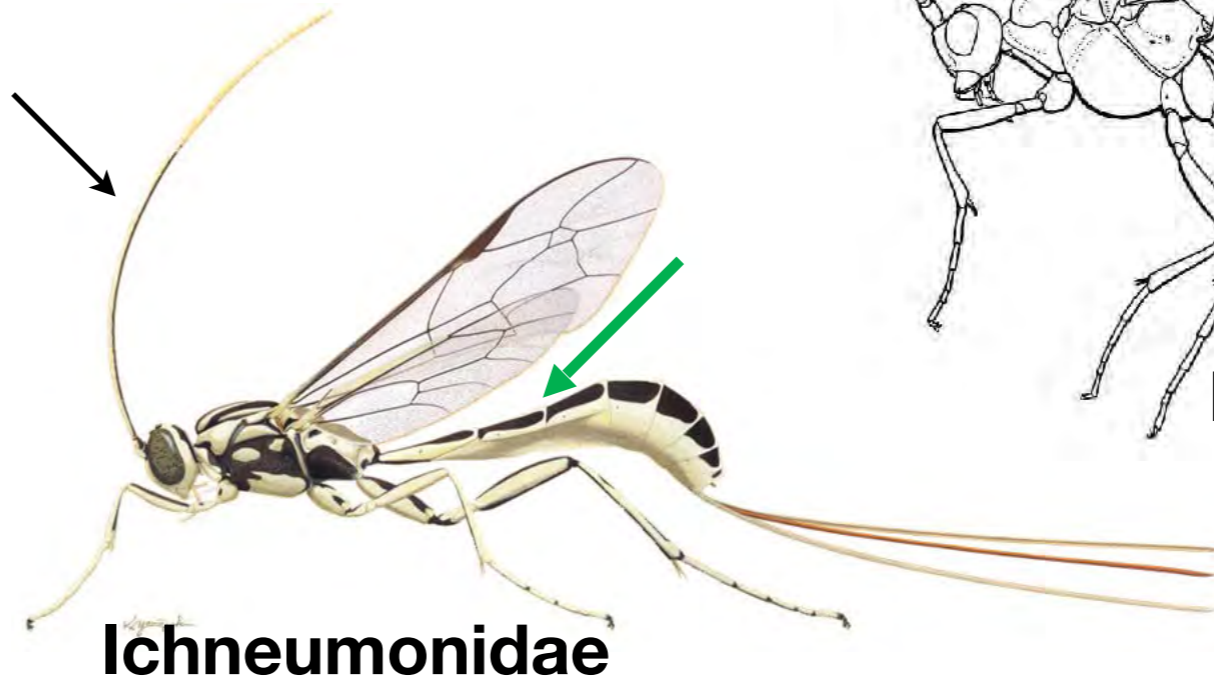
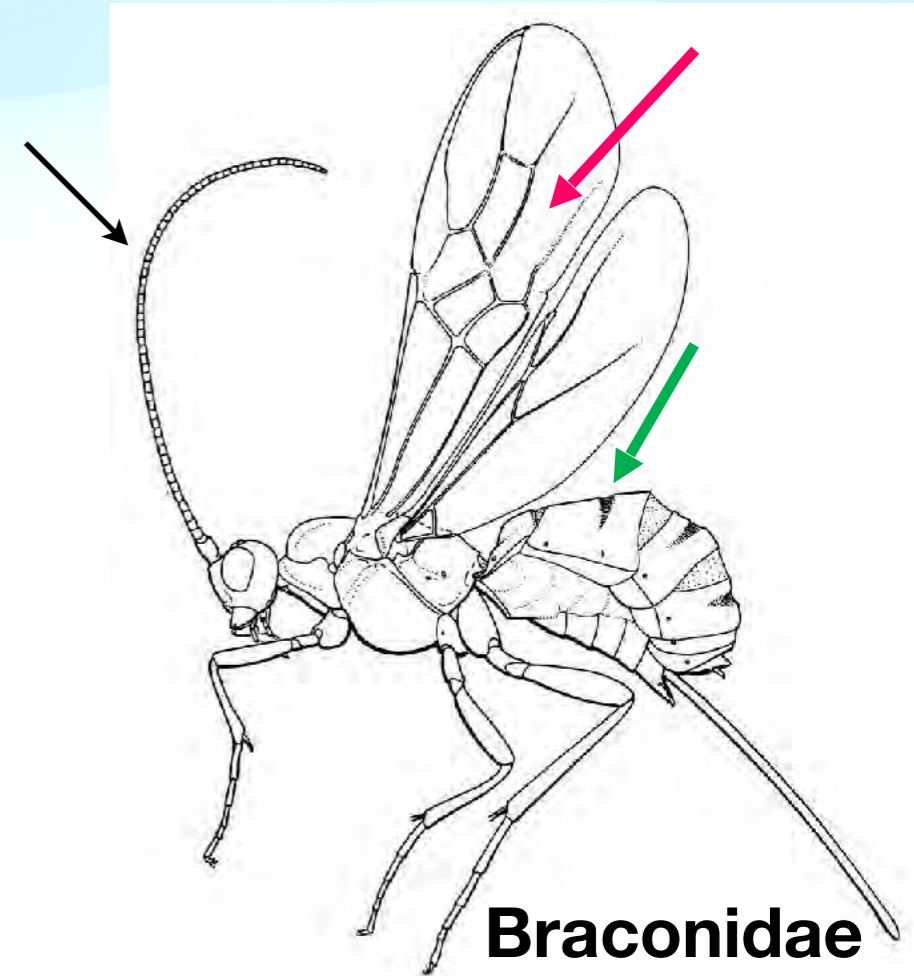
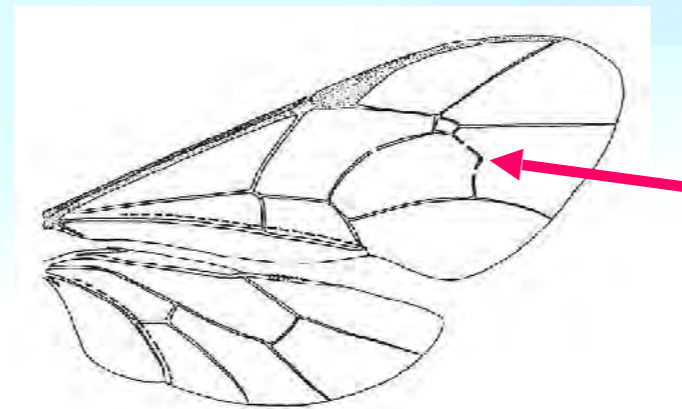
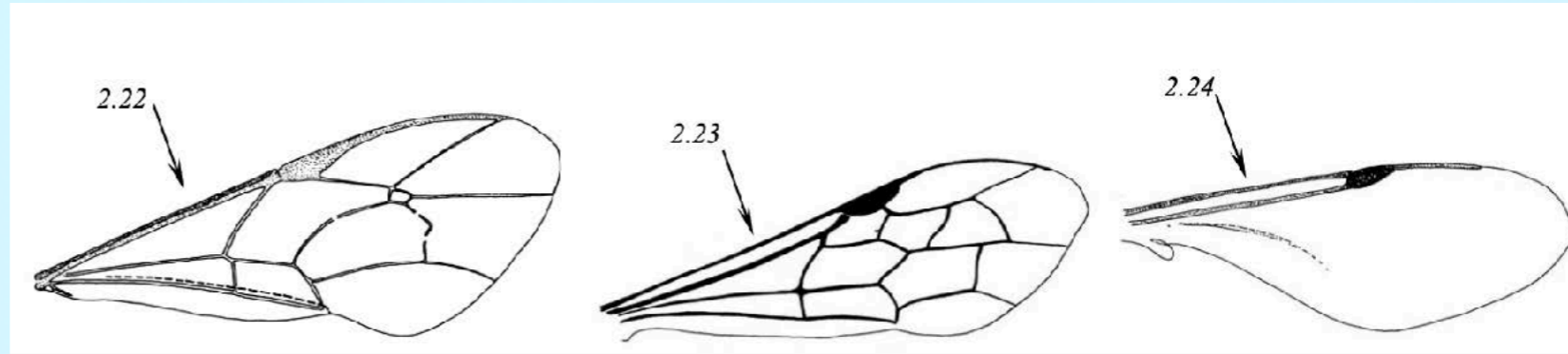
Blaimer et al. 2023



# 2. Braconidae

## Morphology

- Ichneumonoidea: wing venation (costal cell absent -2.22) and antenna filiform.
- Braconidae vs Ichneumonidae: T2+3 (and absence of vein 2m-cu - the second recurrent vein)





# 2. Braconidae

## Biology

- Most Braconidae are larval parasitoids of holometabolous insects (Lepidoptera, Coleoptera and Diptera)

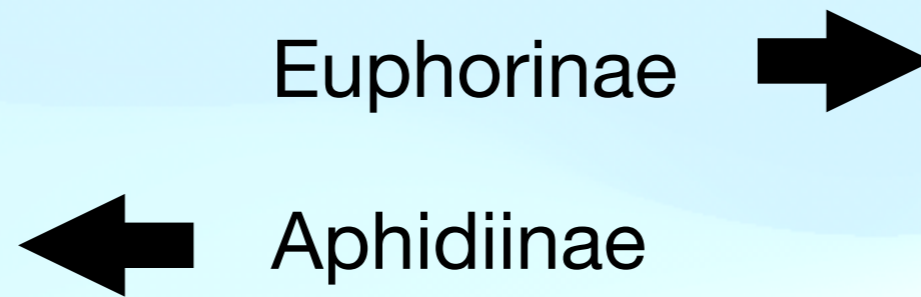




# 2. Braconidae

## Biology

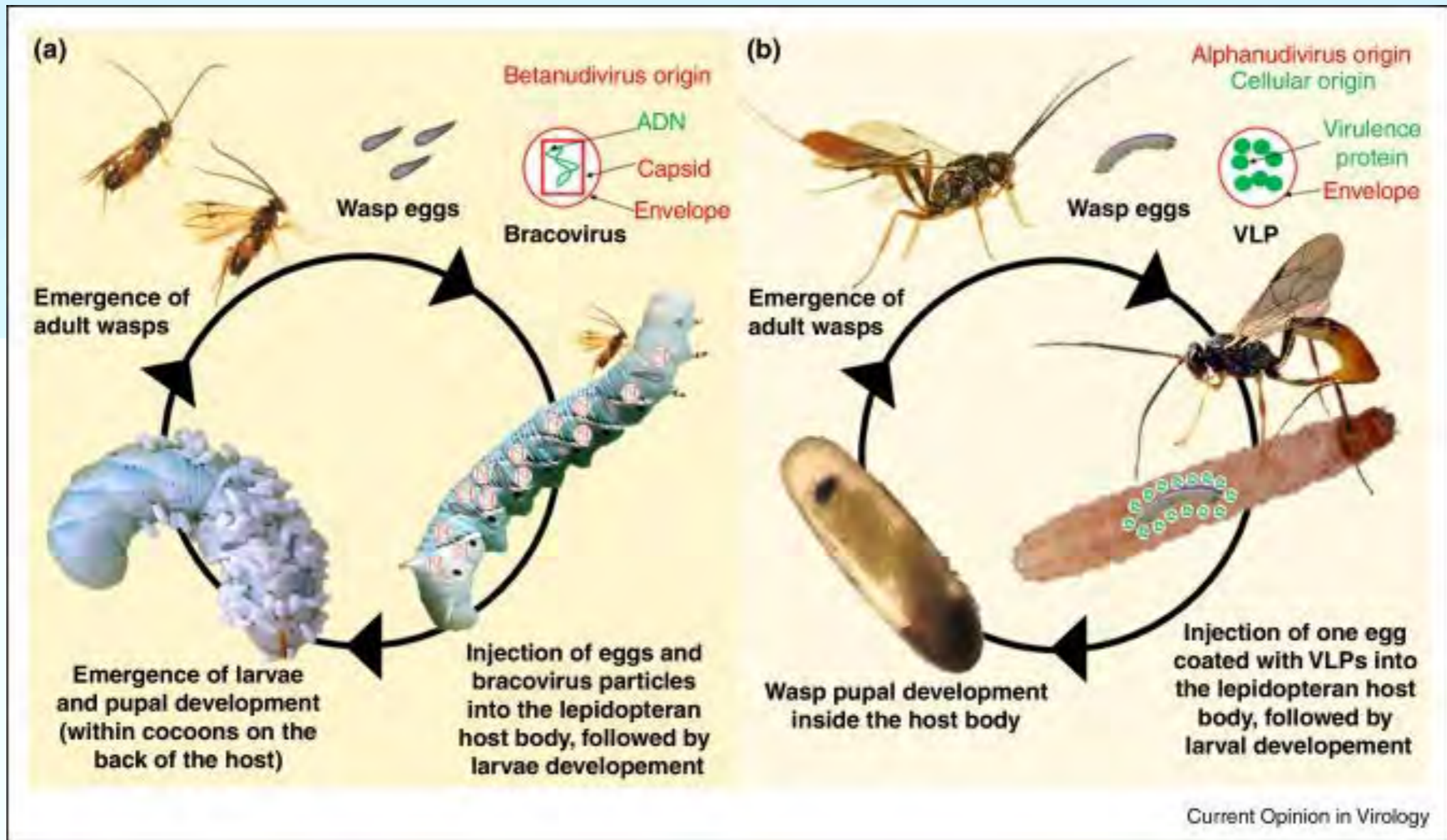
- Exceptions: aphids, adults, phytophagous





# 2. Braconidae

## Polidnavirus

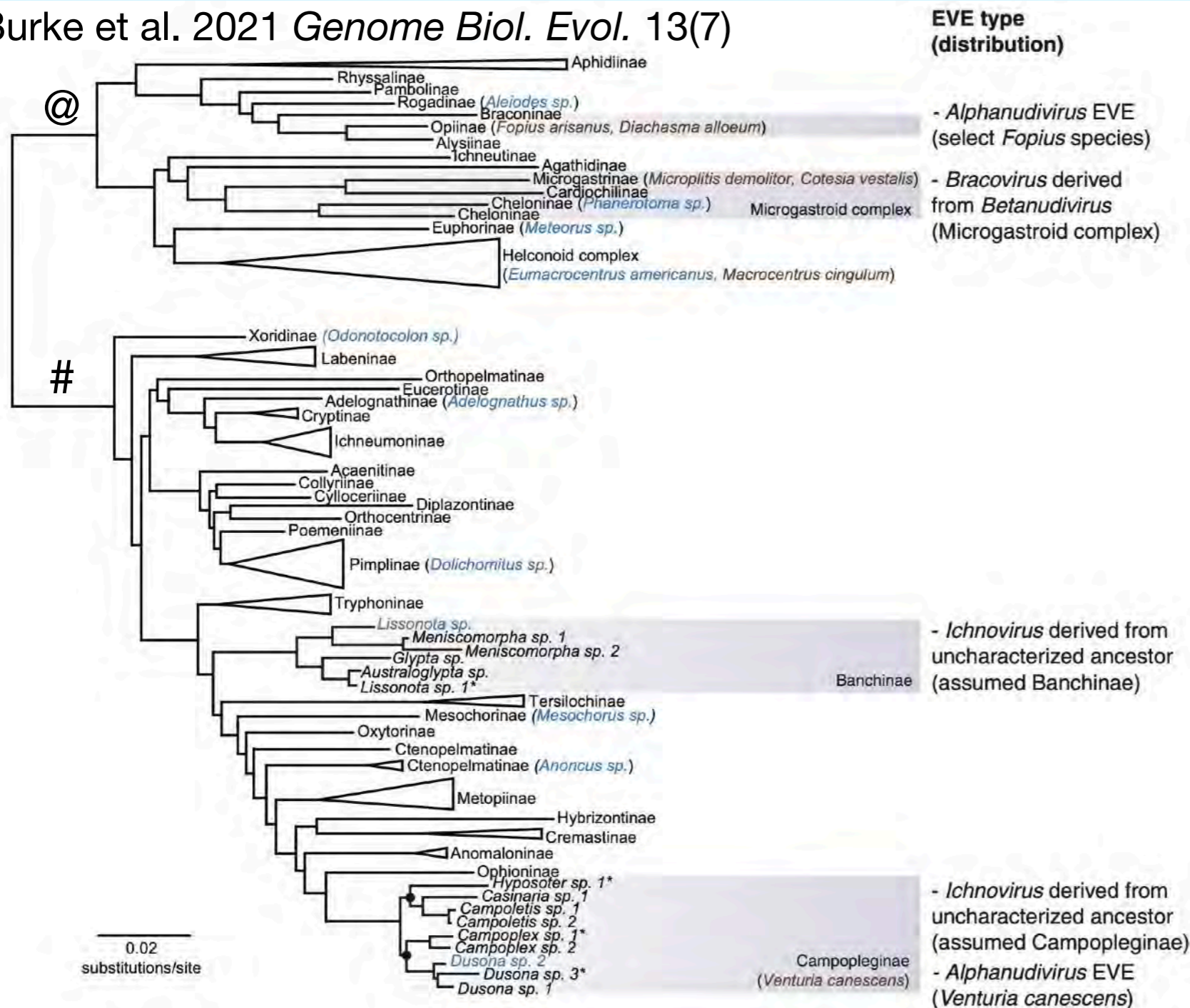




# 2. Braconidae

## Polidnavirus

Burke et al. 2021 *Genome Biol. Evol.* 13(7)



- Polidnavirus are well known from Braco (@) and Ich (#)
- Domestication is widespread (diverse origins) and related to endoparasitism



# 3. The taxonomic gap and the taxonomic impediment

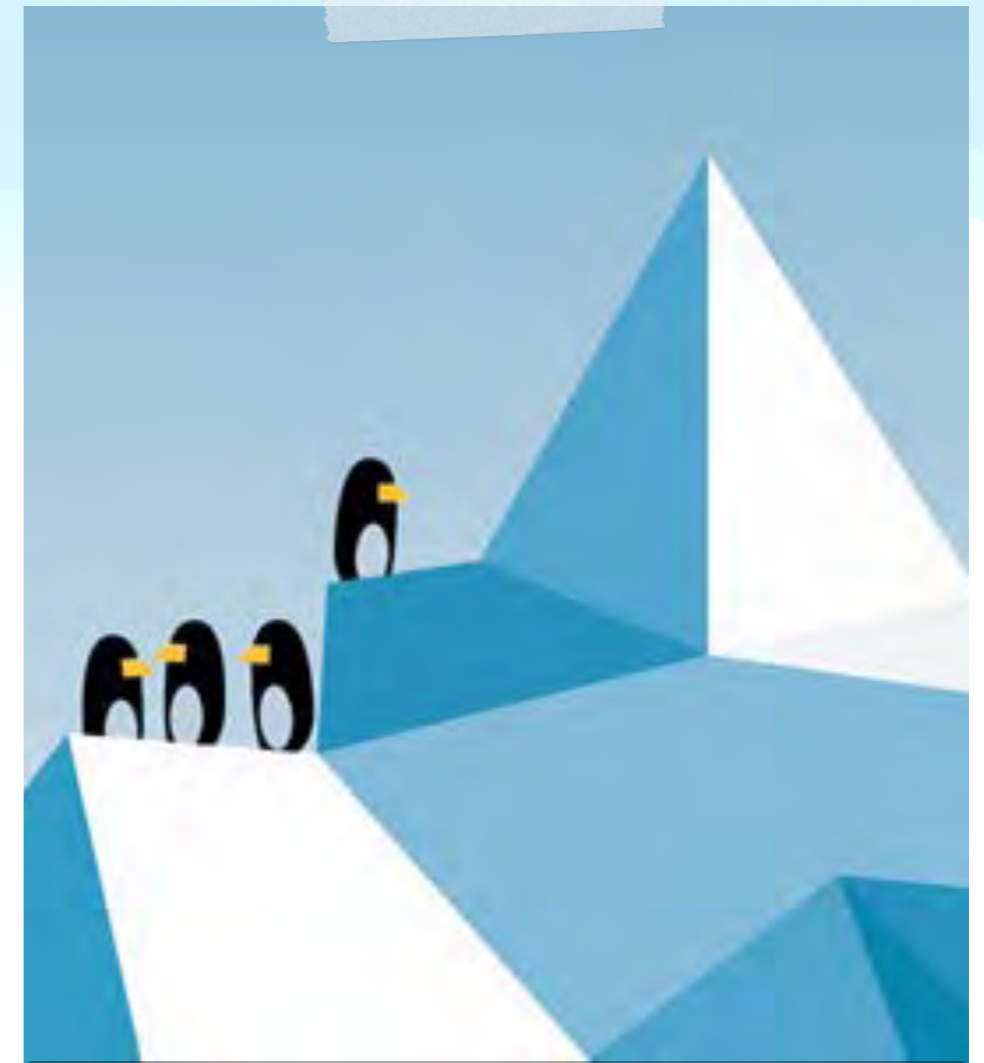
- Most of the **biodiversity remains unknown** (concentrated in the **dark taxa**)
- The biodiversity **crisis is intensifying** - loss of species, habitats and interactions (taxonomic urgency)
- The **arthropods** are essential (ecological roles)
- Biodiversity-based sciences depend on taxonomy
- The **taxonomic impediment**: lack of resources, mainly of taxonomists
- The problem is deeper in the **tropics**: fewer studies, more species
- **How do we quantify (grossly) the gap?**
- **How do we deal with this problem?**

*Zoological Journal of the Linnean Society*, 2021, 193, 381–387.

## EDITORIAL

**The taxonomic impediment: a shortage of taxonomists, not the lack of technical approaches**  
et al.

MICHAEL S. ENGEL<sup>1,2,3</sup>, LUIS M. P. CERÍACO<sup>2,3</sup>, GIMO M. DANIEL<sup>3,4,5</sup>,





# 4. How deep is the gap?

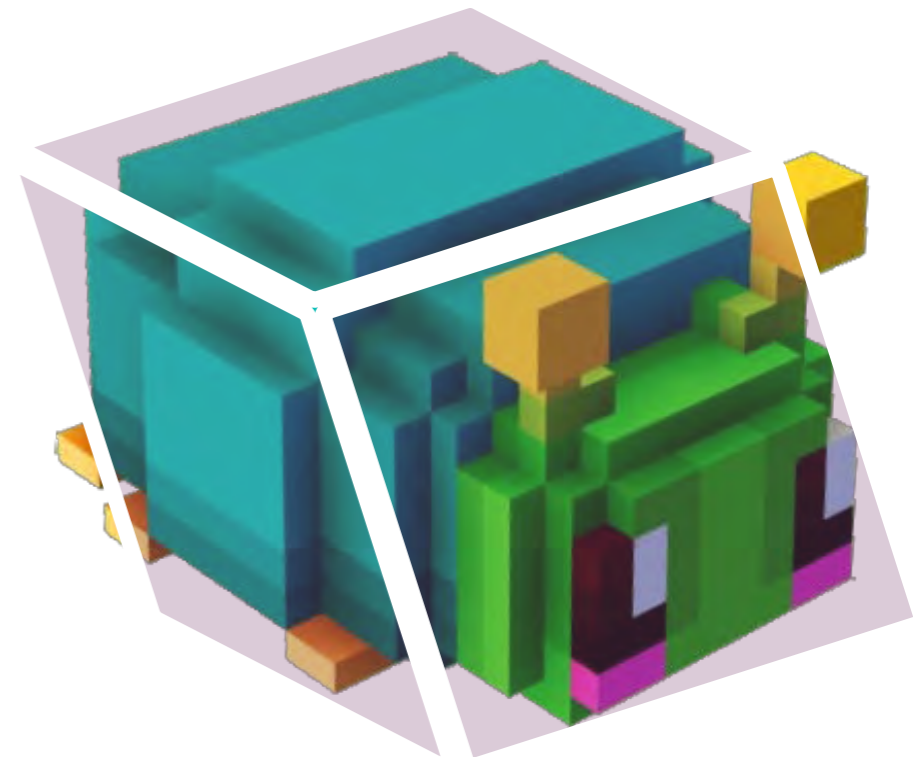
## Estimations - described VS undescribed

Species diversity may be explained in terms of niche availability/creation/differentiation.

Most of a parasitoid species niche is determined by its hosts, as they spend virtually their entire life in it or searching for it.

1:1 proportion — Parasitoid : Host (P:H)

But, of course, things are not so simple





# 4. How deep is the gap?

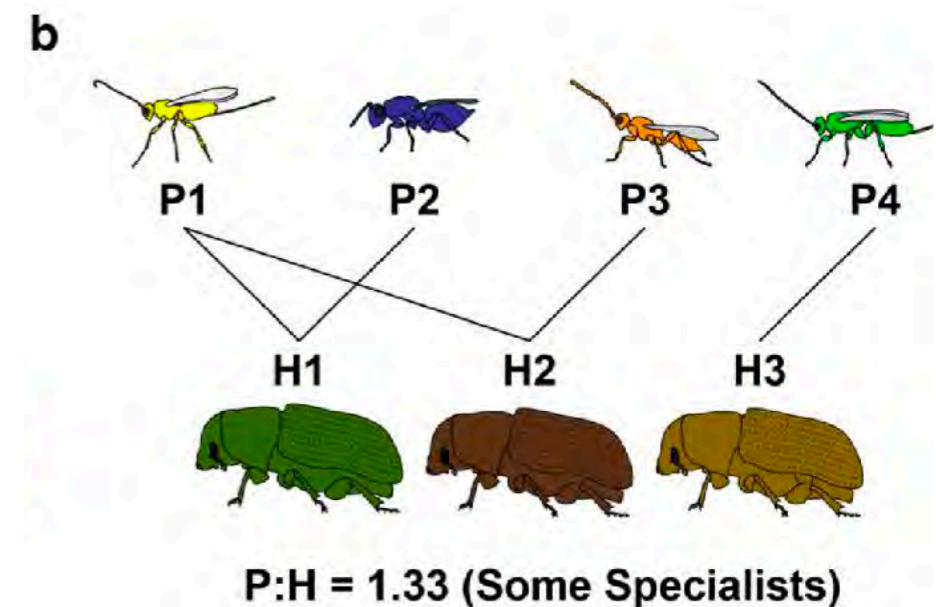
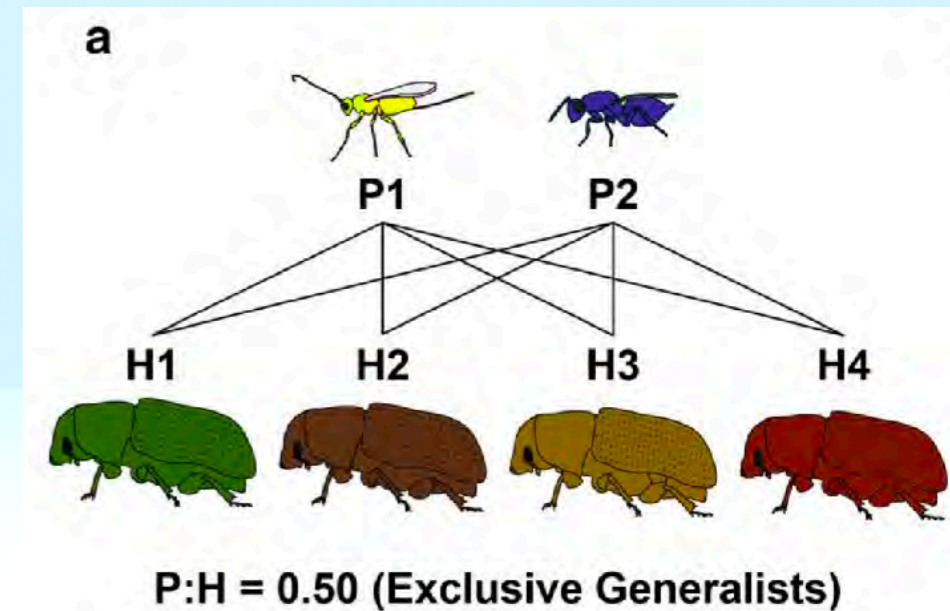
## Estimations - described VS undescribed

Forbes et al. (2018)

Hymenoptera is larger than Coleoptera

P : H based on 4 well-known genera and their parasitoids = 0.95 – 1.83

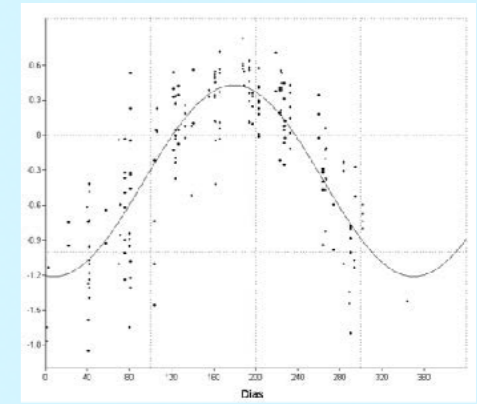
Unrealistic? - ignores the complexity of food webs





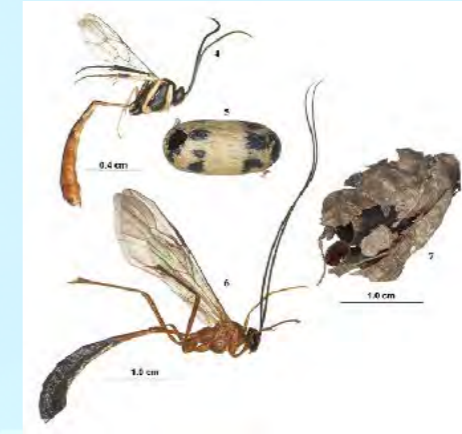
# 4. How deep is the gap?

## Estimations - described VS undescribed

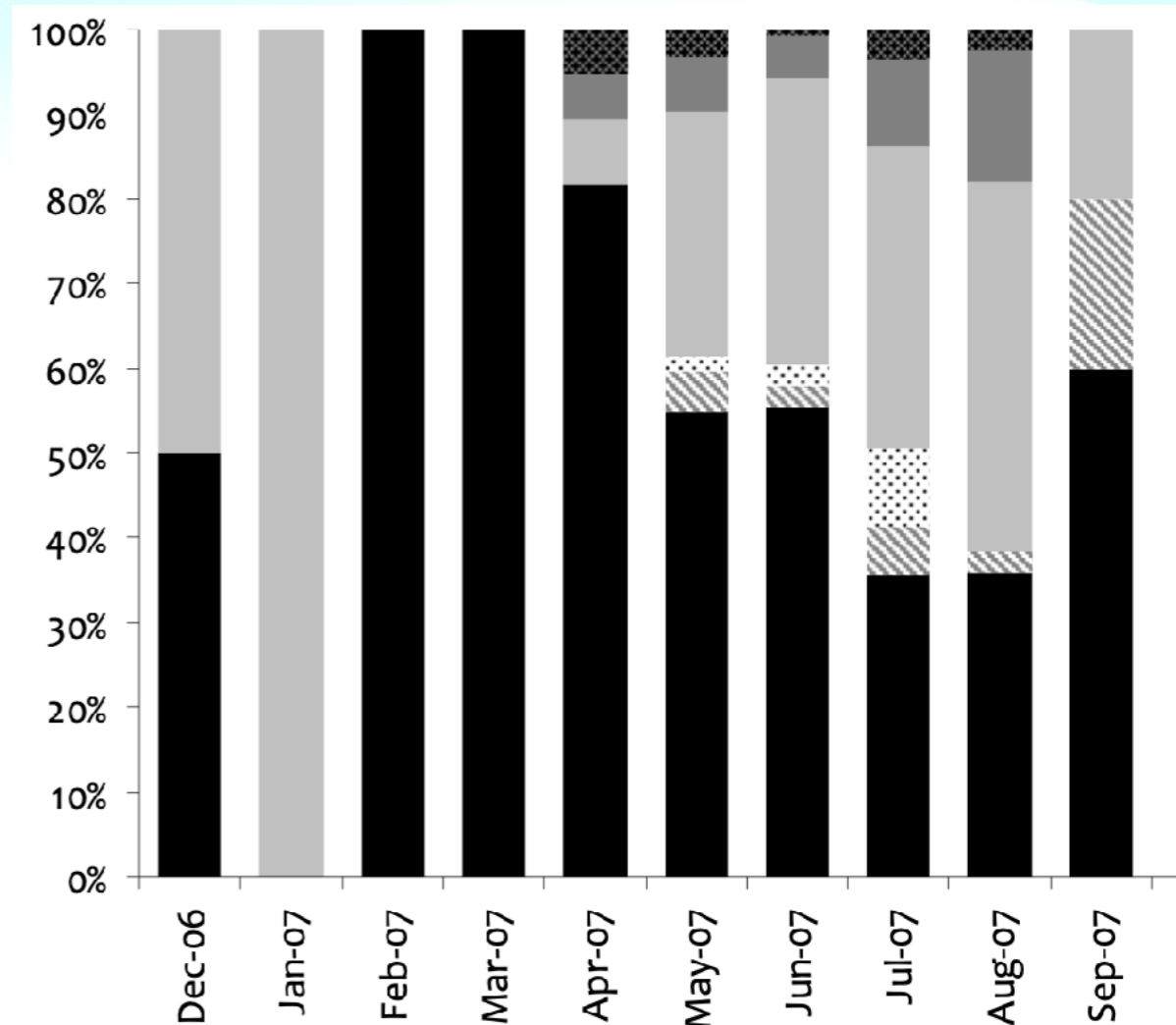


### Real tropical system

1. Based on 2 plant spp.
2. Dominated by 1 herbivore:
  - a. Niche segregation: 21 species in 6 guilds
3. 105 potential hosts (41 actual hosts), 102 primary parasitoids and 21 hyperparasitoids:  
 $P : H = 1.15$  (= Forbes et al. 2018 low estimates)



- Ectoparasitóide
- Larva-Pupa
- Pré-Pupa
- ⊙ Larva tardia
- ▨ Meteorus
- Larva jovem



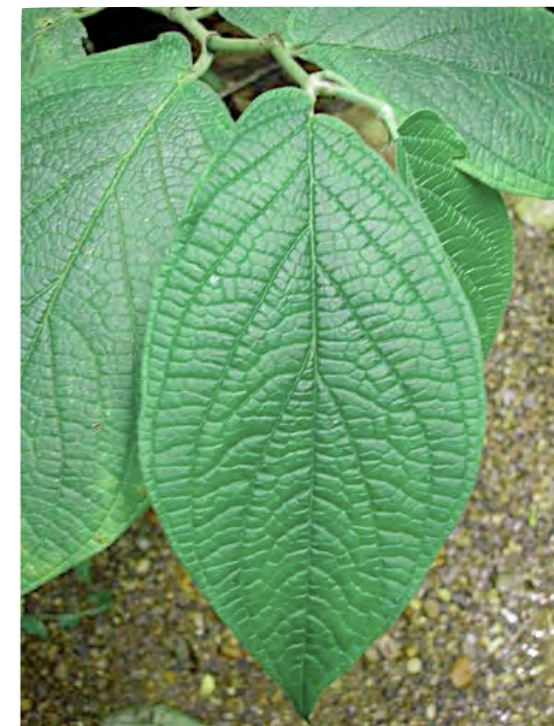
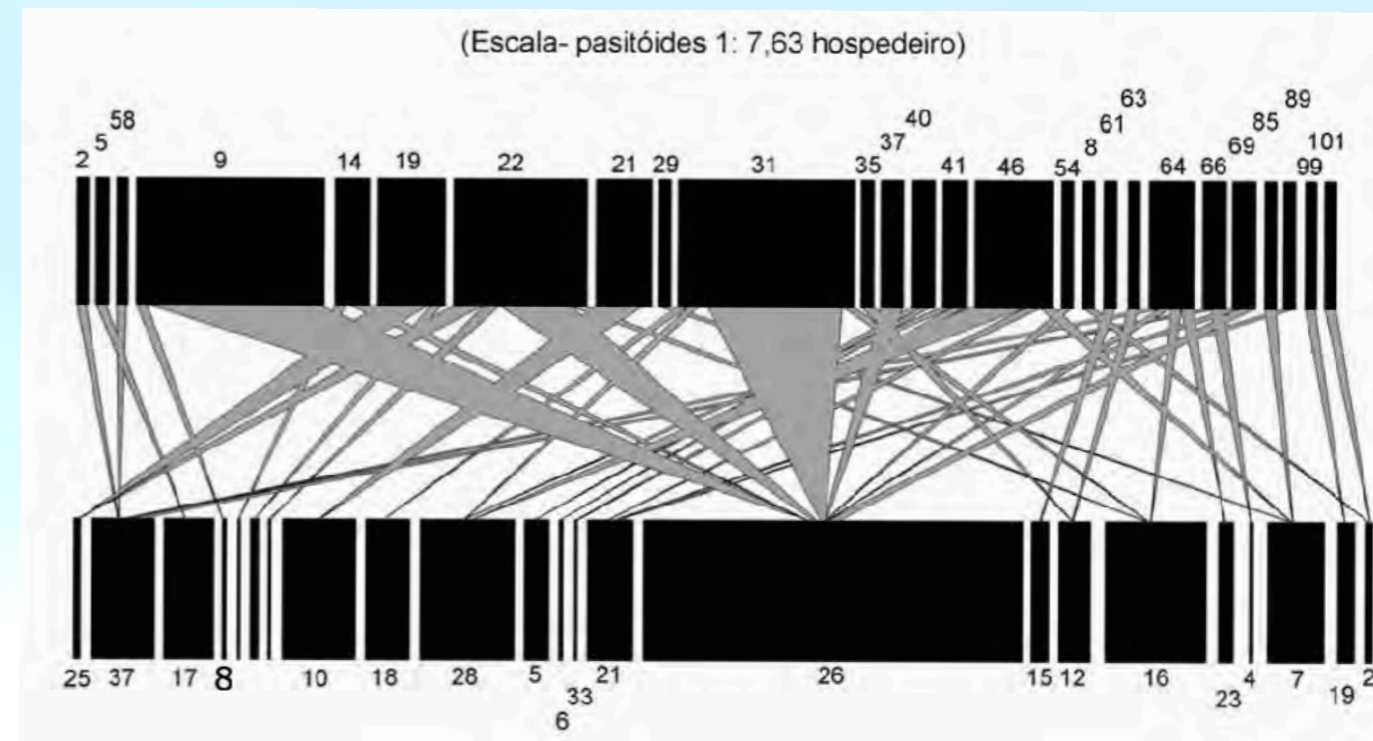


# 4. How deep is the gap?

Estimations - described VS undescribed

## Real tropical system

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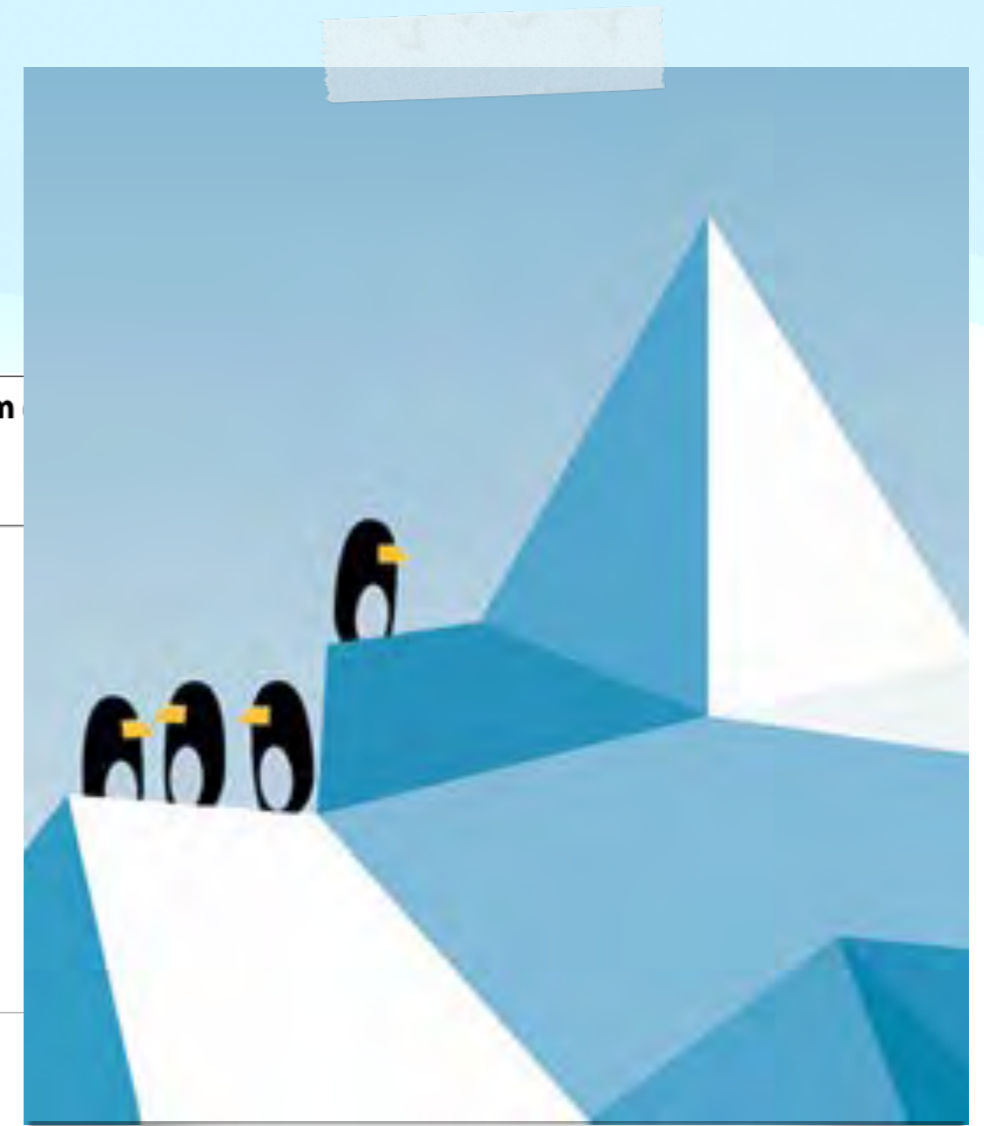




# 4. How deep is the gap?

## Estimations - described VS undescribed

740 thousand to 1 million species of undescribed Hym. parasitoids



Described species ~154,000	High P:H estimates from case studies	Low P:H estimates from studies
Diptera (152,244)	228,366	199,440
Lepidoptera (156,793)	286,931	156,793
Coleoptera (359,891)	494,850	406,677
Non-parasitoid Hymenoptera (~62,000)	79,980	58,900
All other insect orders (335,970)	0 <sup>a</sup>	0 <sup>a</sup>
Total parasitoid Hymenoptera	1,107,487	833,590
Non-parasitoid Hymenoptera (to add to calculated parasitoid numbers)	62,000	62,000
Total Hymenoptera	1,152,127	883,810

Estimates for Microgasterinae: 0.1 P : H  
(Rodriguez et al. 2013) = Brazil = 1-2% spp.

90-92%

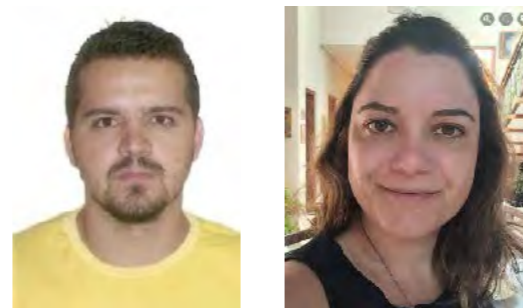
# 4. How deep is the gap?

## Estimations - described VS undescribed

### Examples from intensive surveys

Microgastrinae: 24 genera, 434 morphospecies (1 yr) — Brazil: 23 genera, 126 species. (Gomes 2023)

Orthocentrinae: 13 genera, 127 MOTUS (1 yr) — Brazil - 4 genera, 7 species. (Camargo et al 2021)





# **5. Bridging the taxonomic gap and reducing the taxonomic impediment?**

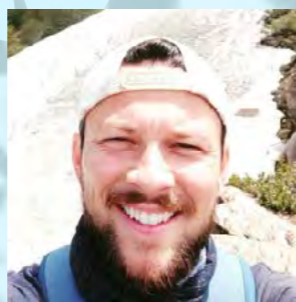
- How do we deal with this problem?

# 5. Bridging the taxonomic gap and reducing the taxonomic impediment?

- How do we deal with this problem?
  - Training/Education in Universities and Collections (1/2)







**Diego Pádua**  
Ichneumonidae/UCM-Chile



**Wesley Colombo**  
Bethylidae/UFES



**Cecília Waichert**  
Pompilidae/UnB



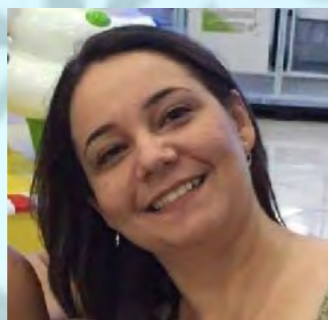
**Daniell Fernandes**  
Ichneumonidae/INPA



**Juliano Nunes**  
Braconidae/UEMG



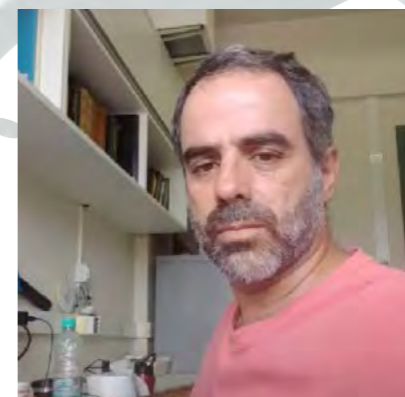
**Eduardo Shimbori**  
Braconidae/UNAM



**Helena Onody**  
Ichneumonidae/UEPI



**Nelson Perioto**  
Eurytomidae/IB



**Marcelo Tavares**  
Chalcididae/UFES



**Celso Azevedo**  
Bethylidae/UFES

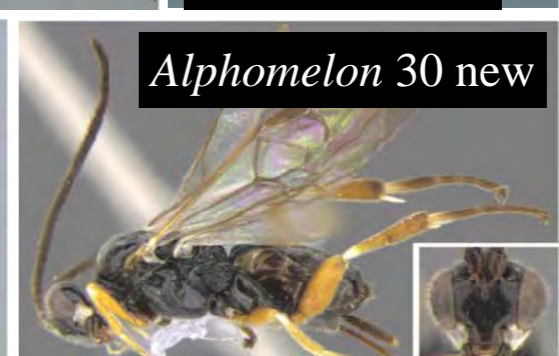
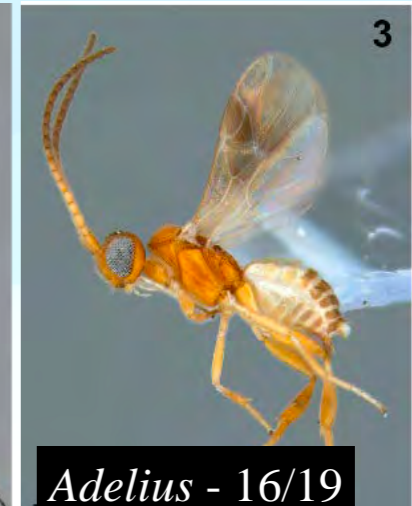


**Angelica Penteado-Dias**  
Braconidae/UFSCar

**Training-Education  
University-Collection**



# 5. Bridging the taxonomic gap and reducing the taxonomic impediment?





# 5. Bridging the taxonomic gap and reducing the taxonomic impediment?

- *Pambolus*

Not Secure — fauna.jbrj.gov.br

CATÁLOGO TAXONÔMICO DA FAUNA DO BRASIL

Search result New Query

**Animalia**

- Arthropoda Gravenhorst, 1843
- Hexapoda Latreille, 1825
- Insecta Linnaeus, 1758
- Hymenoptera Linnaeus, 1758
- Ichneumonoidea Latreille, 1802
- Braconidae Nees, 1811
- Pambolinae Marshall, 1885
- Pambolus* Haliday, 1836
- Pambolus (Phaenodus)* Förster, 1862
- Pambolus (Phaenodus) longicornis* (Enderlein, 1920)

**Information** Statistics

*Pambolus* Haliday, 1836 18327

val d

**Taxonomic Hierarchy**

Animalia → Arthropoda Gravenhorst, 1843 → Hexapoda Latreille, 1825 → Insecta Linnaeus, 1758 → Hymenoptera Linnaeus, 1758 → Ichneumonoidea Latreille, 1802 → Braconidae Nees, 1811 → Pambolinae Marshall, 1885 → *Pambolus* Haliday, 1836

**Species**

*Pambolus (Phaenodus)* Förster

**Origin**

Native

**Endemism**

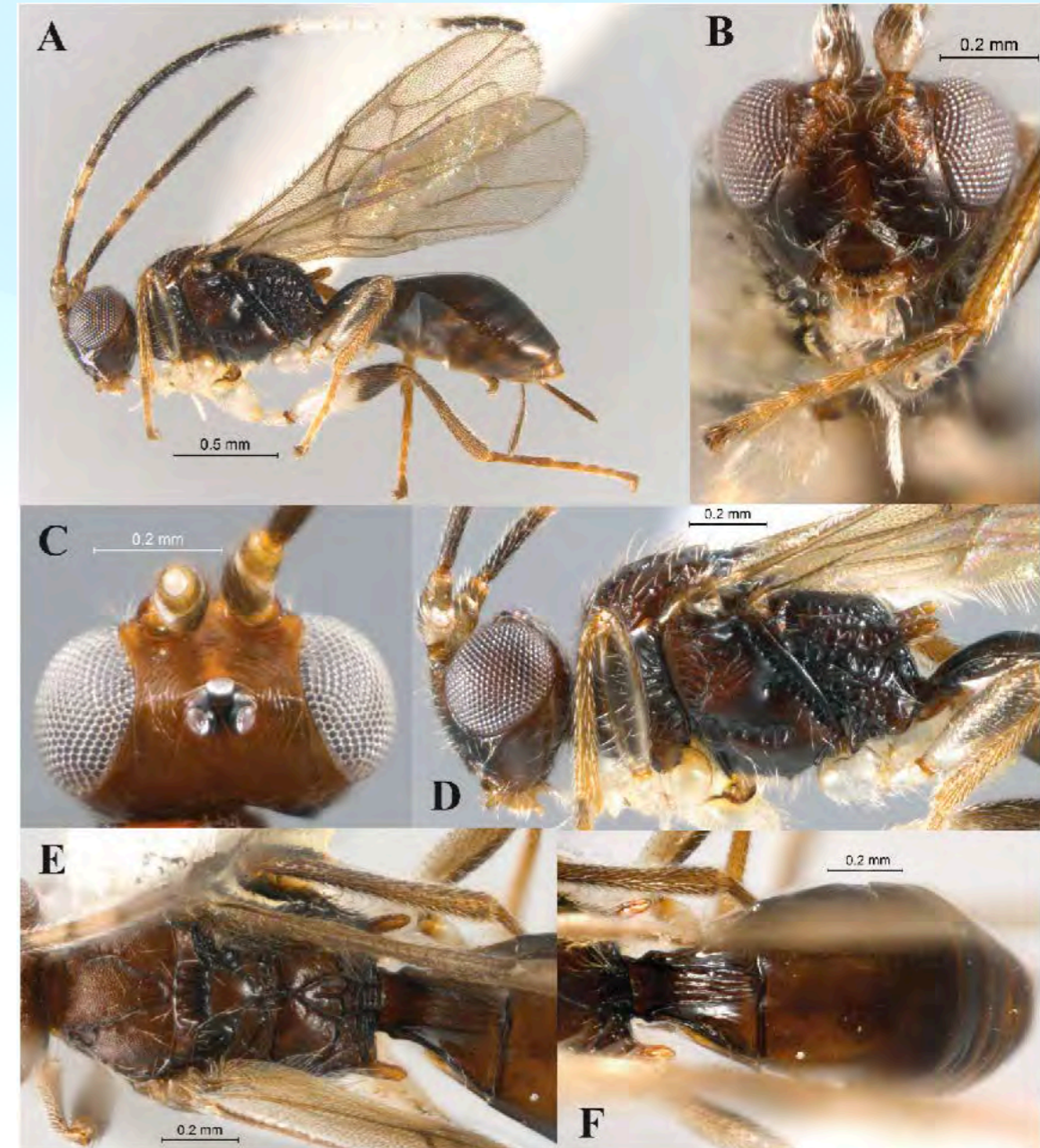
Unknown

**Environments**

Epicontinental

**Distribution**

Geographic distribution



# 5. Bridging the taxonomic gap and reducing the taxonomic impediment?

## Superficial impediment\* in the tropics:

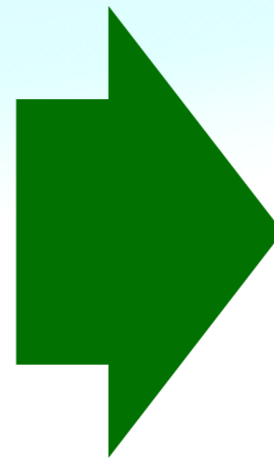
Naturalists expeditions

Many groups have very old treatments only

Superficial descriptions

Very high diversity

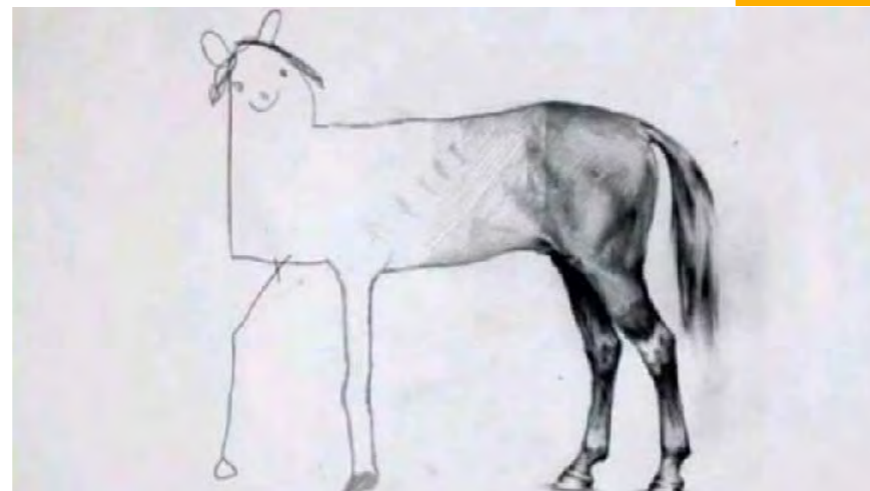
Types unavailable



**Information access: interest and work**

**Morphology: good images, 3D models and modern descriptions**

**Barcoding: museomics in old specimens**





# 5. Bridging the taxonomic gap and reducing the taxonomic impediment?

**A turbo-taxonomic study of Thai *Aleiodes* (*Aleiodes*) and *Aleiodes* (*Arcaleiodes*) (Hymenoptera: Braconidae: Rogadinae) based largely on COI barcoded specimens, with rapid descriptions of 179 new species**

**Butcher et al. 2012:** 1/4 of all names for *Aleiodes* (179 of 698) Thailand

**The Doryctinae (Braconidae) of Costa Rica: genera and species of the tribe Heterospilini**

*Costa Rica:*

**Marsh 2013:** 2/3 all *Heterospilus* names (277 of 416);

**Review of *Apanteles sensu stricto* (Hymenoptera, Braconidae, Microgastrinae) from Area de Conservación Guanacaste, northwestern Costa Rica, with keys to all described species from Mesoamerica**

**Fernandez-Triana et al. 2014:** 186 of 997 *Apanteles*;

**Arias-Penna et al. 2019:** 136 of 213 *Glyptapanteles*

**A species-level taxonomic review and host associations of *Glyptapanteles* (Hymenoptera, Braconidae, Microgastrinae) with an emphasis on 136 new reared species from Costa Rica and Ecuador**

# 6. Biological control - taxonomy

## Integration



- Integrating taxonomy with applied fields, and collaborative work in general, is a good way to reduce the taxonomic impediment
- Search for common goals: results improve, appreciation improves, access to wider funding
- One may think that natural enemies are pretty much known for biological control already, but



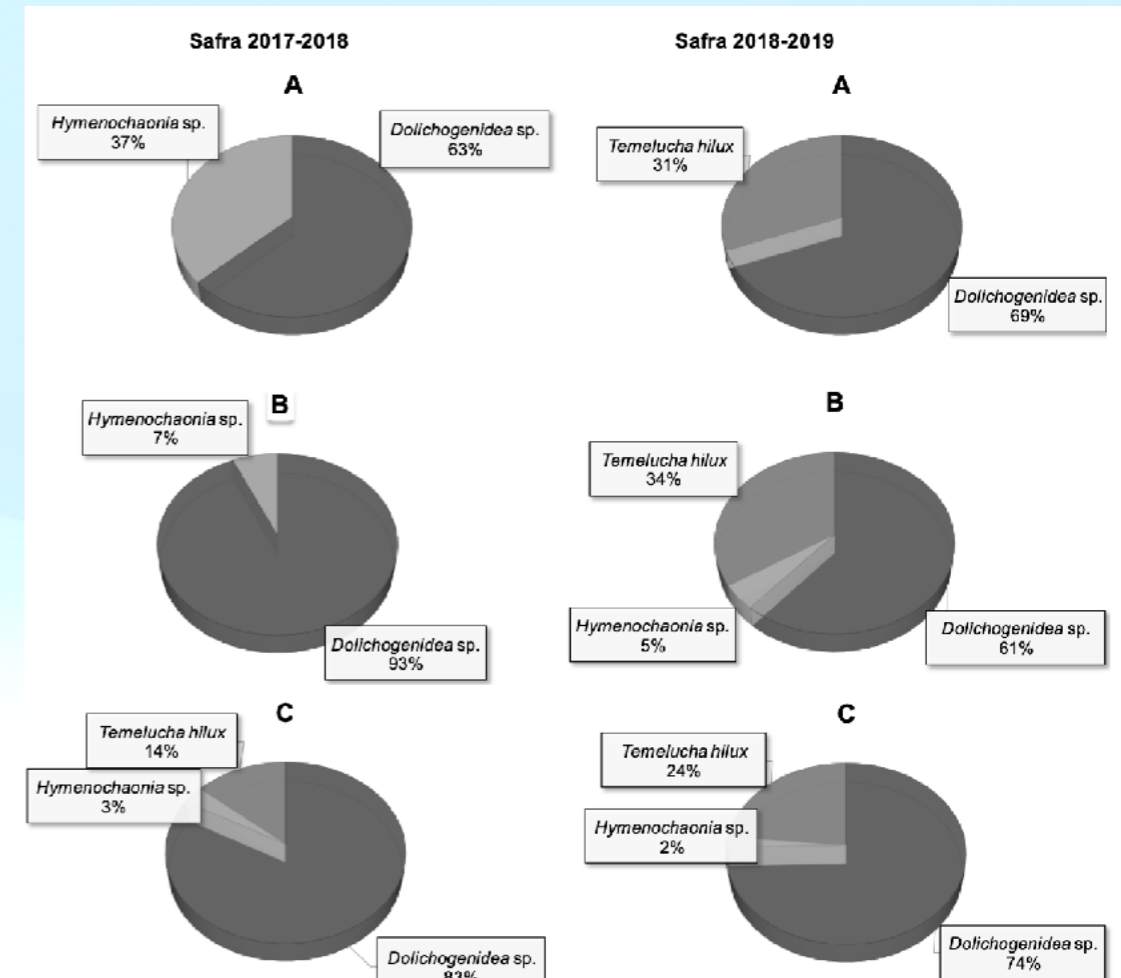


# 6. Biological control - taxonomy

## Integration



*Gymnandrosoma aurantianum* - citrus



Scheunemann T, 2022



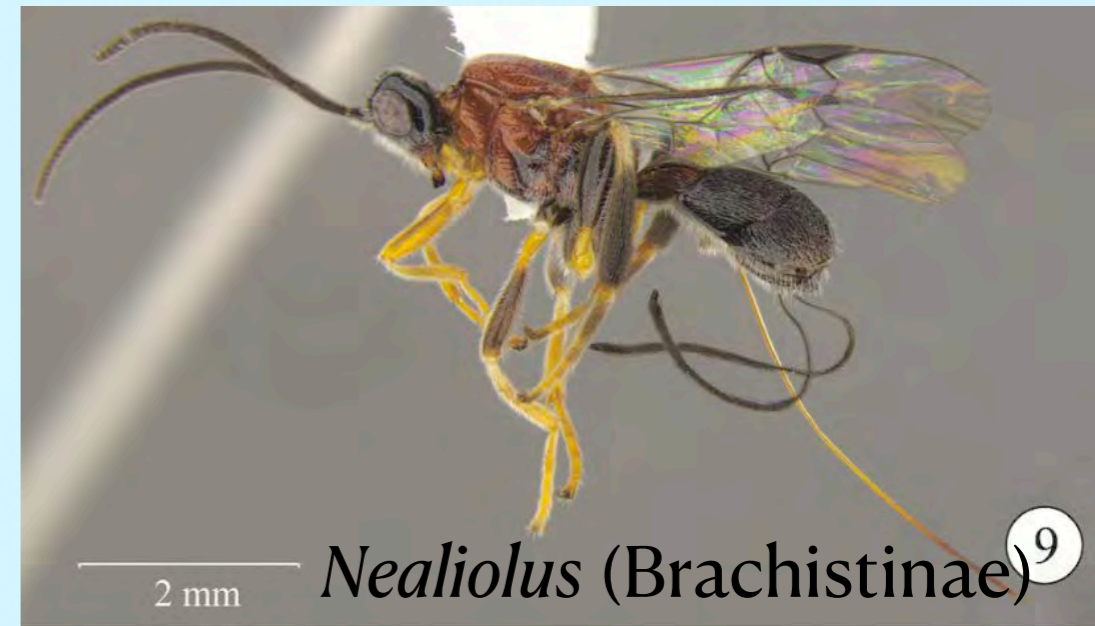
*Dolichogenidea* sp.



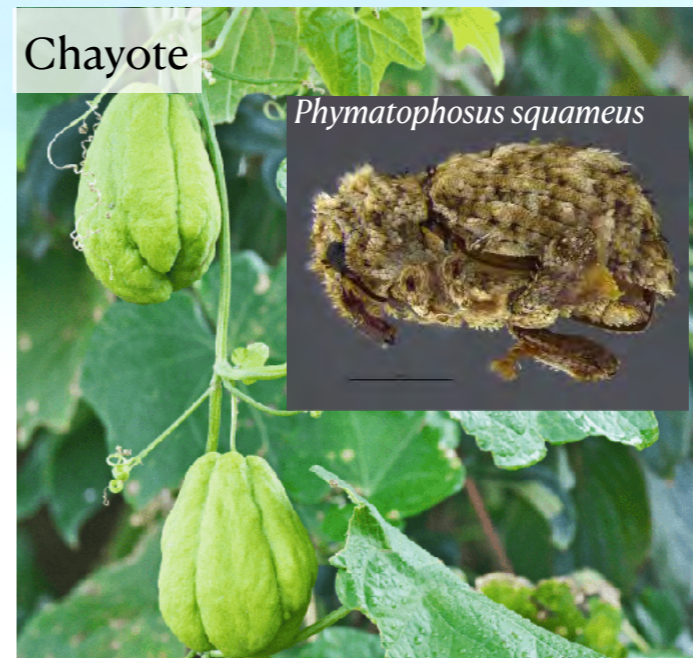
*Palpita forficera* - olive



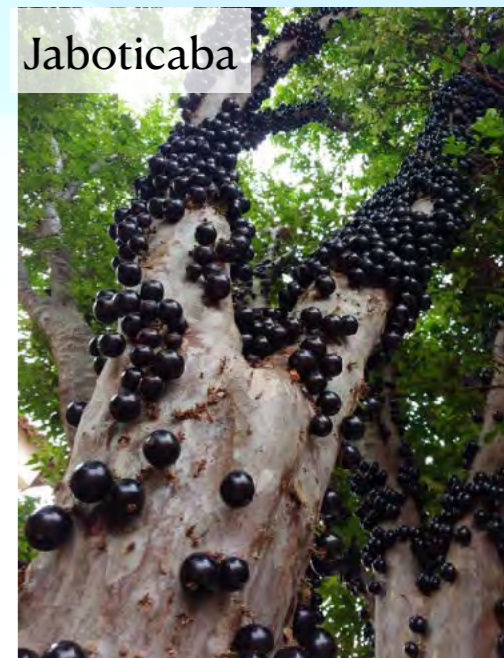
# 6. Biological control Integration



*Nealiolus* (Brachistinae)<sup>9</sup>



*Phymatophosus squameus*



*Plinia cauliflora* (Myrtaceae)



*Apanteles* (Microgastrinae)



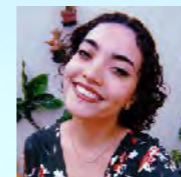
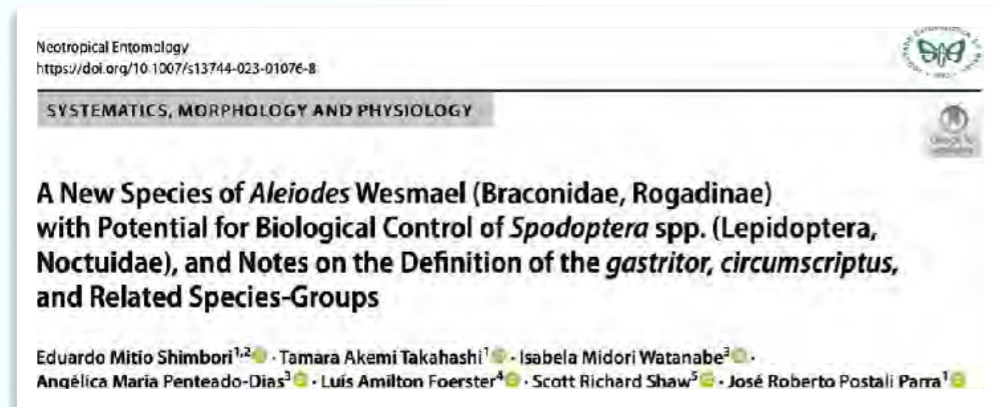
Thanks  
*Apanteles mayochinchi*!!



# 6. Biological control - taxonomy

## Integration

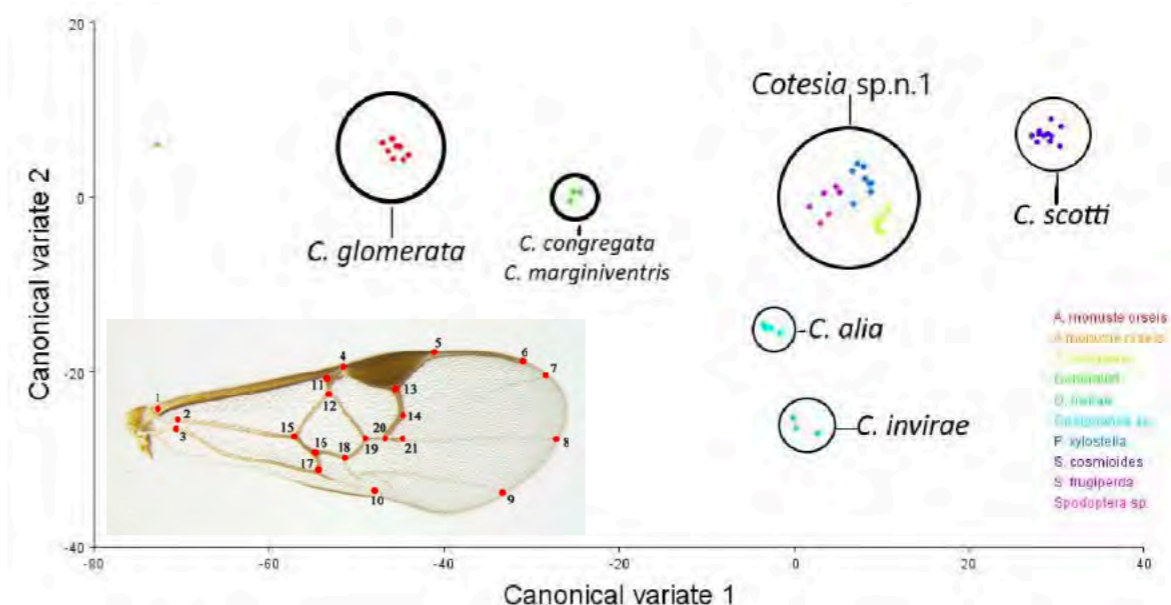
- cryptic species complexes and gamma-level taxonomy



Sarah Garcia



Integrative taxonomy of *Cotesia* Cameron (Hymenoptera: Braconidae) and its potential for biological control in the Neotropical region (2022)





# 6. Biological control - taxonomy

## Integration

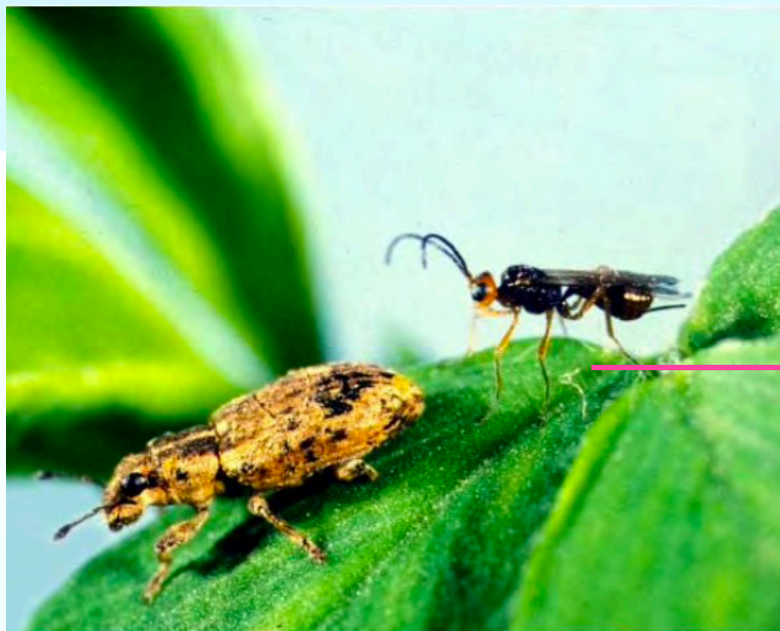
- cryptic species complexes and gamma level taxonomy

NOTES ON THE VARIABILITY OF *MICROCTONUS AETHIOPOIDES* LOAN  
(HYMENOPTERA : BRACONIDAE : EUPHORINAE)

Jean-Paul Aeschlimann

335 avenue Paul Parguel, 34100 Montpellier, France

Moroccan strain



*Sitona discoideus*  
(lucerne weevil)

Irish strain



*Sitona obsoletus*  
(clover root weevil)

intraspecific variation

hybridization/  
speciation

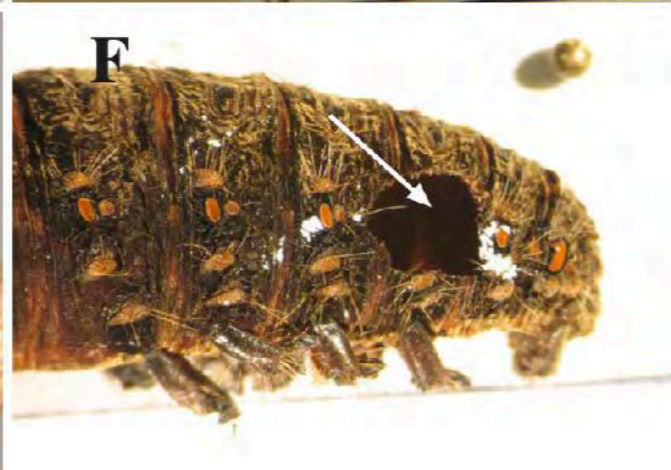
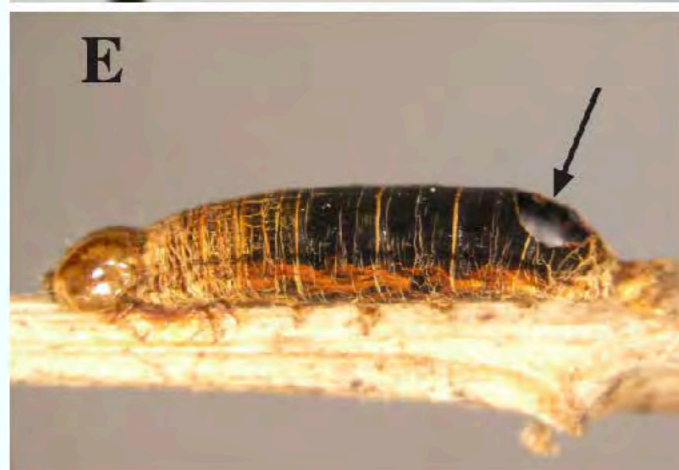
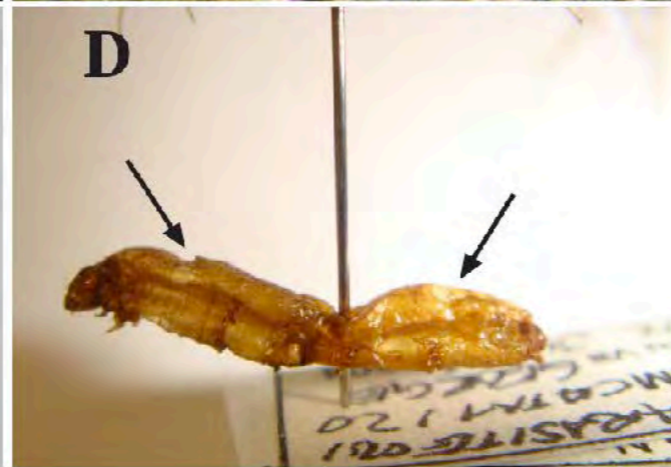
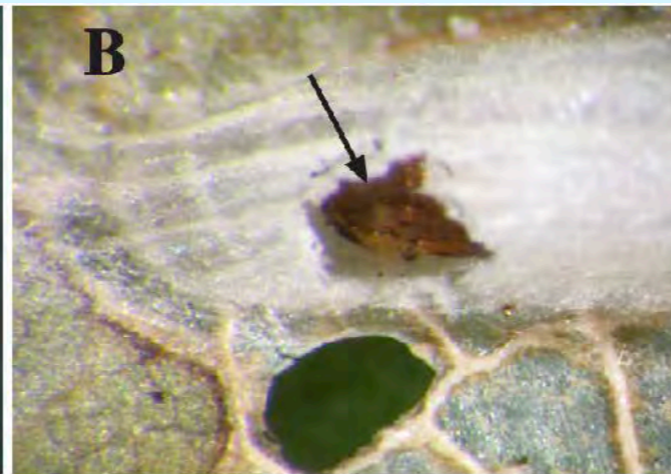
endosymbionts and  
viral endogenization

non-target impacts



# 7. Current projects

## Genomics tools - Rogadinae



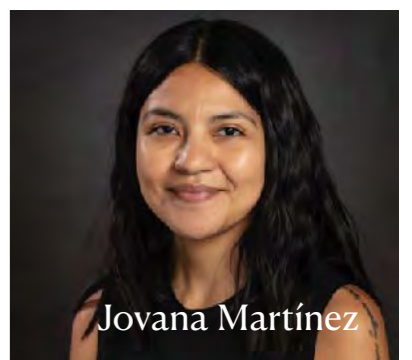


# 7. Current projects

## Genomics tools - Rogadinae



## Phylogenetic systematics and evolution of *Aleiodes* (Braconidae) based on UCEs



**Worldwide sampling (~300 spp)**

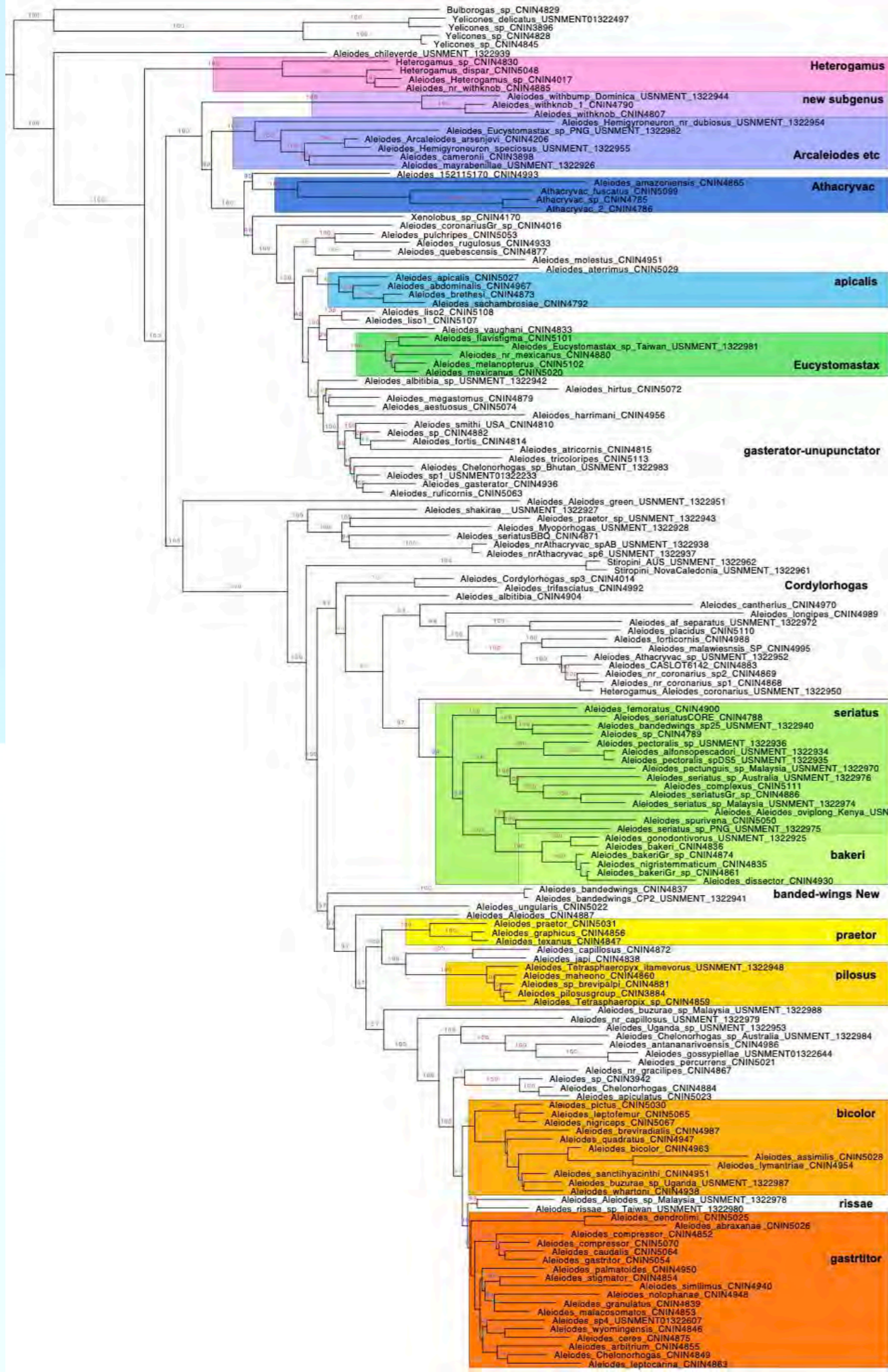
**Whole-genome shotgun sequencing - UCEs**

**Hypothesis: subgenera and species group (>30)**

**Revise systematics at subgeneric level**

**Origin and diversification times and ancestral states**





Geometridae

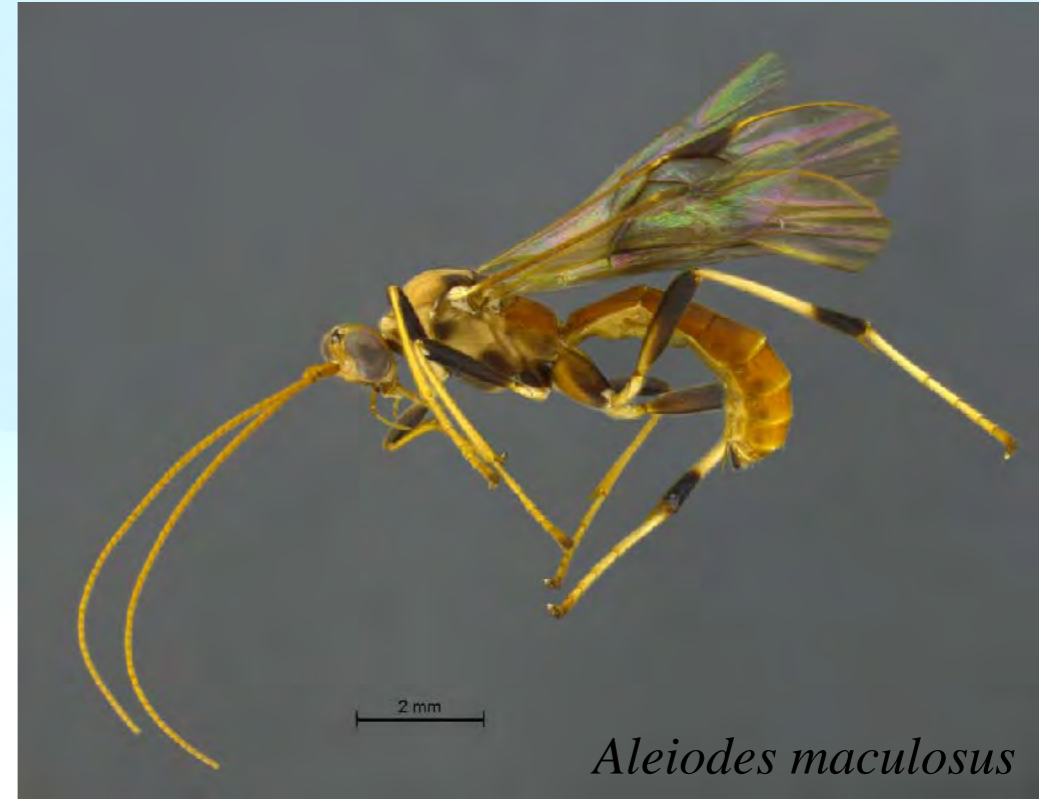
Noctuidae

Erebidae

Sphingidae

Geometridae

Noctuidae



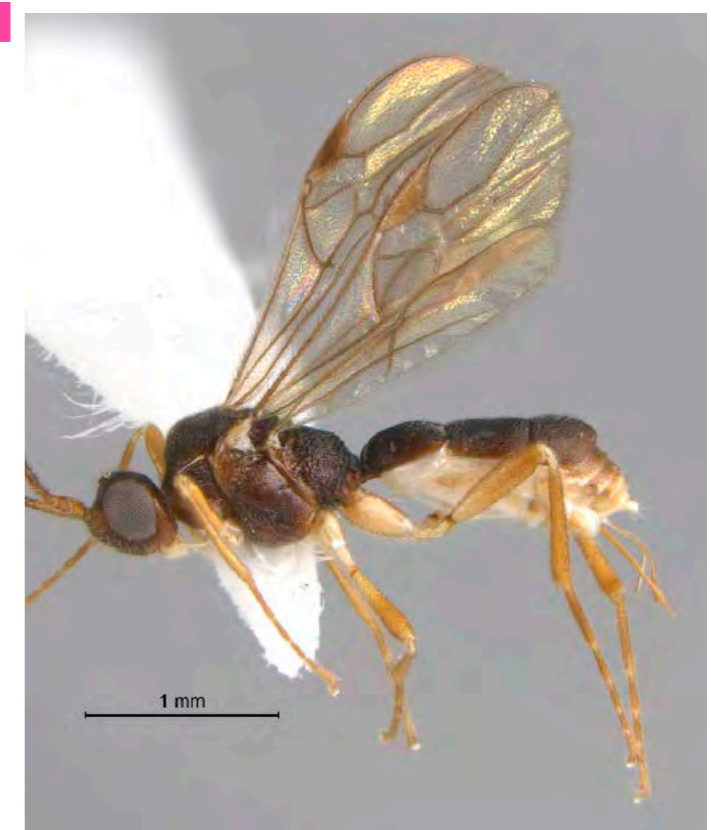
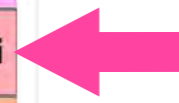
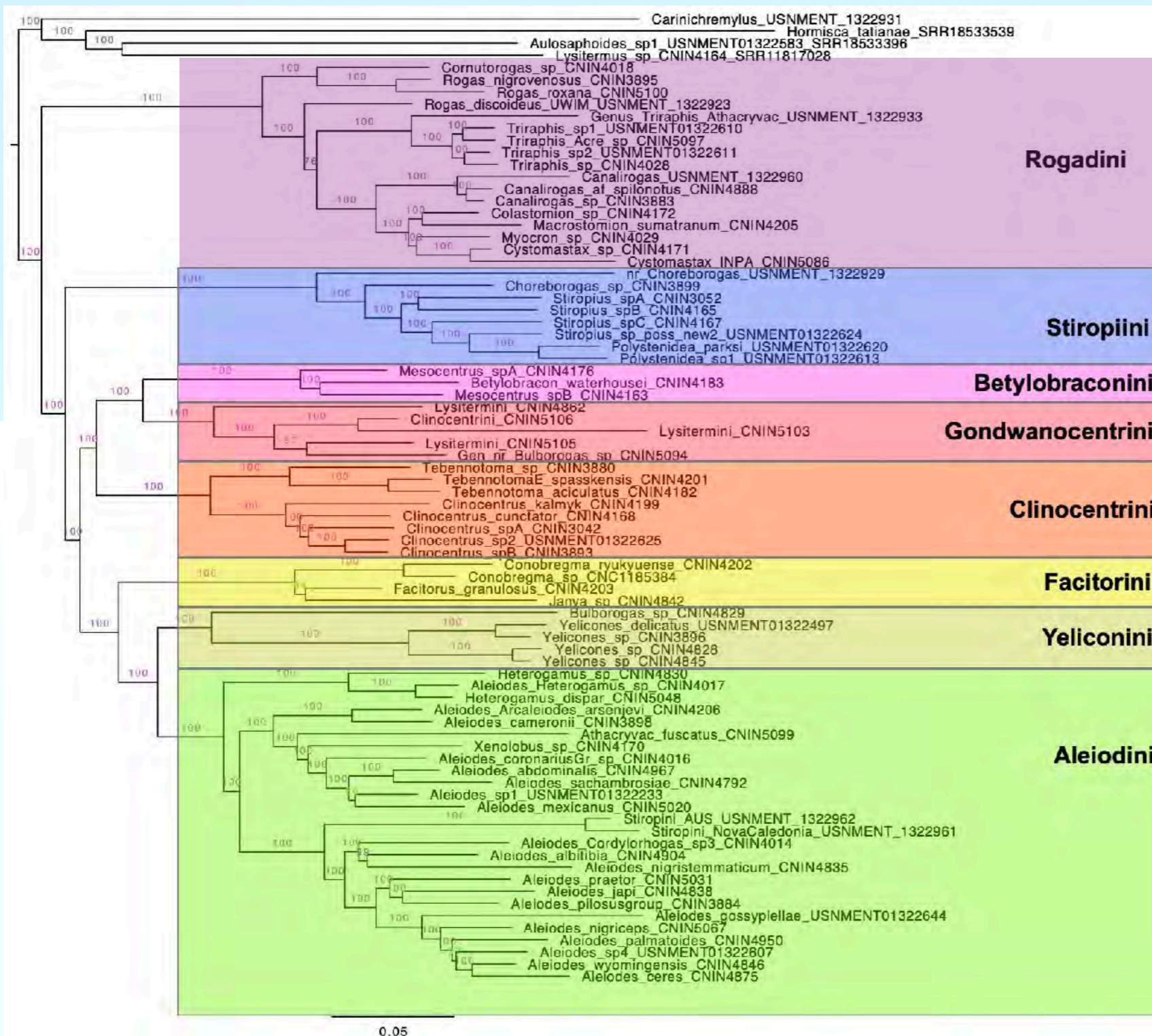
*Aleiodes maculosus*

How to taxonomically treat a genus with worldwide distribution and probably thousands of species?



# 7. Current projects

## Genomics tools - Rogadinae

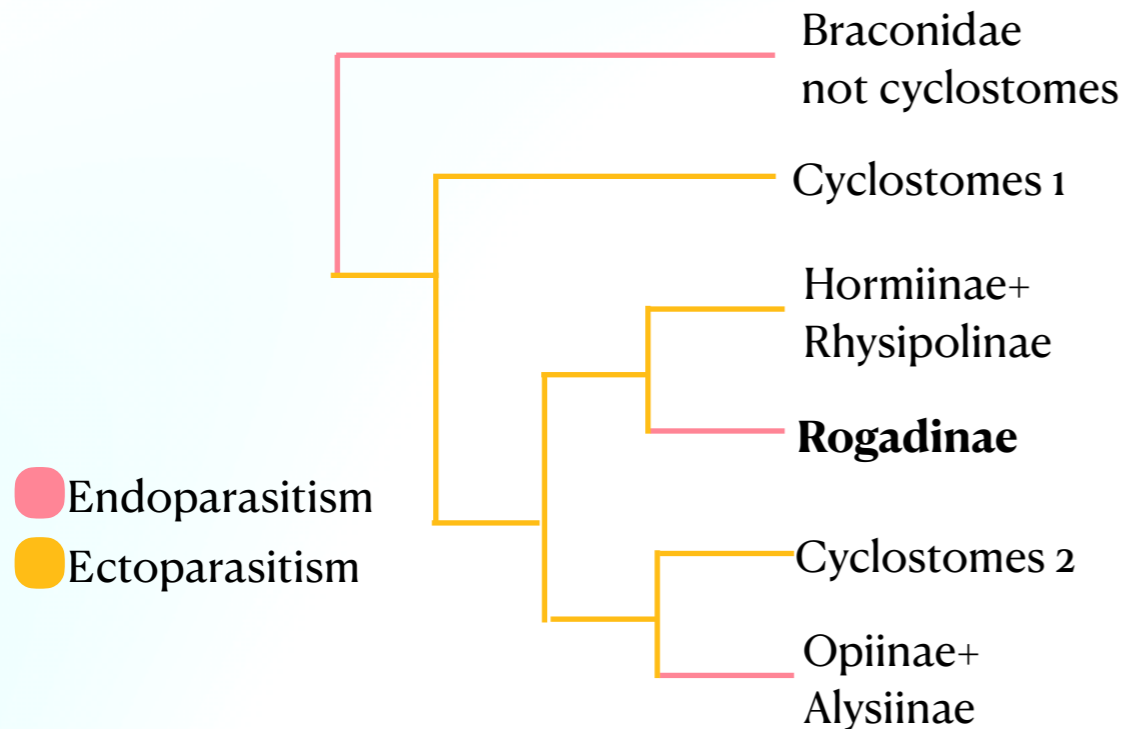
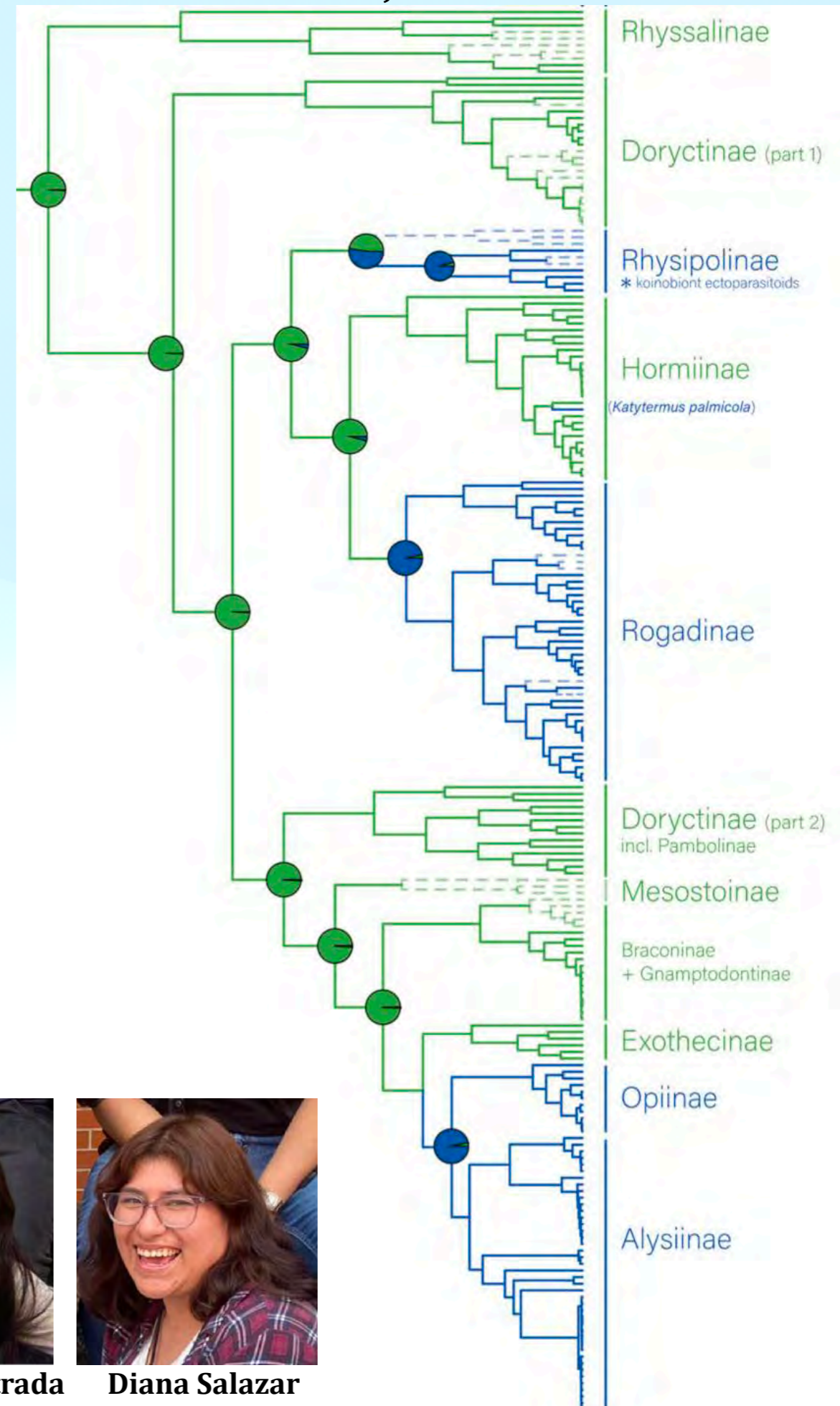




# 7. Current projects

## Genomics tools - Rogadinae

**Evolution of Rogadinae using genomic tools: role of endogenized viral elements and toxin-resistance genes in the transition of life modes and host exploitation.**



Alejandra Estrada



Diana Salazar

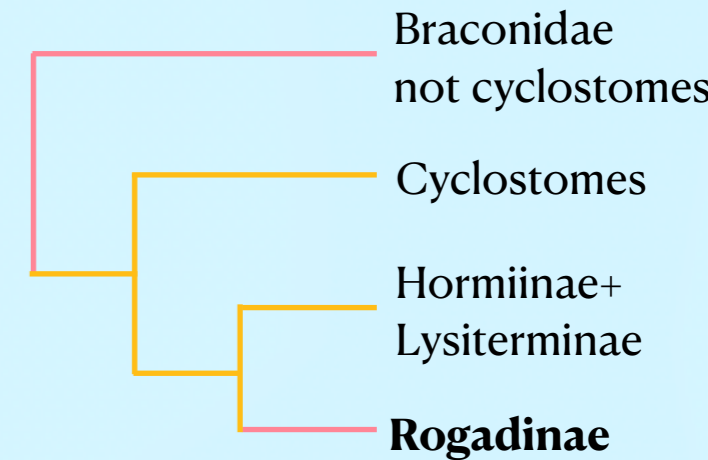


# 7. Current projects

## Genomics tools - Rogadinae

**Evolution of Rogadinae using genomic tools: role of endogenized viral elements and toxin-resistance genes in the transition of life modes and host exploitation.**

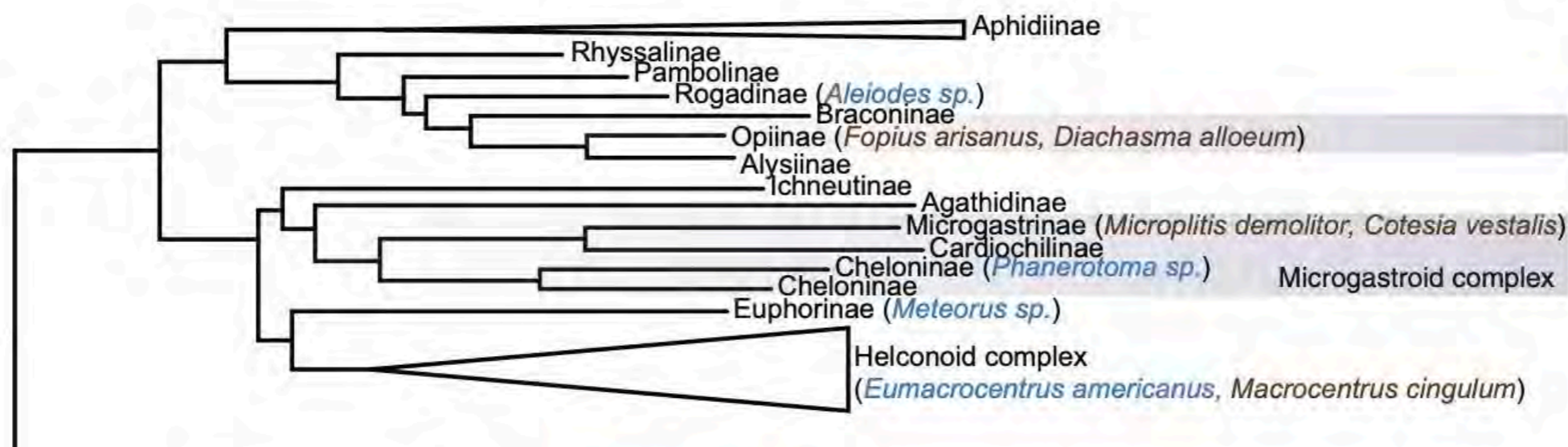
- Endoparasitism
- Ectoparasitism



**Endoparasitoid lifestyle promotes endogenization and domestication of dsDNA viruses**

Benjamin Guinet<sup>1\*</sup>, David Lepetit<sup>1</sup>, Sylvain Charlat<sup>1</sup>, Peter N Buhl<sup>2</sup>, David G Notton<sup>3</sup>, Astrid Cruaud<sup>4</sup>, Jean-Yves Rasplus<sup>4</sup>, Julia Stigenberg<sup>5</sup>, Damien M de Vienne<sup>1</sup>, Bastien Boussau<sup>1</sup>, Julien Varaldi<sup>1\*</sup>

Burke et al. 2021 *Genome Biol. Evol.* 13(7)



**EVE type (distribution)**

- *Alphanudivirus* EVE (select *Fopius* species)
- *Bracovirus* derived from *Betanudivirus* (Microgastroid complex)



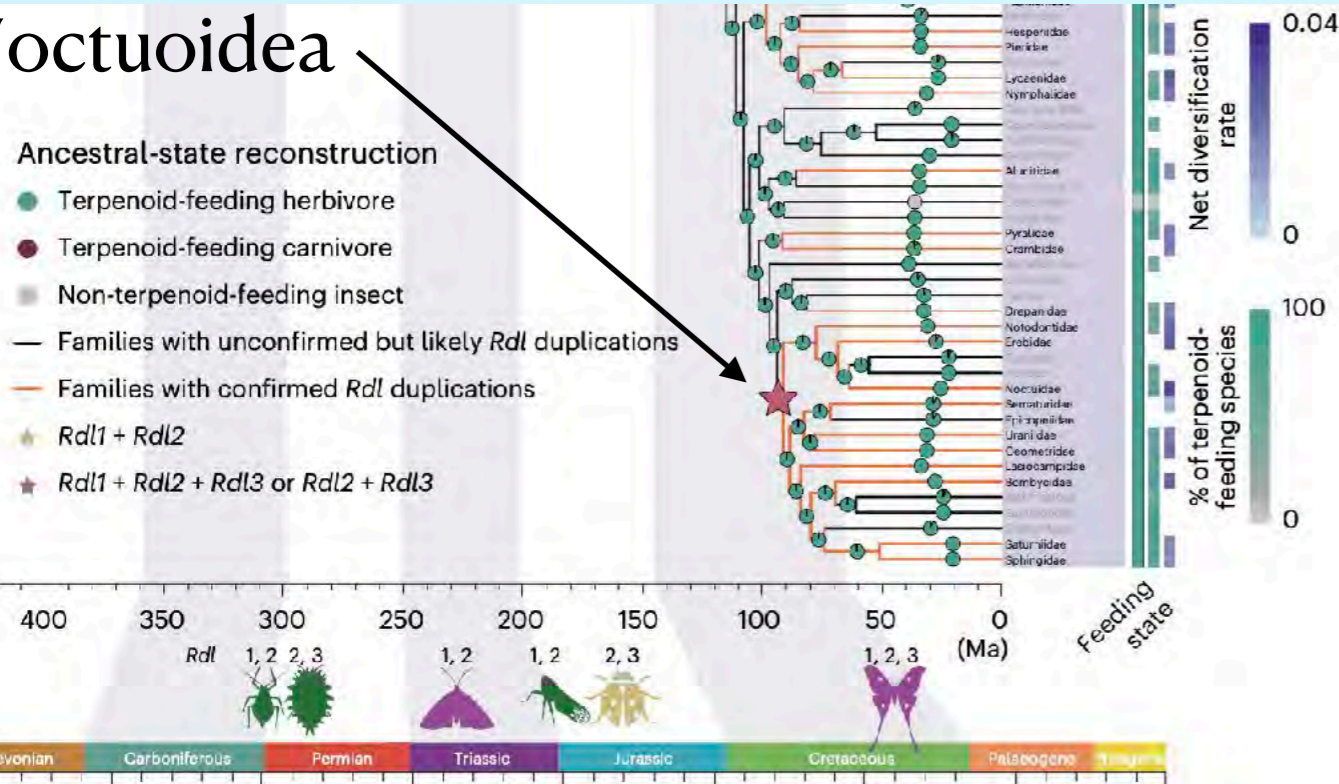
# 7. Current projects

## Genomics tools - Rogadinae

### Noctuoidea

#### Ancestral-state reconstruction

- Terpenoid-feeding herbivore
- Terpenoid-feeding carnivore
- Non-terpenoid-feeding insect
- Families with unconfirmed but likely *Rdl* duplications
- Families with confirmed *Rdl* duplications
- ★ *Rdl1* + *Rdl2*
- ★ *Rdl1* + *Rdl2* + *Rdl3* or *Rdl2* + *Rdl3*



### nature ecology & evolution

## Convergent resistance to GABA receptor neurotoxins through plant-insect coevolution

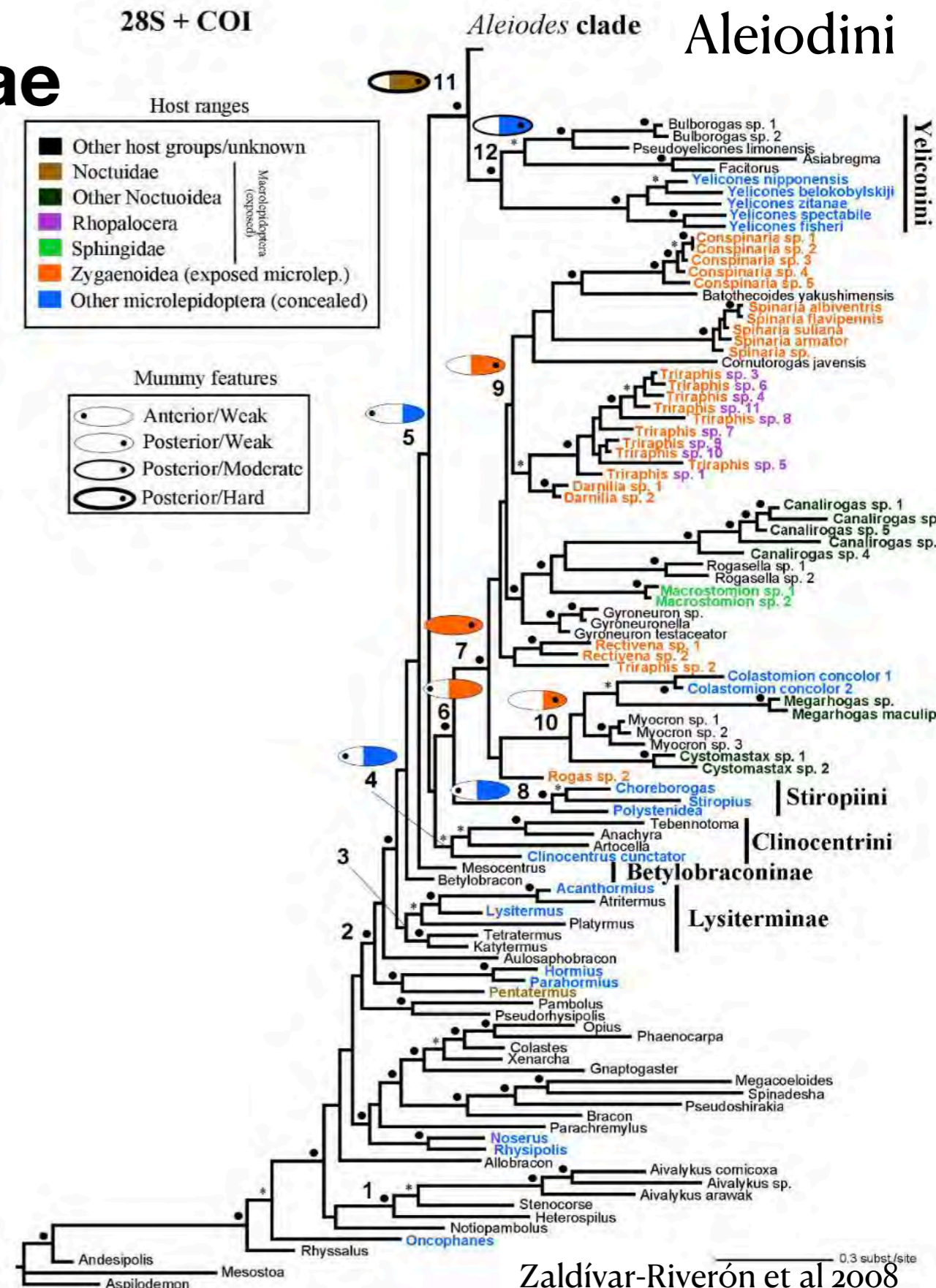
Received: 7 February 2023

Lei Guo<sup>1,5</sup>, Xiaomu Qiao<sup>2,5</sup>, Diler Haji<sup>3</sup>, Tianhao Zhou<sup>1</sup>, Zhihan Liu<sup>1</sup>, Noah K. Whiteman<sup>3,4</sup> & Jia Huang<sup>1</sup>

Accepted: 22 June 2023

Published online: 17 July 2023

<https://doi.org/10.1038/s41559-023-02127-4>



Zaldívar-Riverón et al 2008



# 7. Current projects

## Genomics tools - Rogadinae

- IPM context:
  - Resistance to toxins = resistance to pesticides?
  - Endogenous viral elements and virus biocontrol — interactions in Lepidoptera as preferred targets?



# 7. Current projects

## Genomics tools and barcoding

- Essential tools for bridging gaps
- DNA from museum specimens
- Generating barcodes (legacy genes)
- micro -> macroevolutionary
- Host-parasitoid associations



JHR 97: 29–42 (2024)  
doi: 10.3897/jhr.97.113231  
<https://jhr.pensoft.net>

RESEARCH ARTICLE



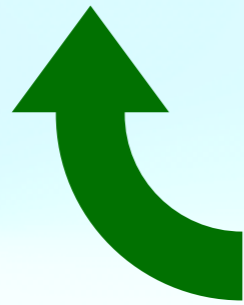
JOURNAL OF  
**Hymenoptera**  
RESEARCH  
The International Society of Hymenoptera Research

**High hymenopteran parasitoid infestation rates in Czech populations of the *Euphydryas aurinia* butterfly inferred using a new molecular marker**

Hana Konvičková<sup>1,2</sup>, Václav John<sup>1,2,3</sup>, Martin Konvička<sup>1,2</sup>, Michal Rindoš<sup>1,2</sup>, Jan Hrček<sup>1,2</sup>



Species discovery/  
identification



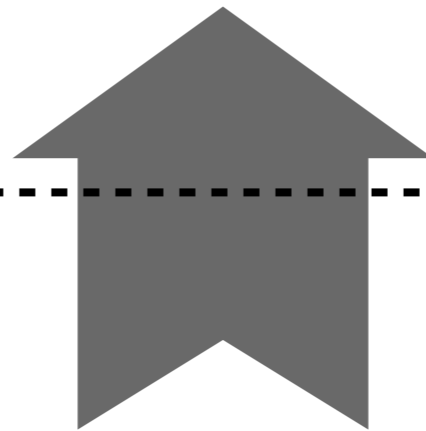
Host-Parasitoid  
associations



Insect monitoring



Barcode  
Reference databases



Taxonomic Impediment

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Legacy descriptions



**Merci  
beaucoup !**



**SHIMBORI@GMAIL.COM**