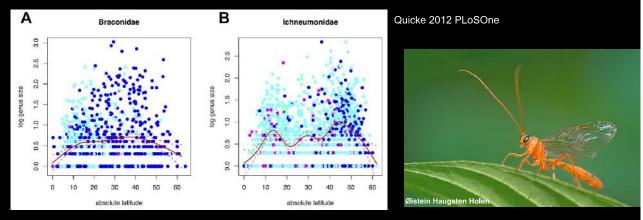


Reversed Latitudinal Biodiversity Gradients or sampling artifact?

Majority of ichneumonid and braconid species described from temperate regions



Discovery of over 200 undescribed orthocentrine species from Ecuador Massive under-description of tropical parasitoid faunas







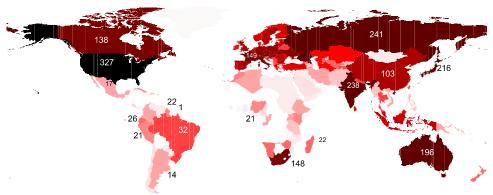
Proc. R. Soc. B doi:10.1098/rspb.2012.1664 Published online

Unprecedented ichneumonid parasitoid wasp diversity in tropical forests

Anu Veijalainen^{1,2,*}, Niklas Wahlberg¹, Gavin R. Broad³, Terry L. Erwin⁴, John T. Longino⁵ and Ilari E. Sääksjärvi¹



- Higher species richness in temperate areas
- Research on gracillariid systematics mainly in Europe, Asia, and North America
- Only one species recorded from French Guiana





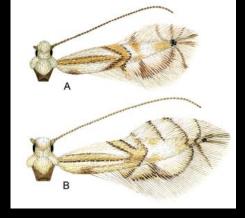
Low diversity in Neotropics: artefact of undersampling?

- 184 gracillariid species recorded in the Neotropics
- 1996 1998, preliminary survey by Davis & Wagner (ALAS) Project
- · La Selva Biological Station a lowland rain forest site in northwestern Costa Rica
- Based on mine morphology and host plant information: 200 species estimated

• Most specious genus *Phyllocnistis* with an estimated 60 species, (only a single previously

known species)

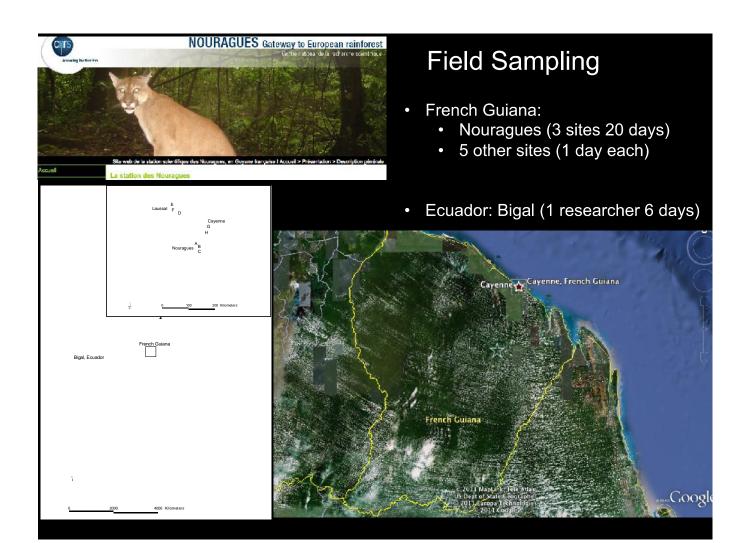




- <u>Hypothesis</u>: Low diversity of Neotropical gracillariids is due to insufficient sampling and strong description deficit
- <u>Problem</u>: how to rapidly assess a local fauna when it may be partly or largely unknown









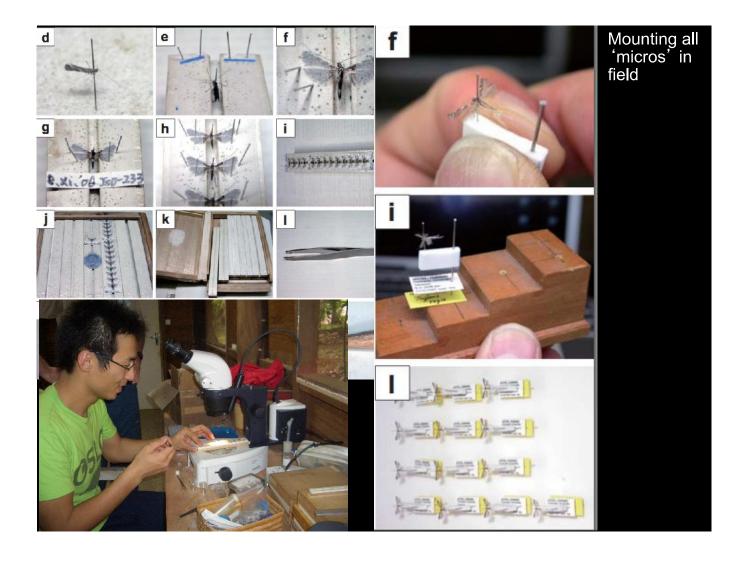


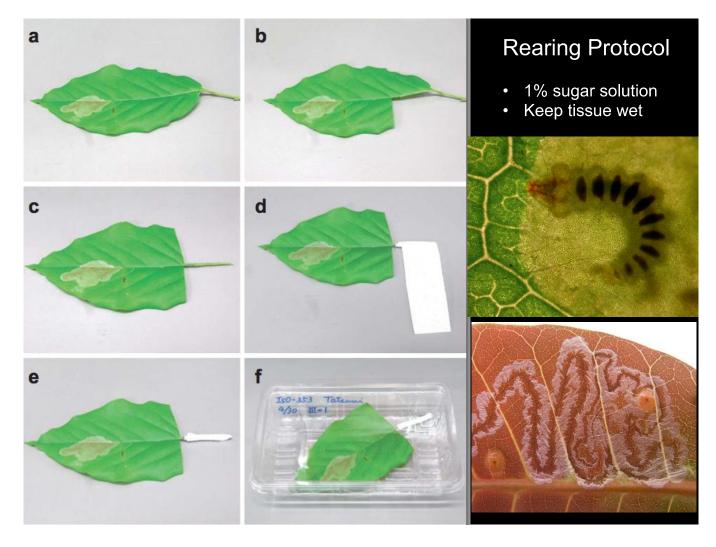


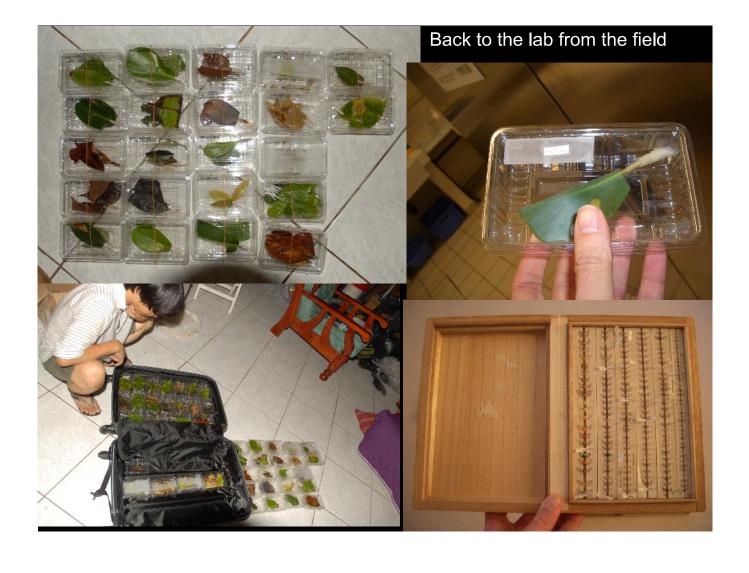


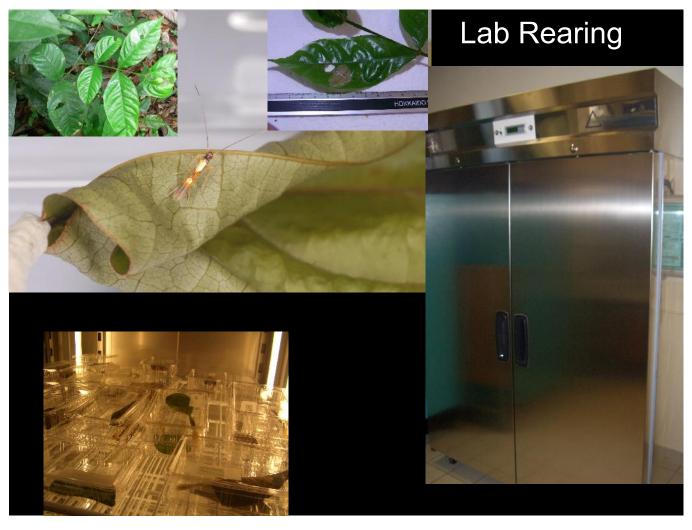














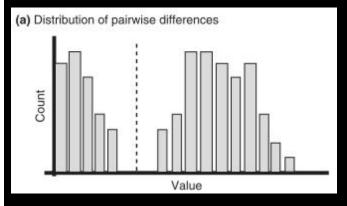


- One hind leg per specimen (adults)
- Barcoding at Guelph, Canada.
- 485 barcodes (94% success per plate):
 - 445 French Guiana
 - 40 Ecuador
- 372 adults, 104 larvae and nine pupae
- •Nearly all barcodes novel to BOLD: only three Eucosmophora had close matches

Species delimitation

Automatic Barcode Gap Discovery





Intraspecific divergence (organisms belonging to the same species) < interspecific divergence (organisms from different species)

MOLECULAR ECOLOGY

Molecular Ecology (2012) 21, 1864-1877

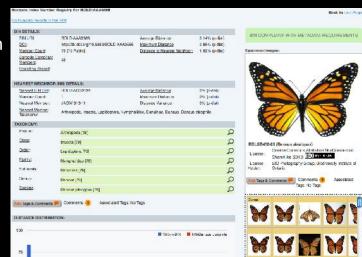
doi: 10.1111/j.1365-294X.2011.0

ABGD, Automatic Barcode Gap Discovery for primary species delimitation

N. PUILLANDRE, * A. LAMBERT, † S. BROUILLET †§ and G. ACHAZ †§

Species delimitation: Refined Single Linkage algorithm

- Rapid automated Operational Taxonomic Units (OTUs) recognition system
- Each OTU is assigned a Barcode Index Number (BIN)
- Persistent registry : DOI number
- 274 000 BIN web pages available: http://www.boldsystems.org/bin



OPEN & ACCESS Freely available online

PLOS ONE

A DNA-Based Registry for All Animal Species: The Barcode Index Number (BIN) System

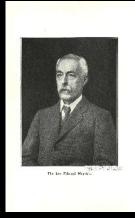
Sujeevan Ratnasingham1*, Paul D. N. Hebert1,2

1 Biodiversity Institute of Ontario, University of Guelph, Guelph, Ontario, Canada, 2 Department of Integrative Biology, University of Guelph, Guelph, Ontario, Canada

Morphological Analysis



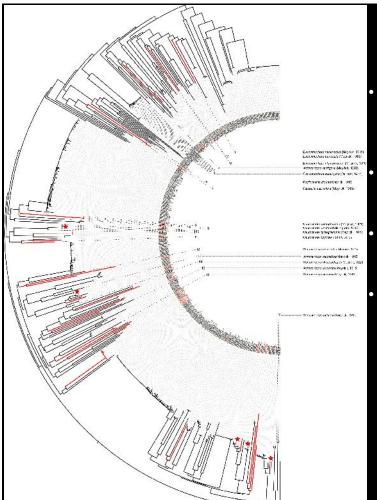




- Identification of species based on wing patterns (monographic revisions and taxonomic literature): morphospecies
- Meyrick's Types at BMNH London
- 64% of 184 Neotropical species described by Meyrick before 1930's
- 150 species examined from types/literature.

VII. Descriptions of South American Micro-Lepidoptera. By E. Meyrick, B.A., F.R.S.

[Read May 5th, 1915.]



Species delimitation:

ABGD: 136BINs: 151

Identification using morphology:

• 17 tentative species names

Nouragues: 108 species 64 (59.3%) are represented by singletons

 85% of the species collected as adults do not fit any of the 150 species we examined from types/ literature.

Lees et al 2013 Molecular Ecology Resources

Spatial and Temporal Turnover

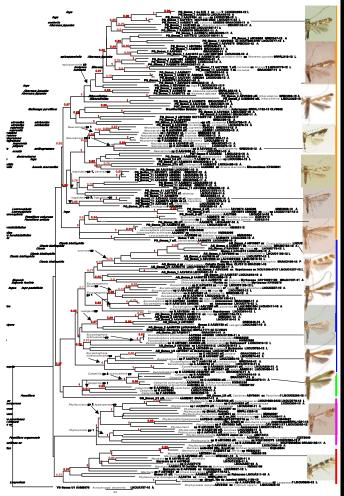
Table 1 Local species richness for the nine sampling sites

| 170 175 | | |
|--------------------|---------------------------------------|---|
| Number of barcodes | Number of BINS | Number of ABGD |
| 132 | 72 | 67 |
| 65 | 20 | 20 |
| 166 | 41 | 39 |
| 21 | 10 | 9 |
| 6 | 5 | 5 |
| 2 | 2 | 2 |
| 30 | 12 | 12 |
| 24 | 10 | 11 |
| 40 | 27 | 27 |
| | of barcodes 132 65 166 21 6 2 30 24 | of barcodes of BINS 132 72 65 20 166 41 21 10 6 5 2 2 30 12 24 10 |

- Groupings by the two delimitation methods were remarkably congruent
- Five BINs occur in both Ecuador and French Guiana: vast distribution ranges

- High temporal turnover:
 - Out of 72 BINs at Nouragues camp (site A) only eight BINs (11.1%) in common between January and September

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Phylogenetic signal

Parectopa group

Maximum likelihood analysis segregated our specimens among five of the six major lineages of Gracillariidae as defined by Kawahara et al. (2011) based on 21 nuclear protein-coding genes

Acrocercops group

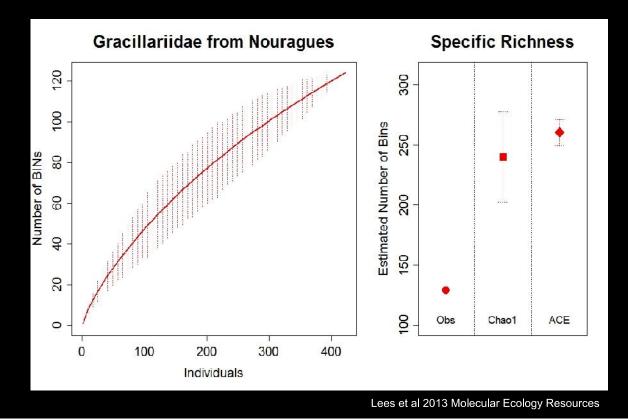
Gracillariinae

Phyllocnistinae

Lithocolletinae + Marmara

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Species richness estimates for Nouragues: 240 species (Chao1) - 260 species (ACE). As many as in the whole of Europe



Conclusions

- High alpha diversity: 108 species at one site (Nouragues).
- Frequency of singletons (59.3%) at Nouragues twice as high as average for tropics (32%):
 - Very incomplete Inventory: 240–260 species at Nouragues are likely underestimates
- we found as many species as described on the whole continent within the past 150 years.
- DNA barcoding allows researchers to overcome the taxonomic impediment: rapid biodiversity assessments in poorly documented regions
- The reverse LBG found in Gracillariidae is an artefact of insufficient sampling and underdescription

Perspectives

- Formal taxonomic treatment could take years using traditional methods (i.e. genitalia dissections, wing and SEM preparations).
 - Species description pipelines
- Improve sampling effort: Too many singletons
- 16 000 tree species in Amazonia: 16 000 Gracillariid species?

Acknowledgements



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Rodolphe Rougerie (INRA)





Atsushi Kawakita (Univ Kyoto)





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Everyone at the GUELPH lab