

Biodiversity and evolution in Neotropical mimetic butterflies

Maël Doré

PhD Thesis Defense

June 1st 2023



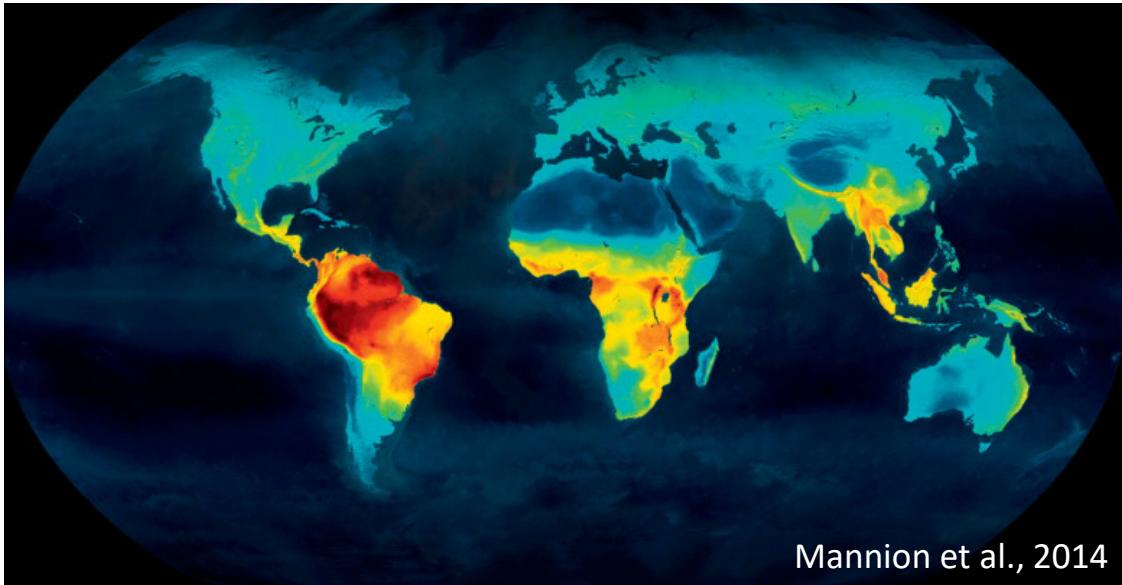
Credit photos: C. Jiggins



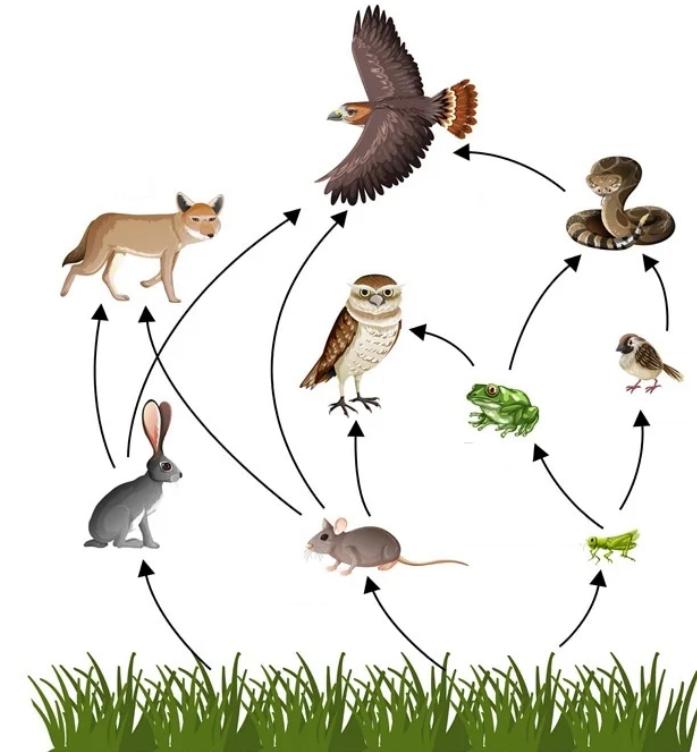
Context

Biodiversity

Biological units



Interactions



Credit: Shutterstock.com

Context

Credits: Thomas Kline



- Mutualistic interactions:** +
- Cooperative hunting
 - Plant facilitation
 - Müllerian mimicry

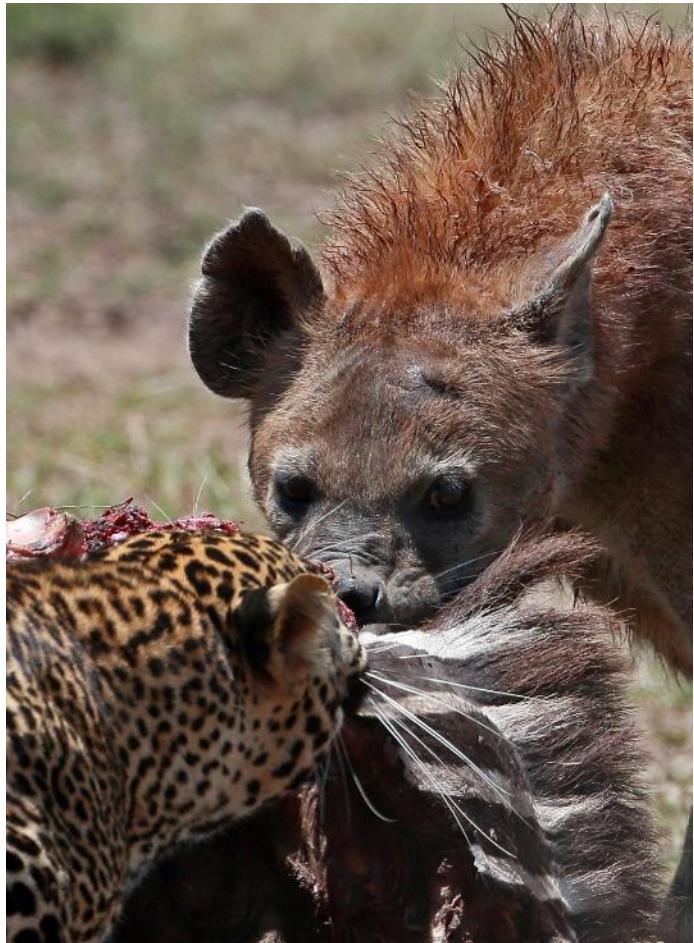


Credits: Amanda R. Liczner



Credits: Jason L. Brown

Context



Credit: Caters News Agency



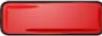
Credit: Campillo Rafael



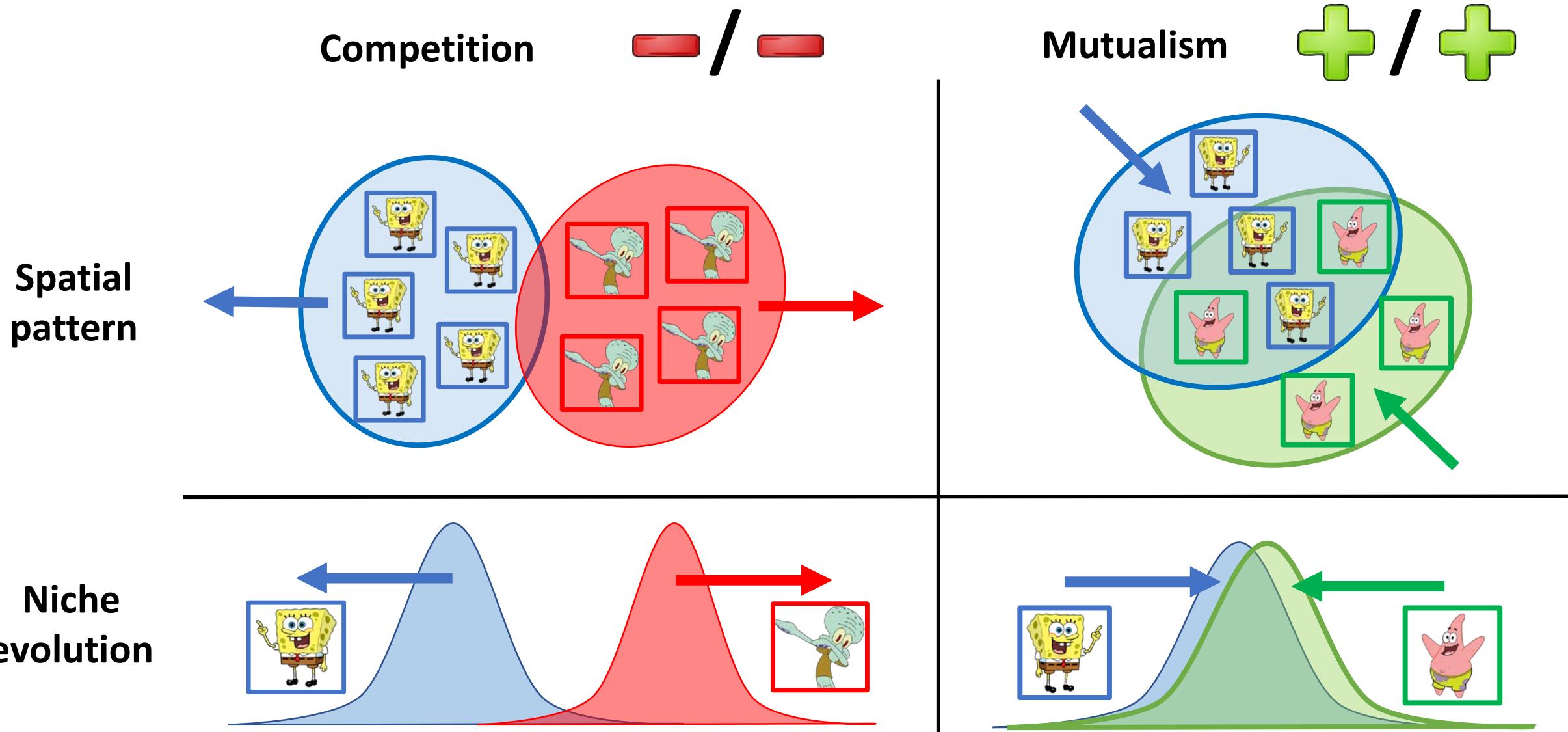
Credits: Anytka Olkova & Larry Myers

Exploitative competition:

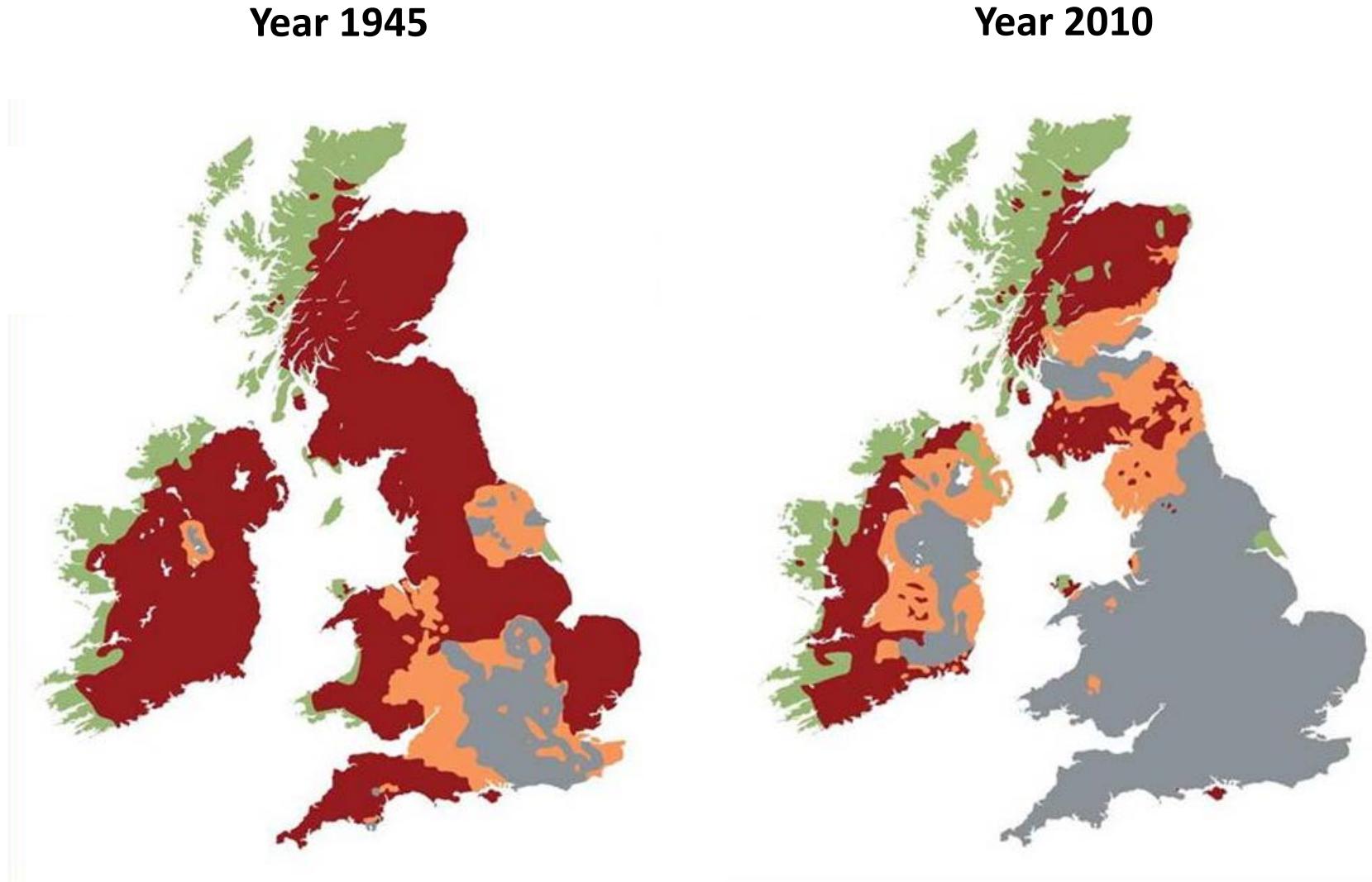
- for local resources
- for habitats
- for space



Context



Context



Source: WTSWW

Question & Hypotheses

How **mutualistic interactions** affect the **structure** and **evolution** of biodiversity at the **macroecological scale**?

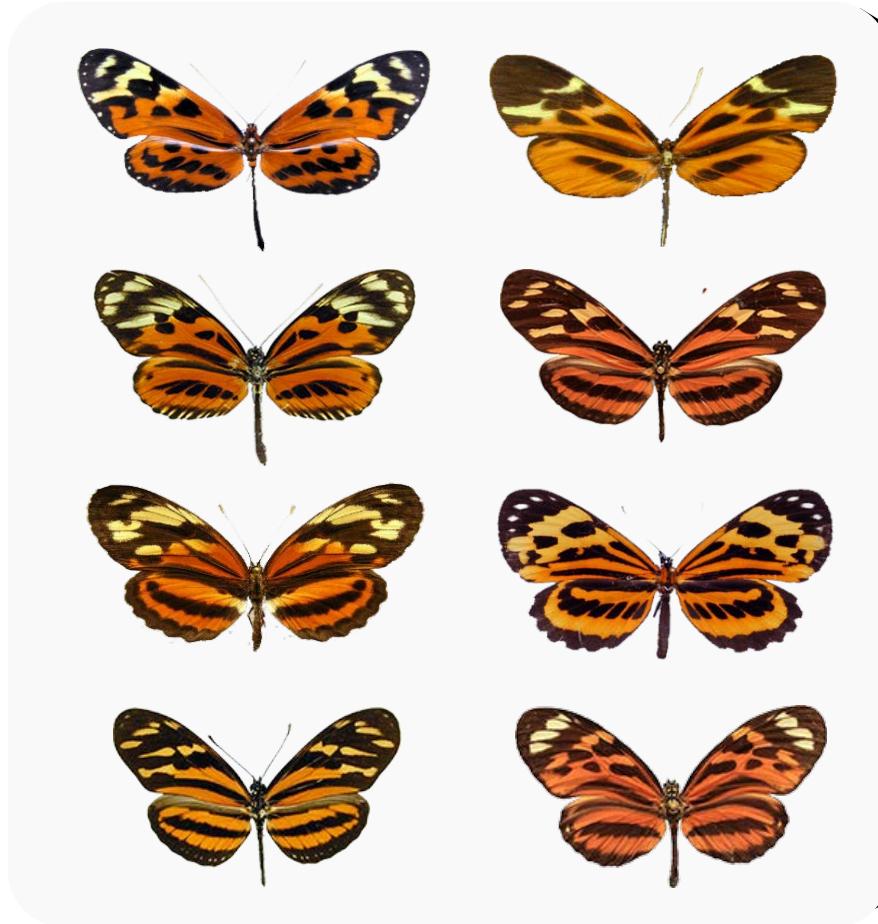
**Spatial
pattern**

Promote the large-scale **cooccurrence** of mutualistic species

**Niche
evolution**

Drive the **convergence** of the niche of mutualistic species

Study system: Müllerian mimicry

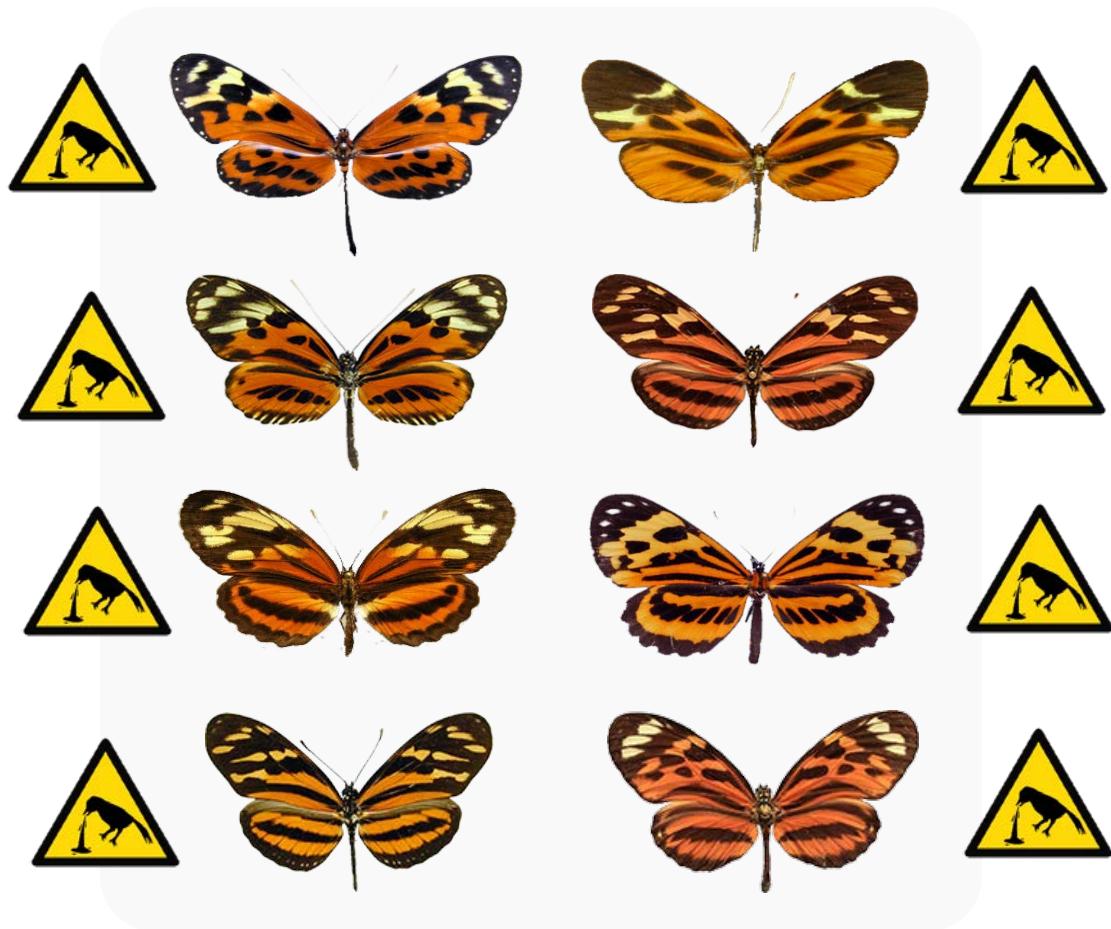


A mimicry ring

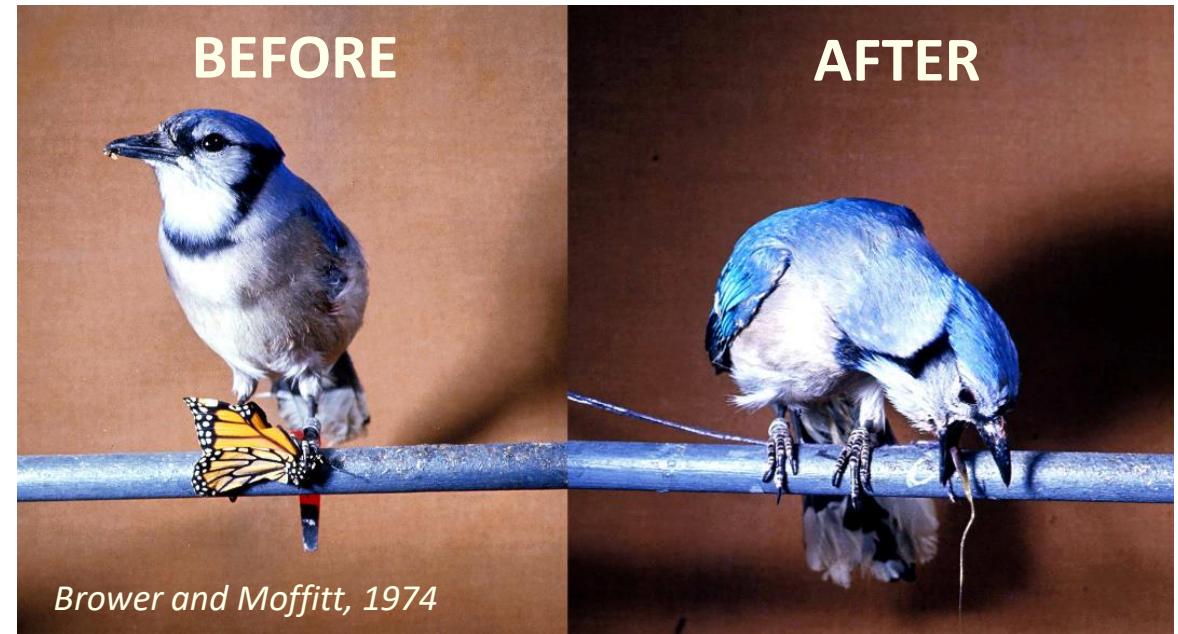


Credits: Google Earth

Study system: Müllerian mimicry



A mimicry ring



Brower and Moffitt, 1974



Study system: Müllerian mimicry

+ Let a_1 and a_2 be the numbers of two distasteful species of butterflies in some definite district during one summer, and let n be the number of individuals of a distinct species which are destroyed in the course of a summer before its distastefulness is generally known. If both species are totally dissimilar, then each loses n individuals. If, however, they are undistinguishably similar, then the first loses $\frac{a_1 n}{a_1 + a_2}$, and the second $\frac{a_2 n}{a_1 + a_2}$. The absolute gain by resemblance is therefore for the first species $n - \frac{a_1 n}{a_1 + a_2} = \frac{a_2 n}{a_1 + a_2}$; and in a similar manner for the second, $\frac{a_1 n}{a_1 + a_2}$. This absolute gain, compared with the occurrence of the species, gives for the first, $1_1 = \frac{a_2 n}{a_1 (a_1 + a_2)}$, and for the second species, $1_2 = \frac{a_1 n}{a_2 (a_1 + a_2)}$, whence follows the proportion, $1_1 : 1_2 = a_2^2 : a_1^2$.

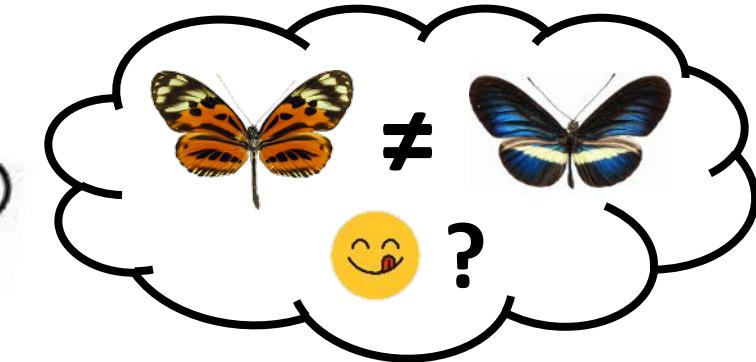
Müller, 1879



Fritz Müller
(1821 – 1897)

Study system: Müllerian mimicry

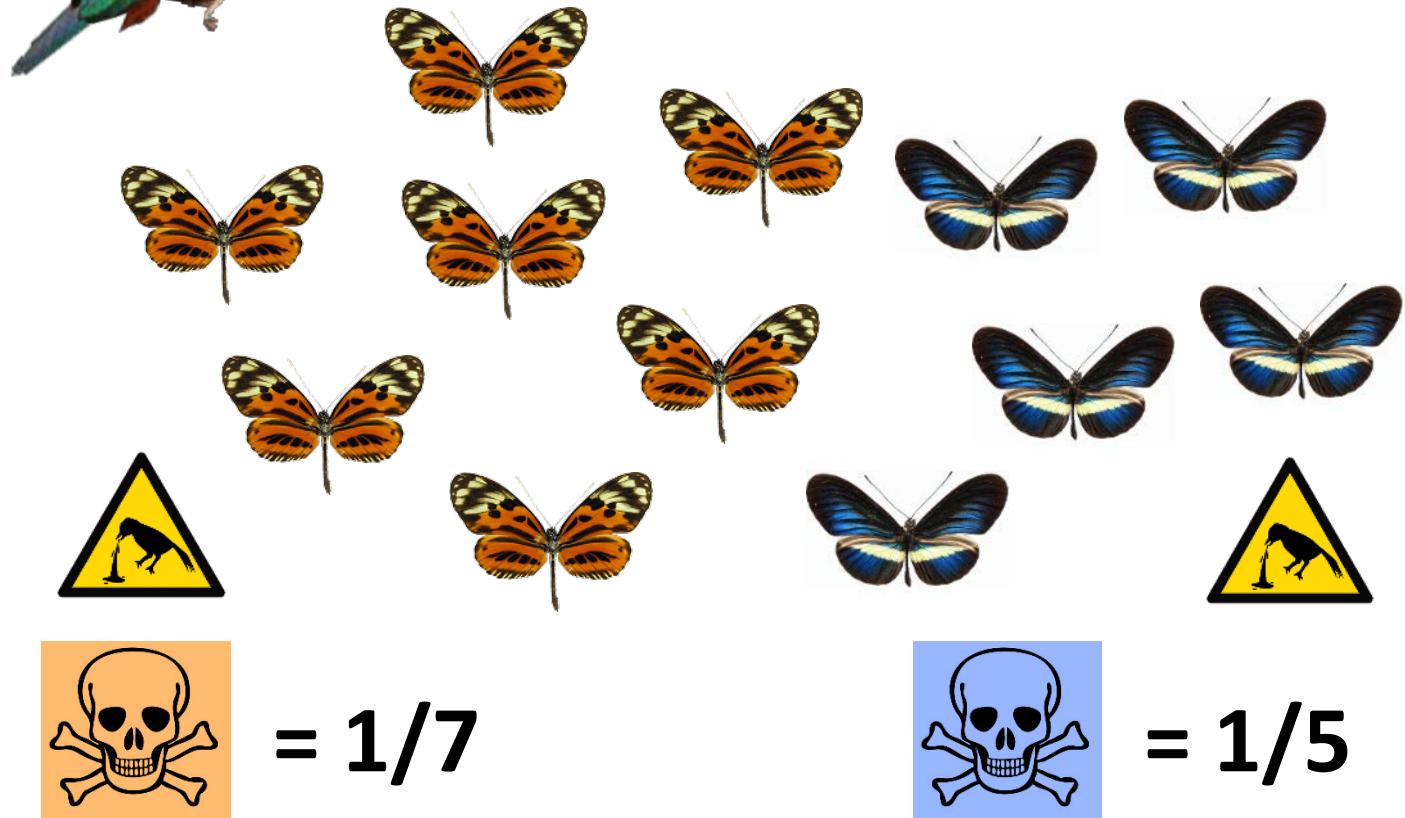
1/ Positive frequency-dependent selection



2/ Advantage for similarity

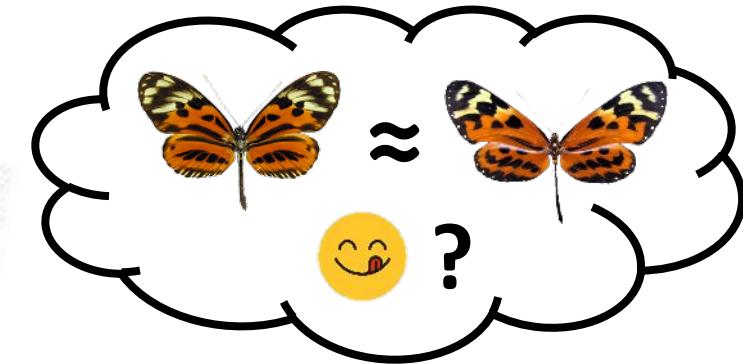
Consequences:

- Local pattern convergence
=> mimicry rings
- Mutual benefit from cooccurrence
=> mutualistic interactions



Study system: Müllerian mimicry

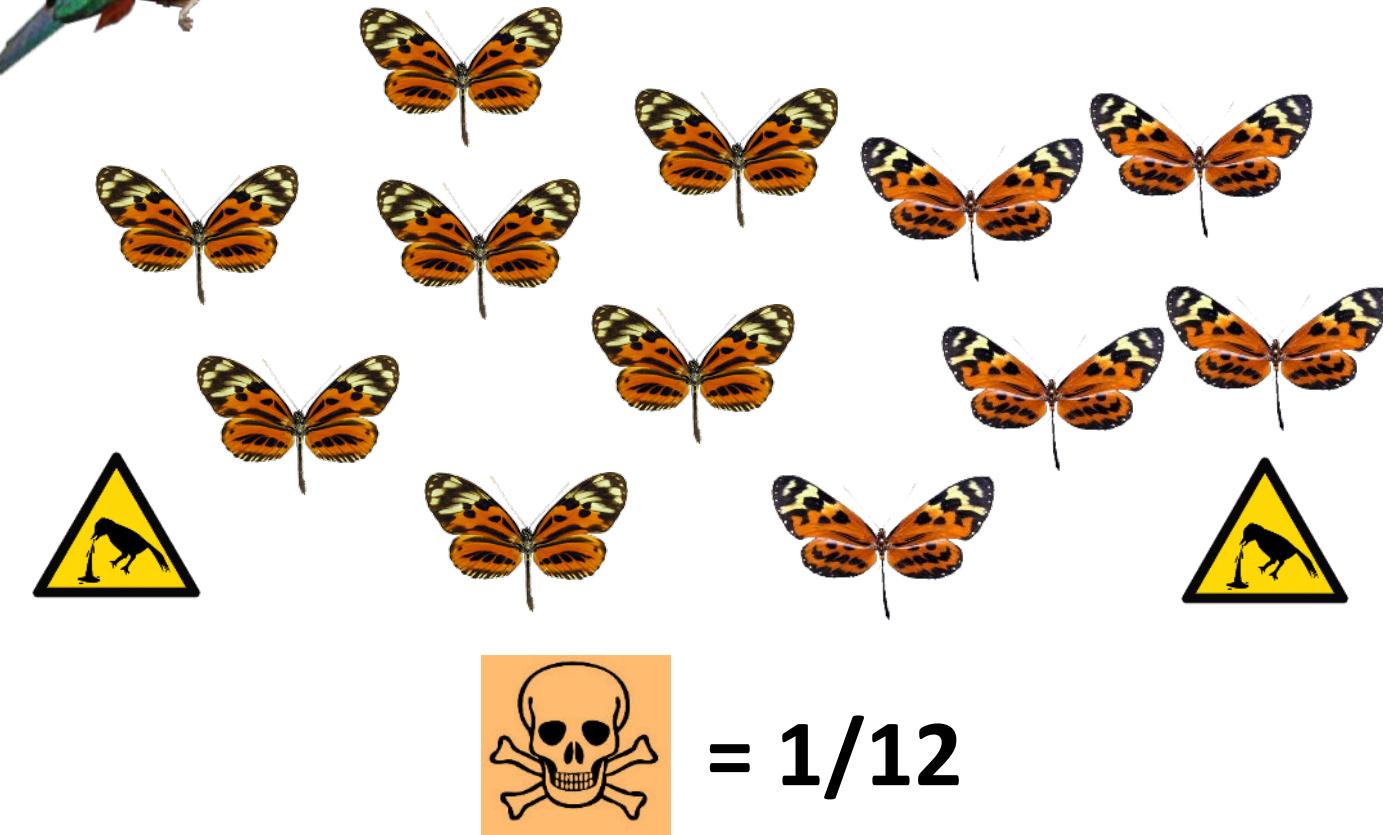
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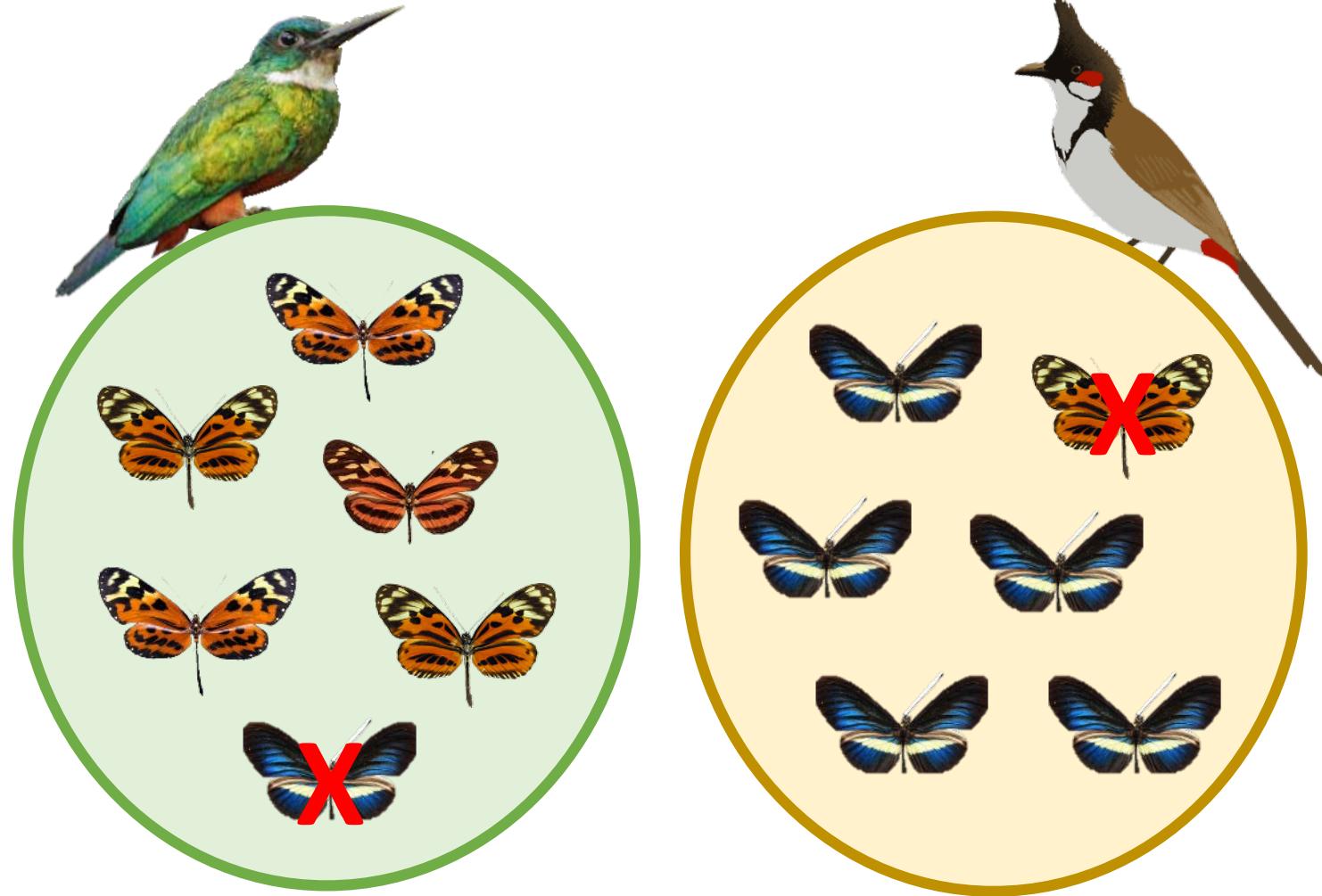
Study system: Müllerian mimicry

1/ Positive frequency-dependent selection

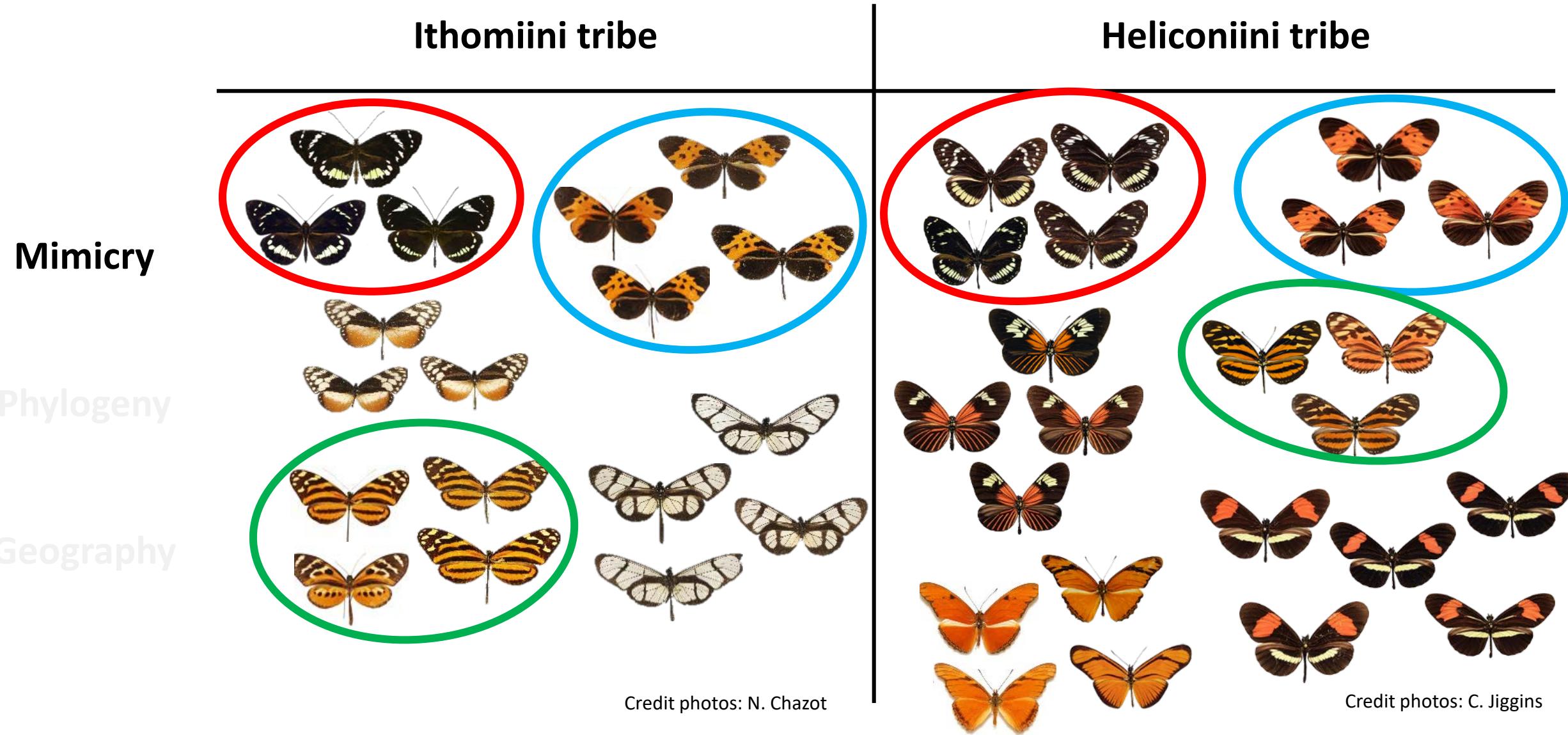
2/ Advantage for similarity

Consequences:

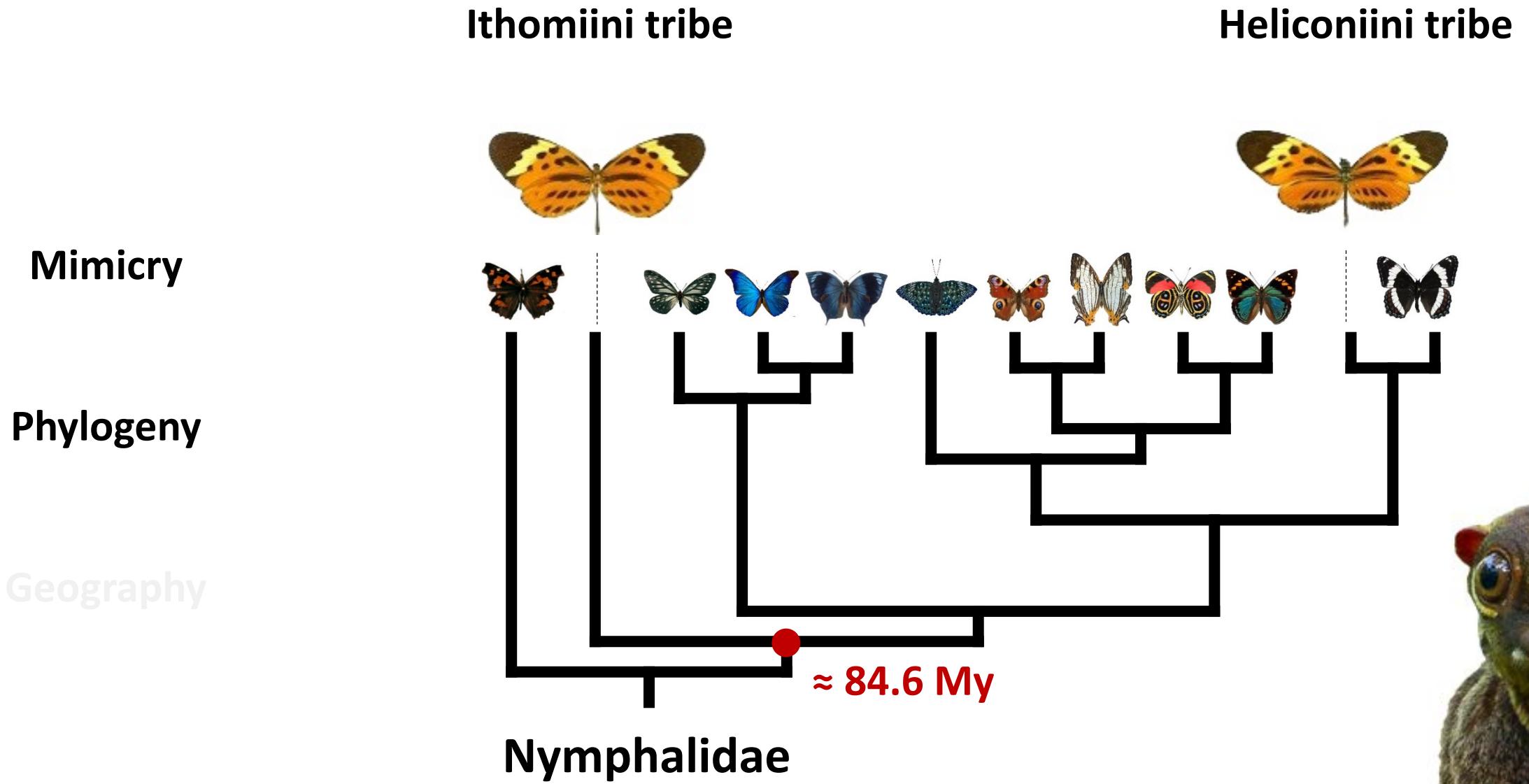
- Local pattern **convergence**
=> **mimicry rings**
- Mutual benefit from cooccurrence
=> **mutualistic interactions**



Study system: Neotropical butterflies



Study system: Neotropical butterflies



Study system: Neotropical butterflies

Mimicry

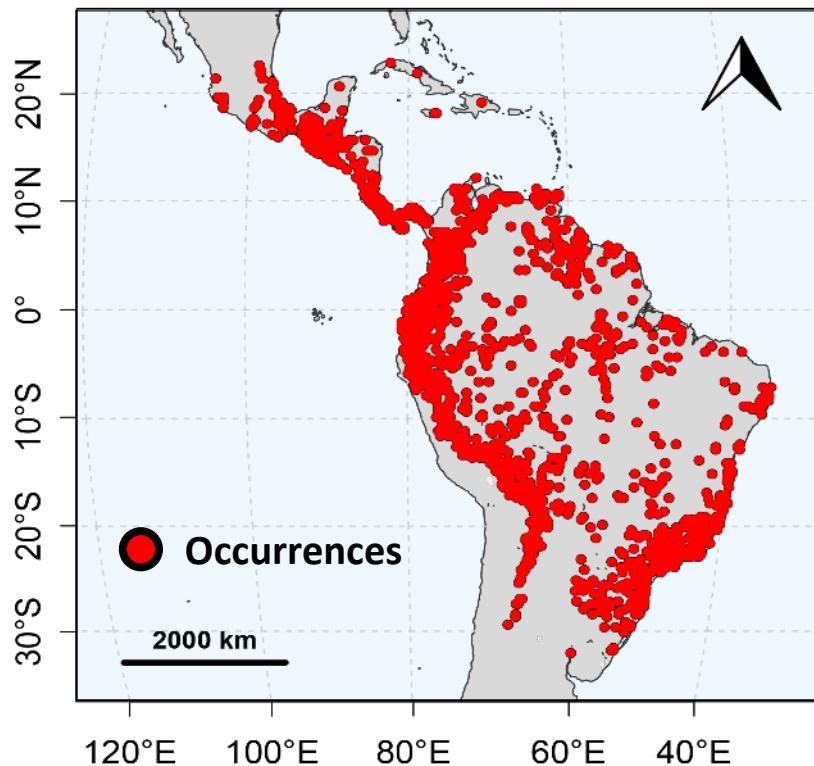
Phylogeny

Geography

Ithomiini tribe

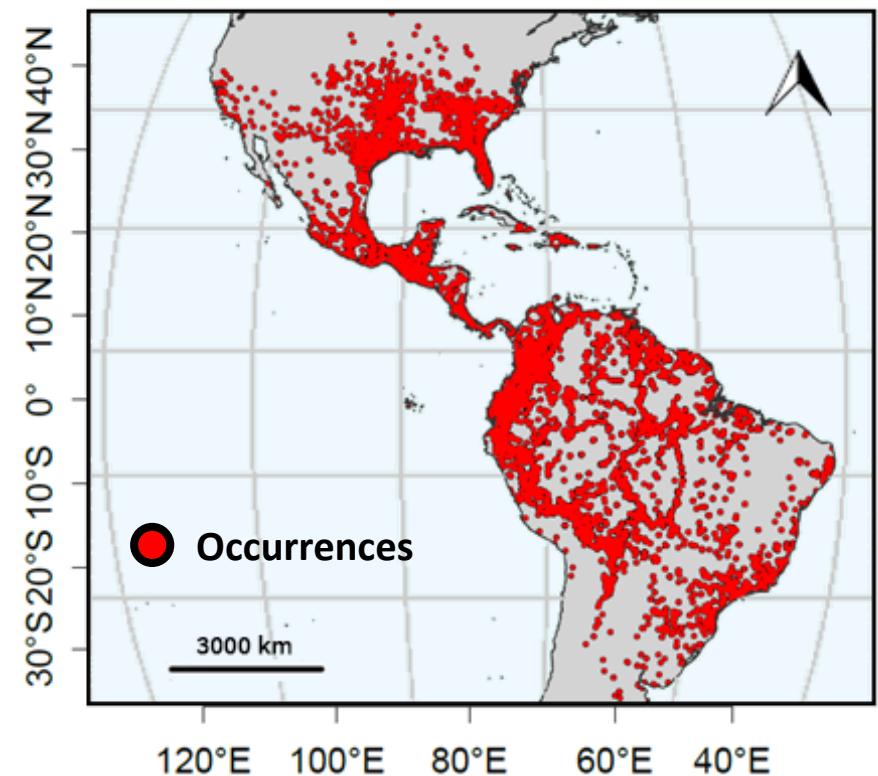
Heliconiini tribe

Occurrence map



Doré et al., 2022

Occurrence map



Perochon et al., in prep

Study system: Neotropical butterflies

Ithomiini tribe

Heliconiini tribe

Mimicry

Phylogeny

Geography



Credit Photo: M. Chouteau

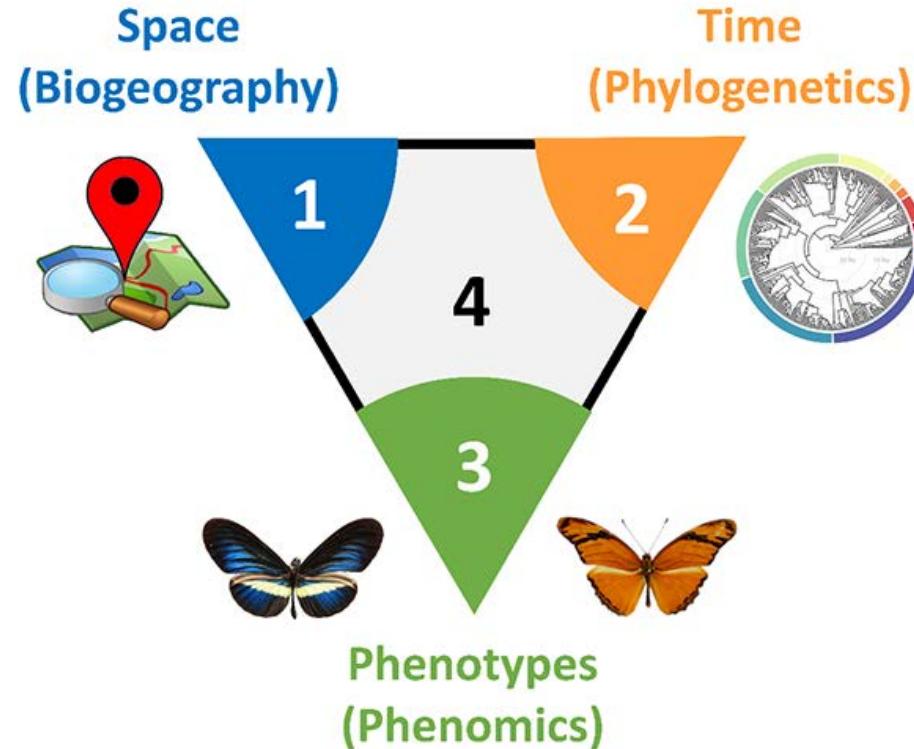
Outlines and objectives

CHAPTER 1

Map biodiversity patterns

CHAPTER 3

Quantify phenotypic similarity in wing patterns



How mutualistic interactions affect the structure and evolution of biodiversity at the macroecological scale?

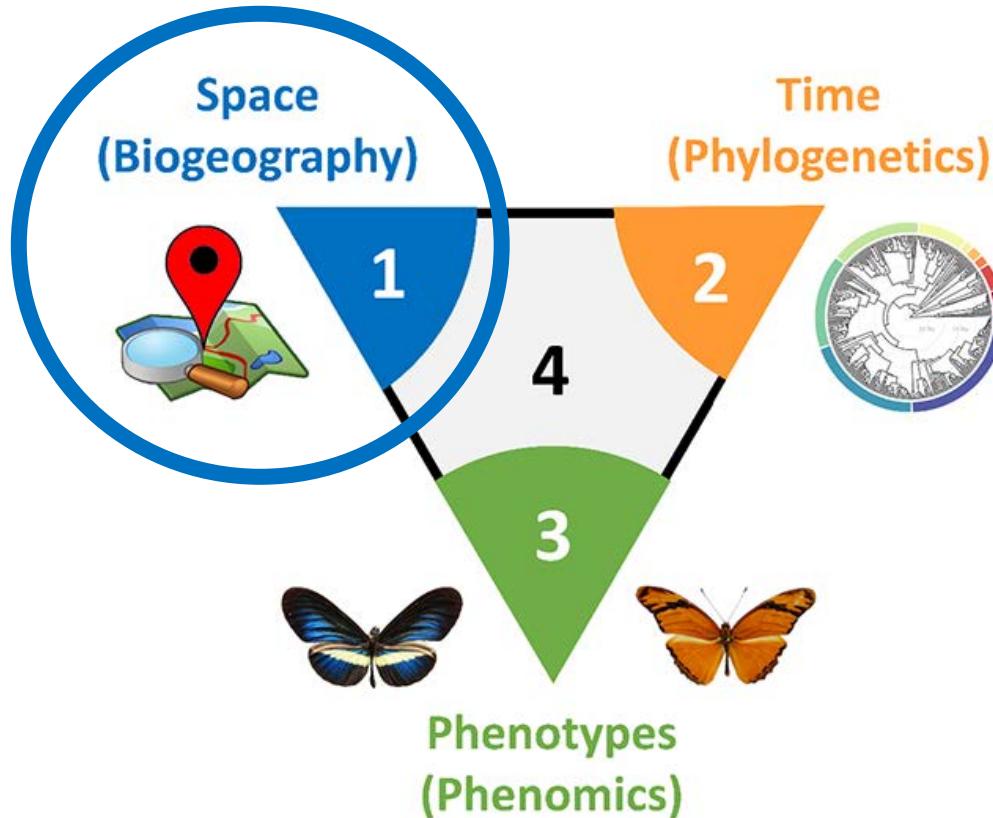
CHAPTER 2

Resolve deep evolutionary relationships

CHAPTER 1: Anthropogenic pressures coincide with Neotropical biodiversity hotspots in a flagship butterfly group

CHAPTER 1

Map biodiversity patterns



Reference:

Doré, M., Willmott, K., Leroy, B., Chazot, N., Mallet, J., Freitas, A. V. L., Hall, J. P. W., Lamas, G., Dasmahapatra, K. K., Fontaine, C., & Elias, M. (2022). Anthropogenic pressures coincide with Neotropical biodiversity hotspots in a flagship butterfly group. *Diversity and Distributions*, 28(12), 2912–2930. <https://doi.org/10.1111/ddi.13455>

Stage M2: Heliconiini



Eddie Pérochon

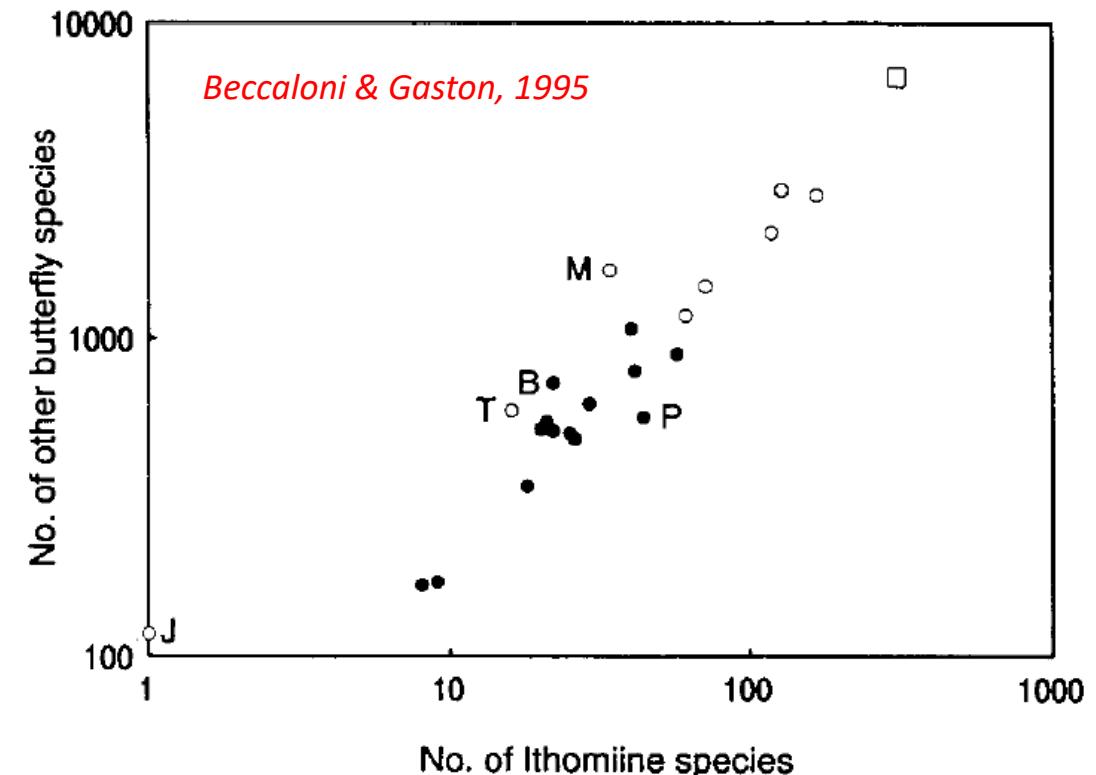
Context & Objectives

Ithomiini as indicator species

- For forest habitat quality
(Brown Jr 1997; Uehara-Prado & Freitas 2009)
- For overall butterfly diversity
(Beccaloni & Gaston, 1995)

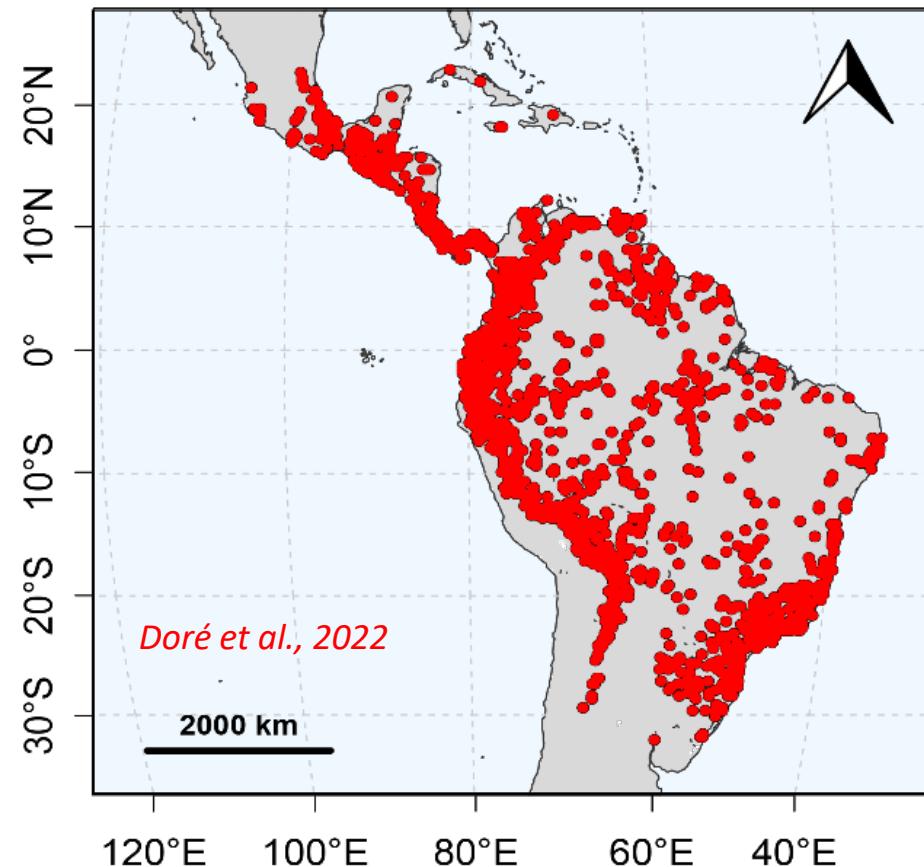
Objectives

- 1/ Map species distributions
- 2/ Map biodiversity patterns for ithomiine (and heliconiine)
- 3/ Identify risk areas and refuge areas against anthropogenic threats



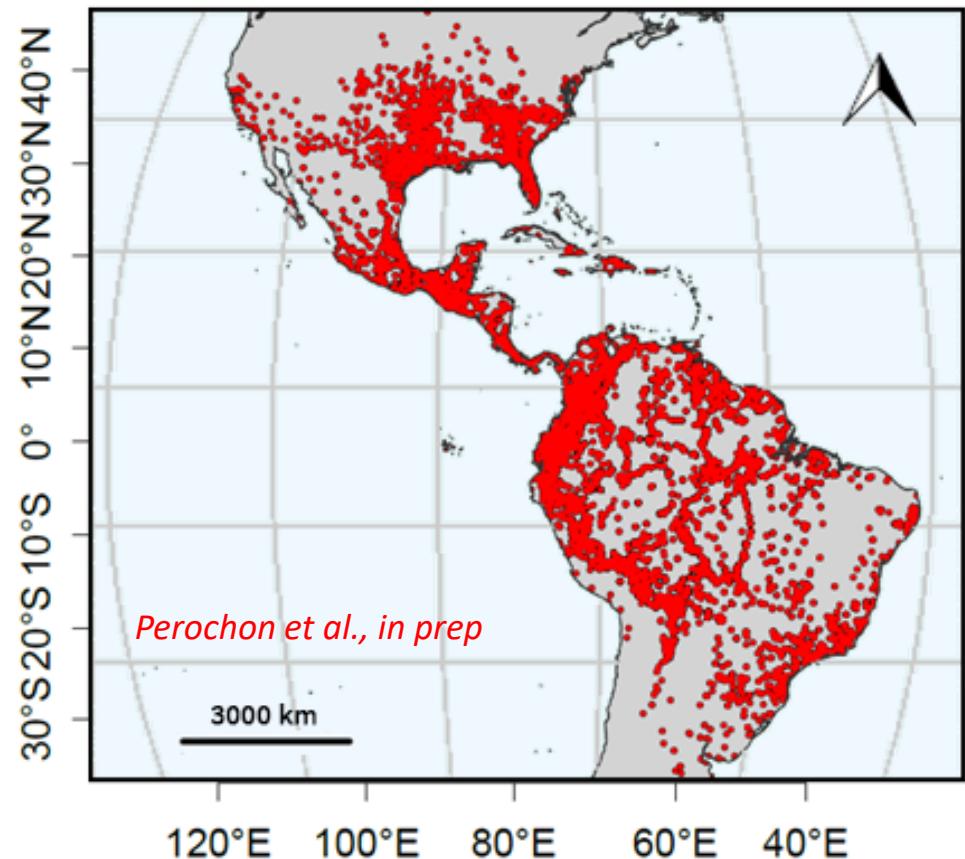
Available data

Ithomiini tribe (396 species)



28,986 occurrences across 1,834 sites

Heliconiini tribe (77 species)



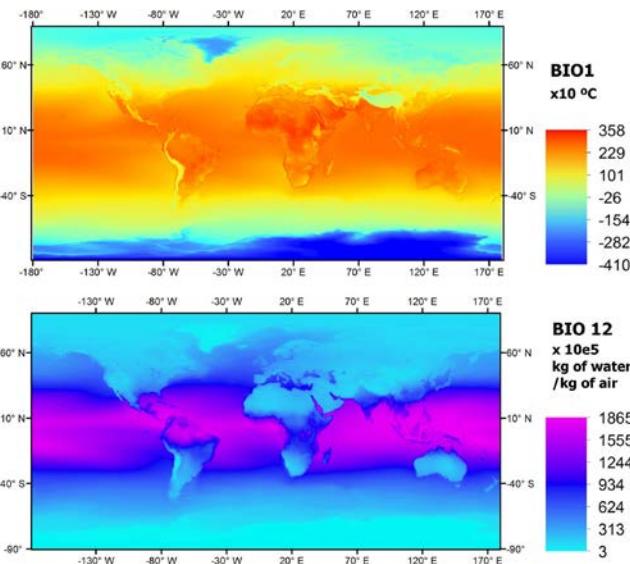
77,577 occurrences across 4,168 sites

Available data

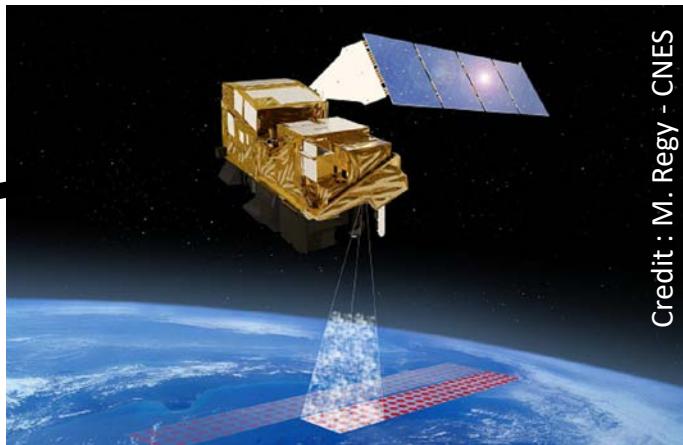
Climate data

MERRAclim v.2.0

- Annual temperature
- Humidity levels
- Temperature seasonality
- Humidity seasonality



Vega et al., 2017



Credit : M. Regy - CNES

Vegetation cover

GFCC v.3.0



Sexton et al., 2013

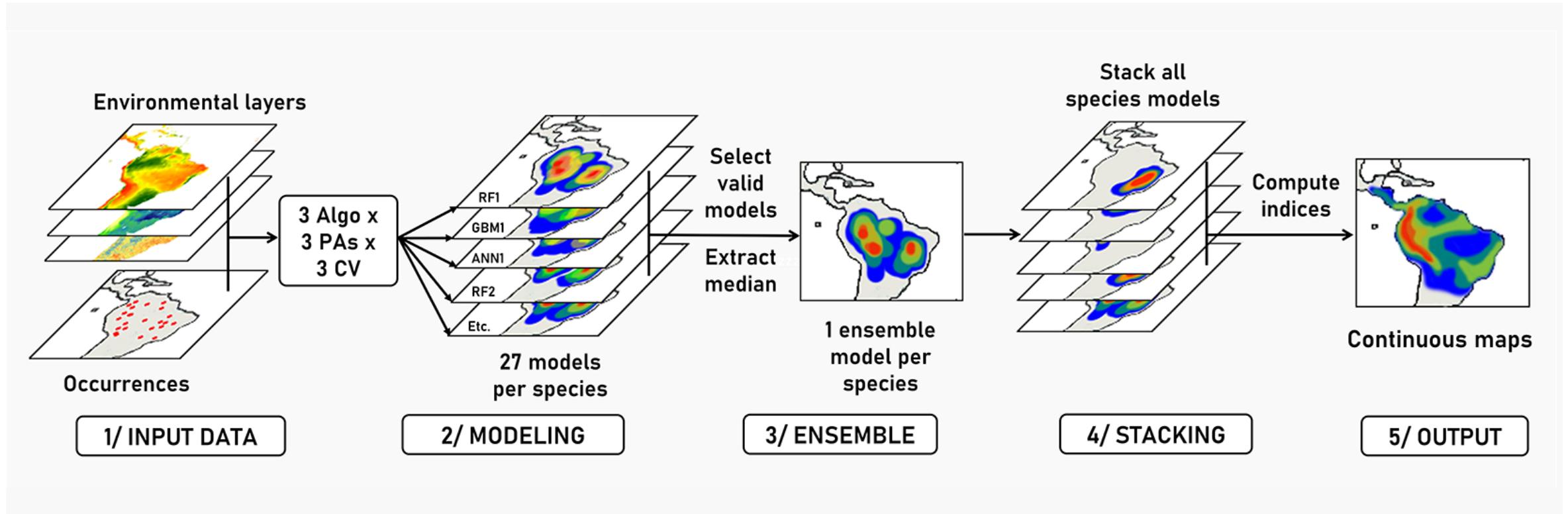
Elevation

SRTM Dataset v.4.1



Farr et al., 2007

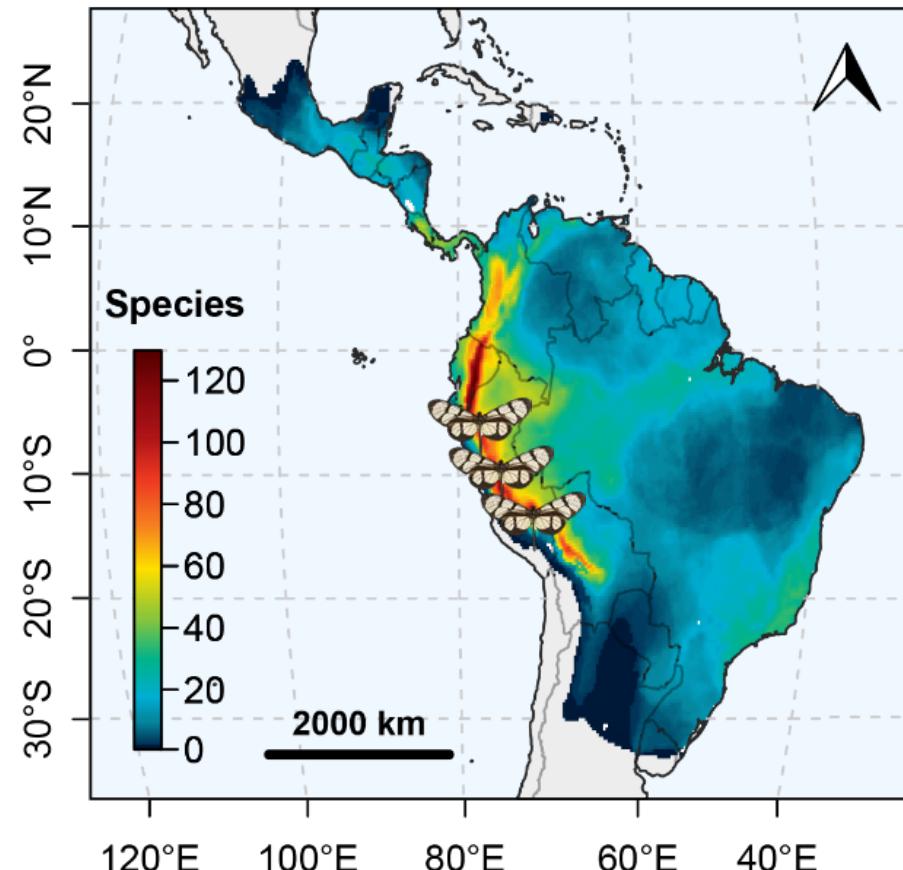
Species Distribution Modeling (SDM)



SDM → Species distribution maps → Diversity indices

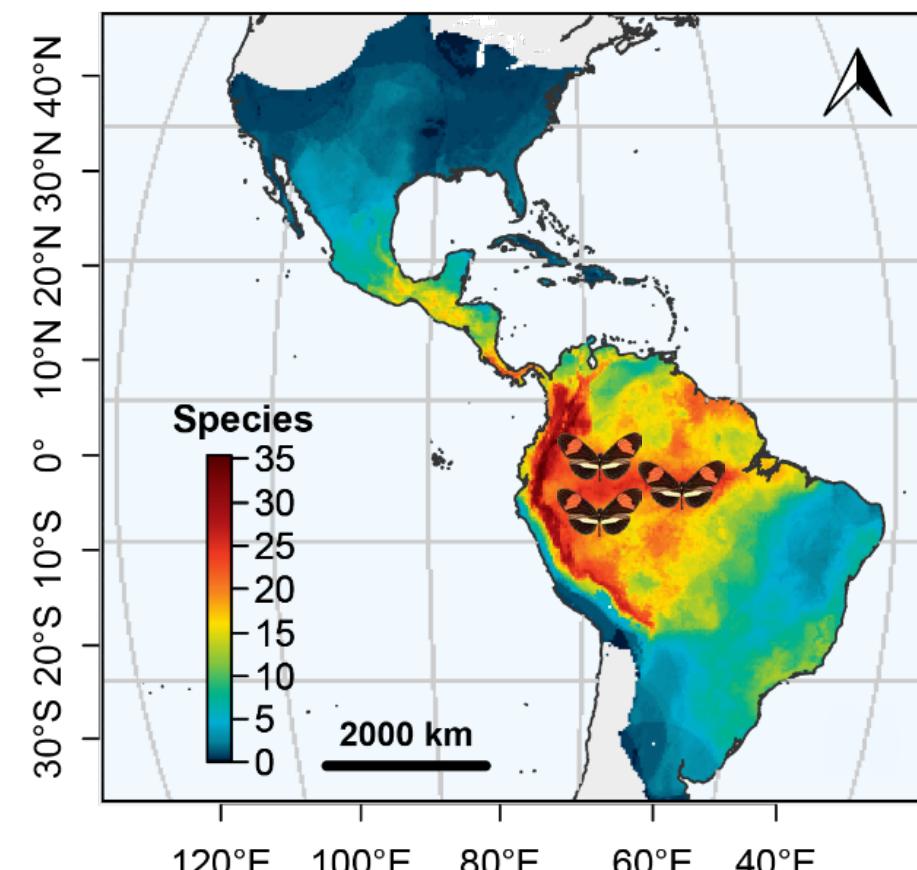
Biodiversity patterns

Ithomiini
(396 species)



Species richness

Heliconiini
(77 species)



Doré et al., 2022

Pérochon et al., in prep

Anthropogenic threats

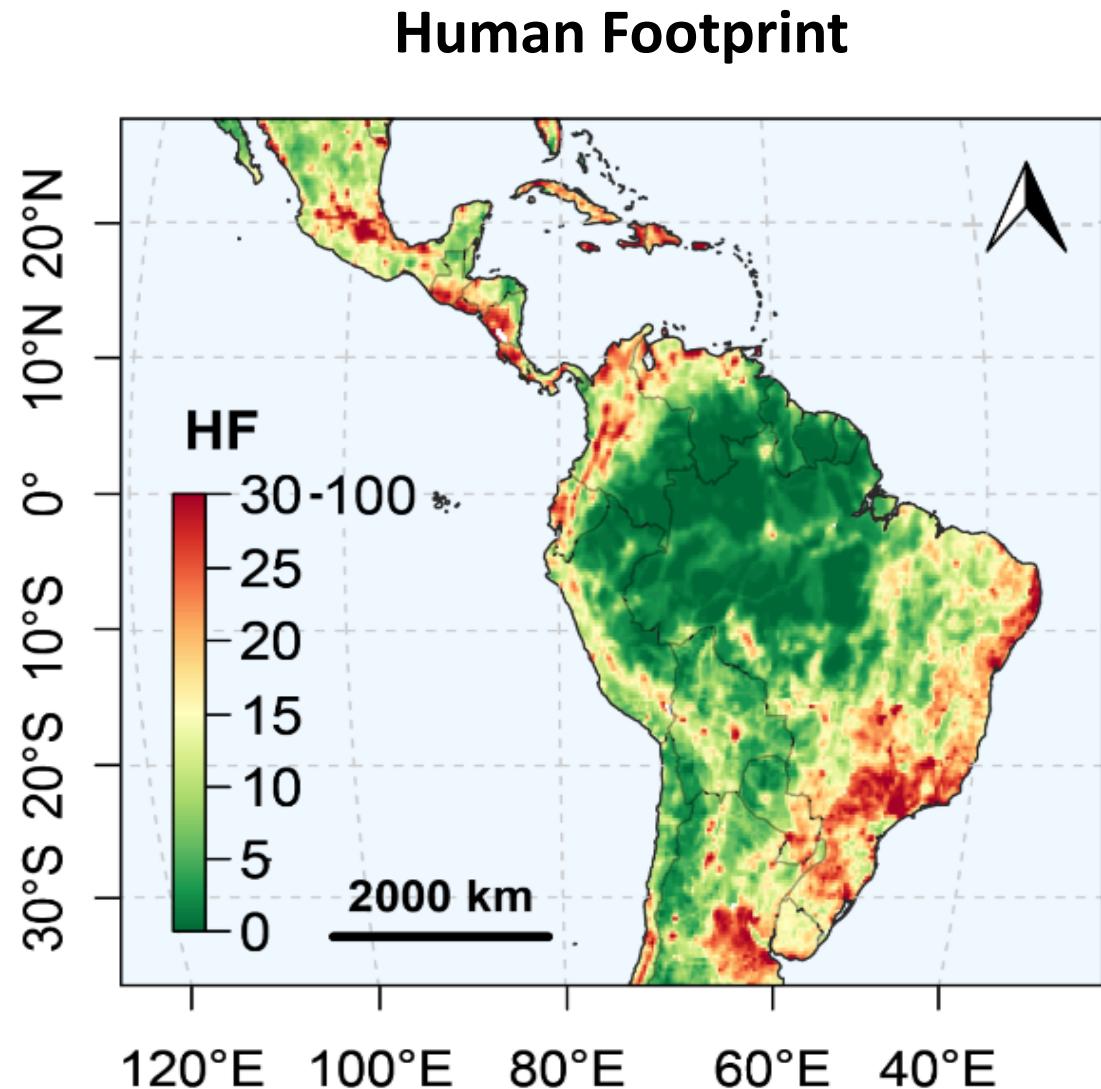
Identify **risk areas** and **refuge areas** against anthropogenic threats

Current anthropogenic threats

≈

Human Footprint

- (1) Human population density
- (2) Landcover: urban, crop, pasture, wilderness
- (3) Proximity to transport infrastructures: railways, major roadways, and navigable waterways
- (4) Night-time light pollution



Venter et al., 2016

Anthropogenic threats

Identify **risk areas** and **refuge areas** against anthropogenic threats

Current anthropogenic threats

≈

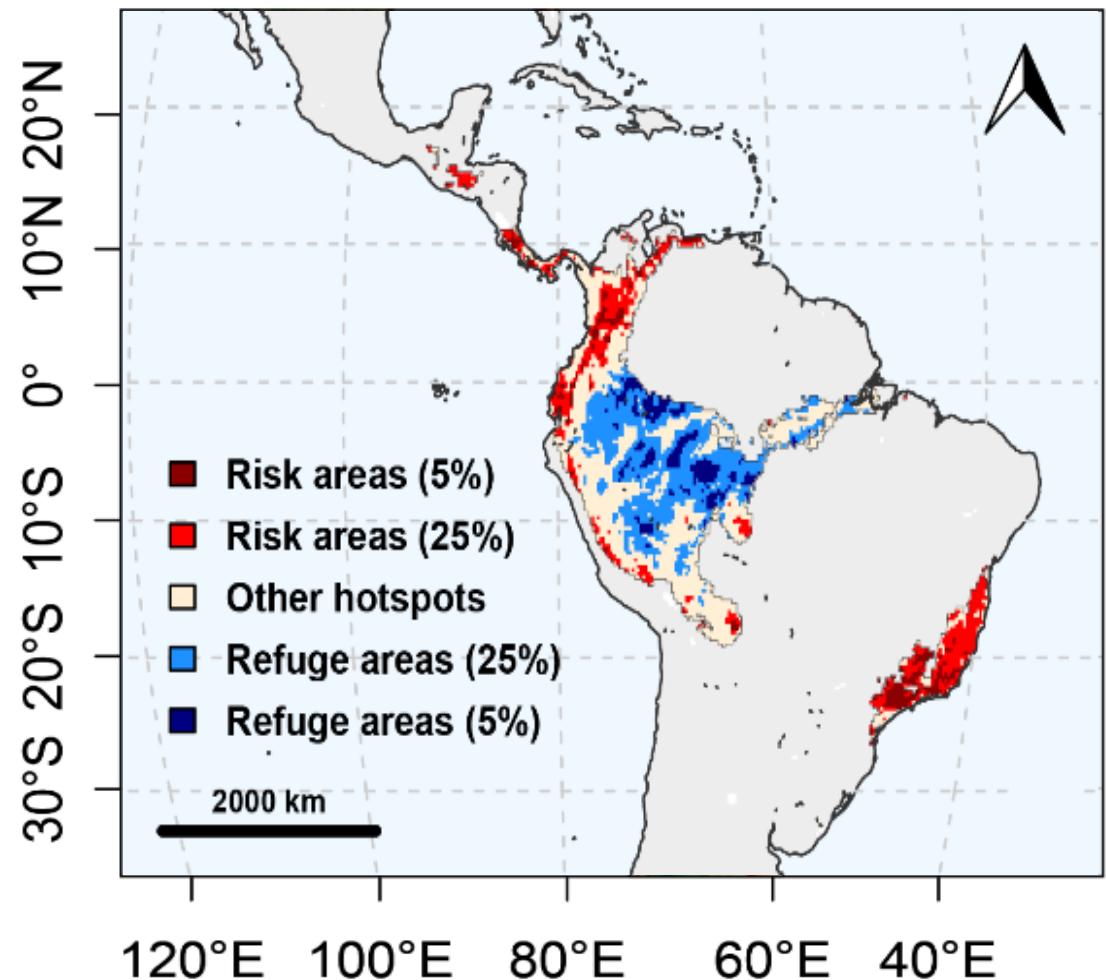
Human Footprint

Risk areas: Tropical Andes, Central America, Brazilian Atlantic Forest

Potential **refuge area**: Western Amazon (?)

Limits: Low redundancy with other areas
Climate change and deforestation

Species richness
Top 25% hotspots



Doré et al., 2022

Anthropogenic threats

Identify **risk areas** and **refuge areas** against anthropogenic threats

Current anthropogenic threats

≈

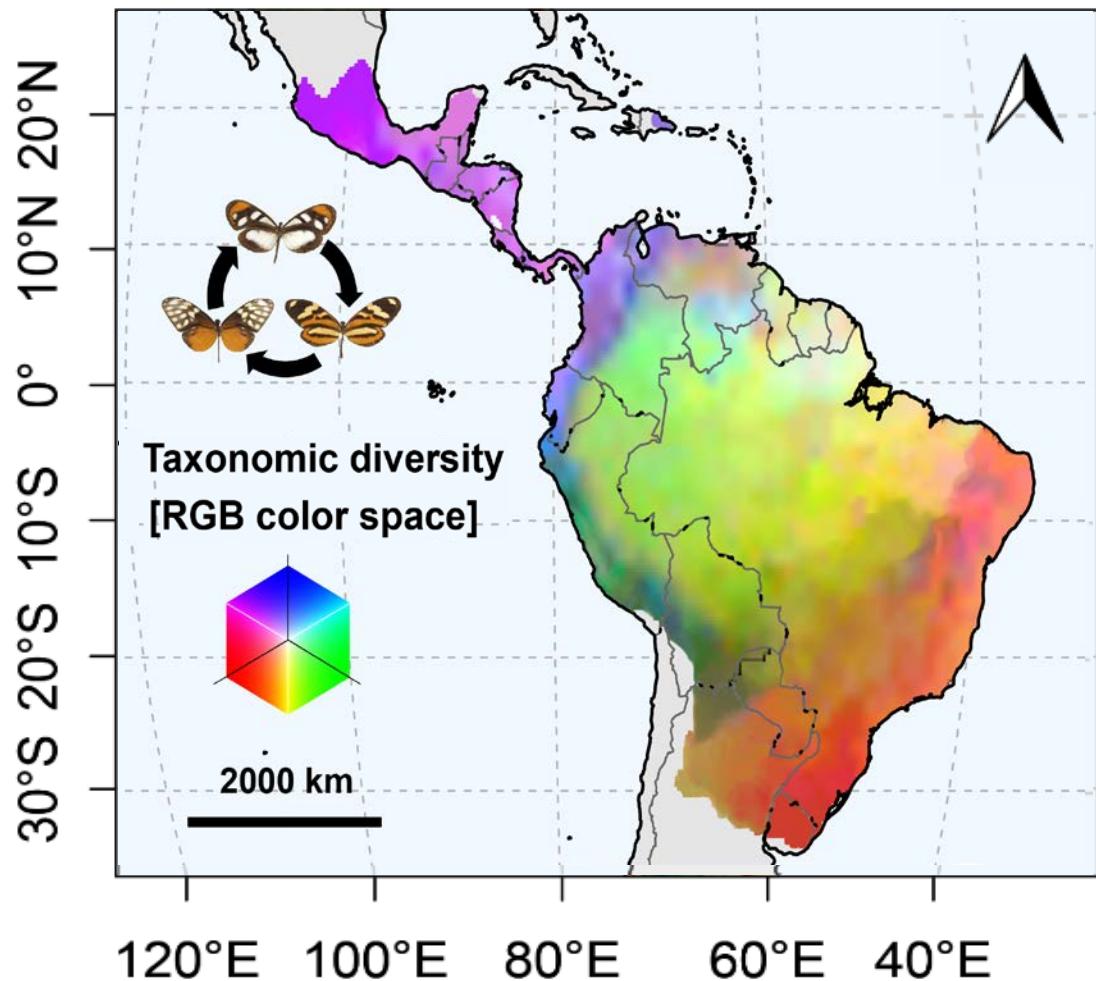
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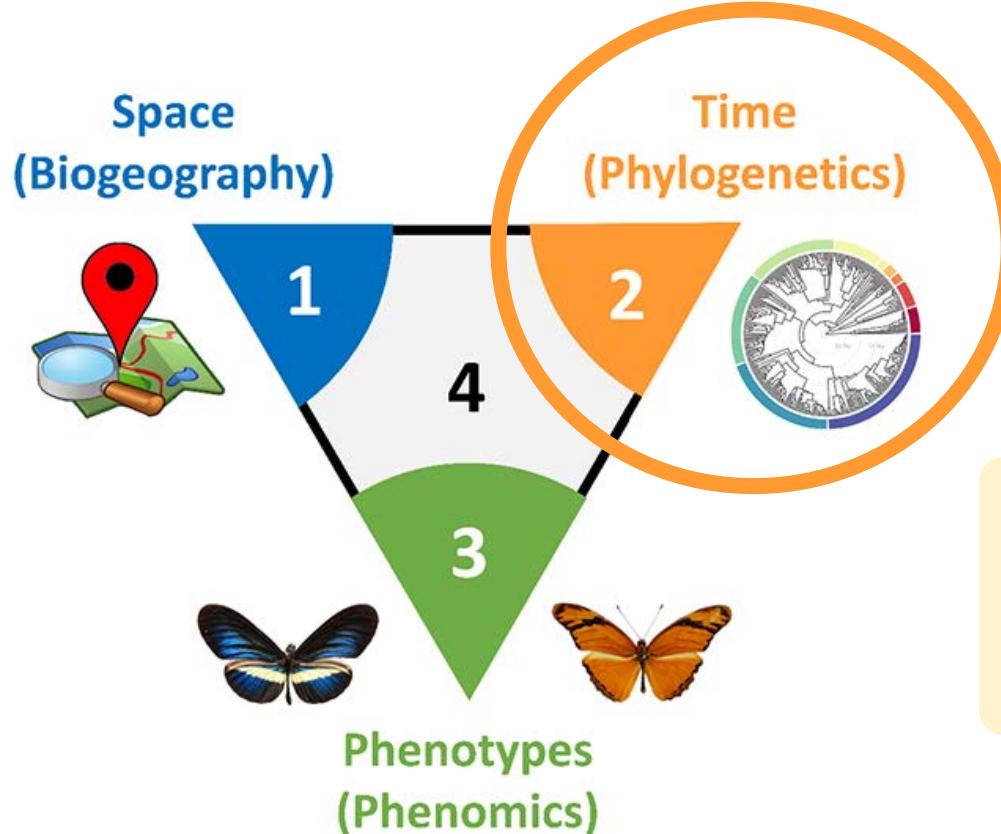
Potential **refuge area**: Western Amazon (?)

Limits: **Low redundancy** with other areas
Climate change and deforestation

Taxonomic β -Diversity



CHAPTER 2: Phylogenomics resolve deep evolutionary relationships in clearwing butterflies

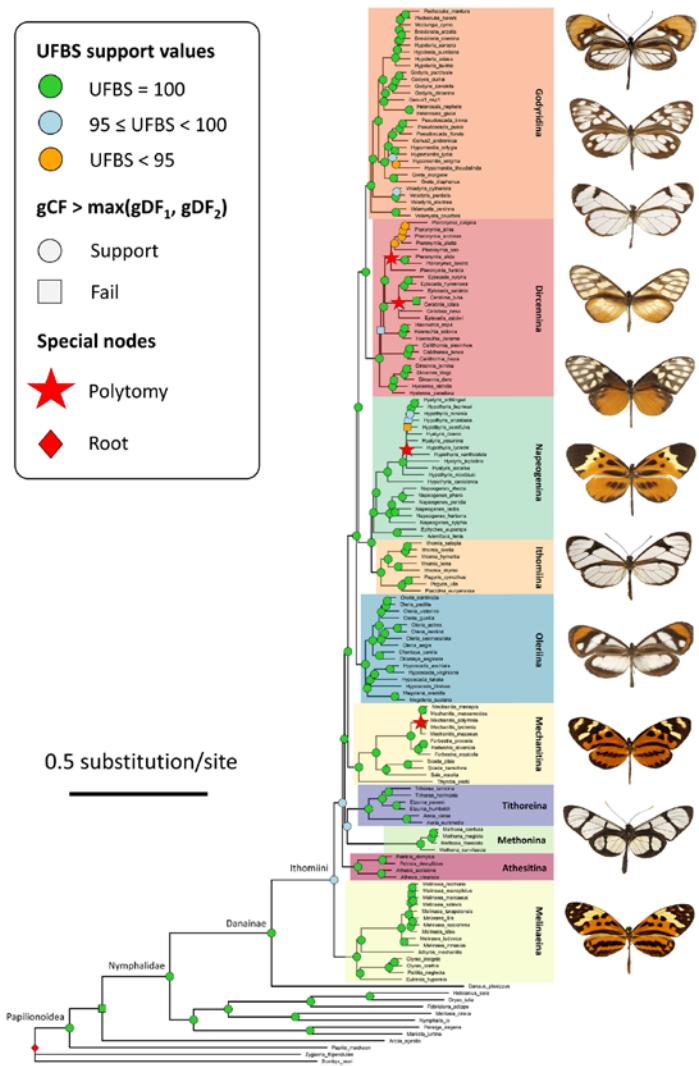


CHAPTER 2
Resolve deep
evolutionary relationships

Reference:

Doré, M., Gauthier, J., Allio, R., Grishin, N., Meier, J., Chazot, N., Willmott, K., & Elias, M. (2023). Phylogenomics resolve deep evolutionary relationships in clearwing butterflies. *In prep.*

From a WGS-based backbone...



153 ithomiines + 12 outgroups

39 % of extent species
100% of extent genera

7.8 M sites across 11,012 genes

... to a comprehensive Time-calibrated phylogeny

356 ithomiines + 12 outgroups

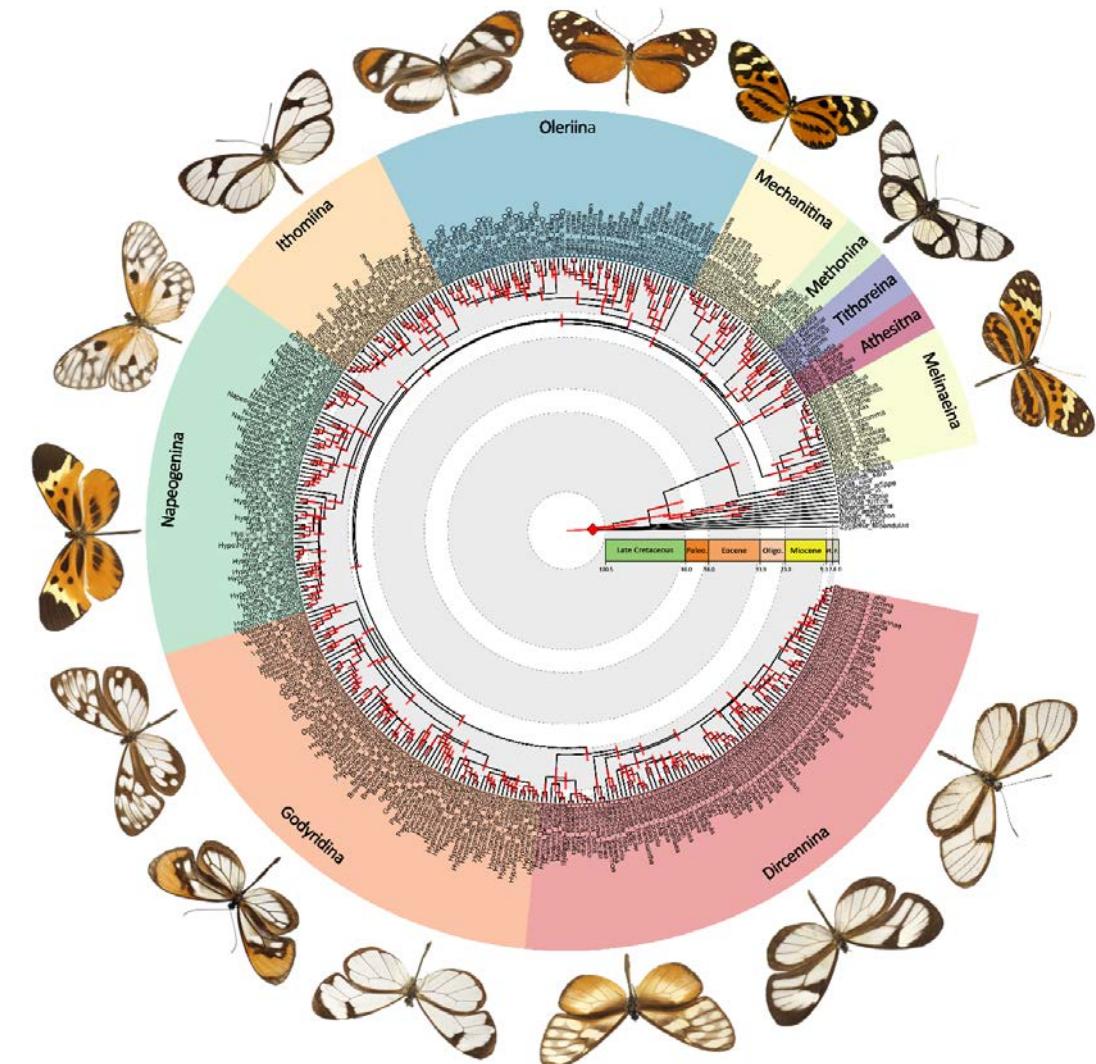
90 % of extent species

100% of extent genera

9,930 sites across 8 genes

Fixed backbone from WGS phylogeny

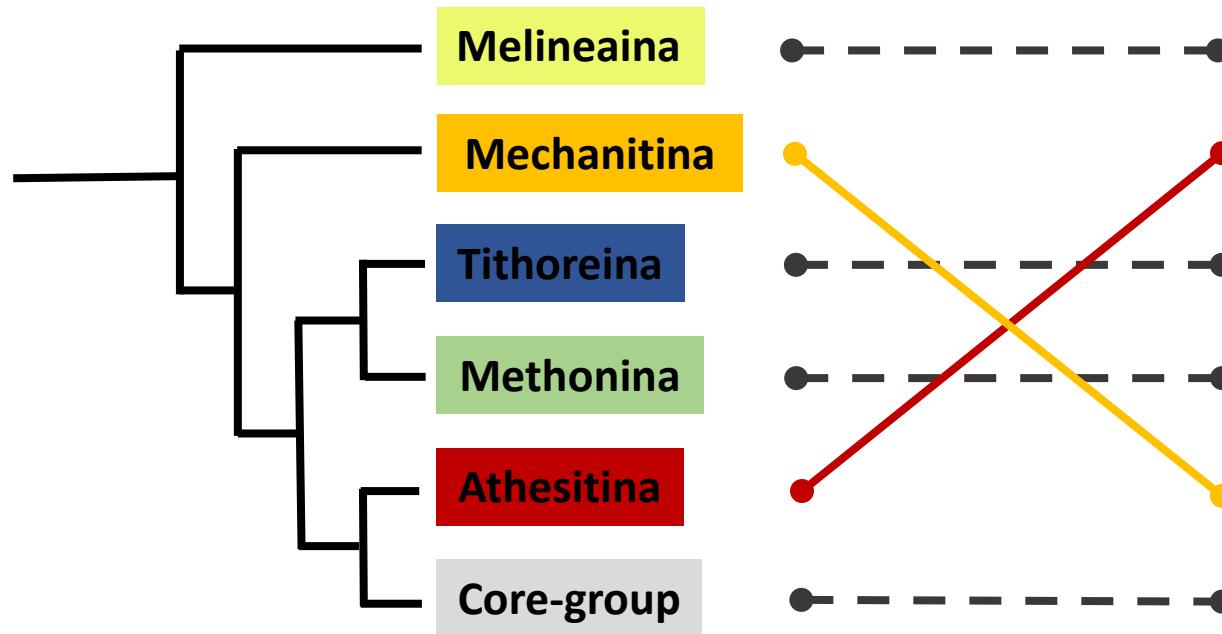
Secondary calibrations
from host-plant ages
and higher-level phylogenies



Phylogenomics resolving conflicts

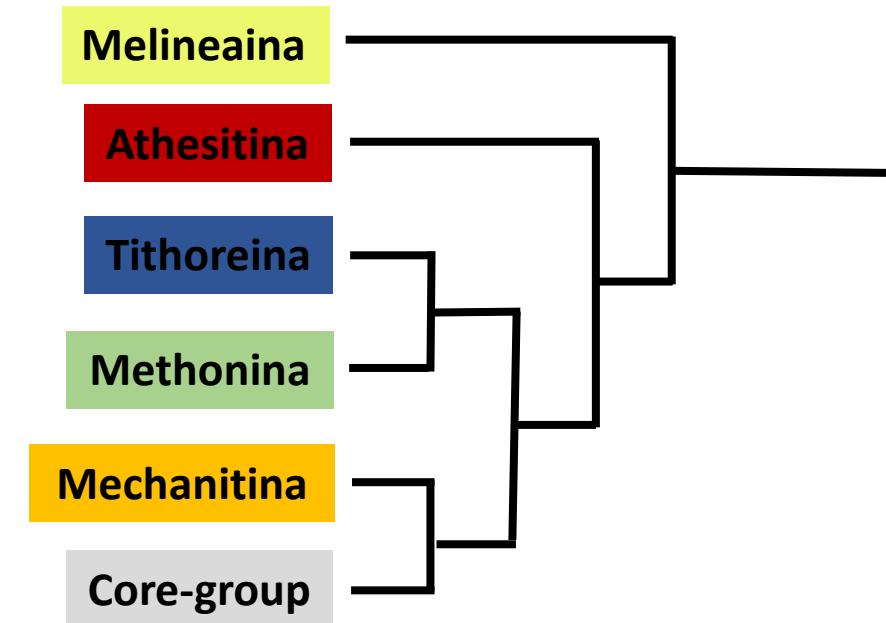
Sanger sequencing

Chazot *et al.*, 2019



Phylogenomics

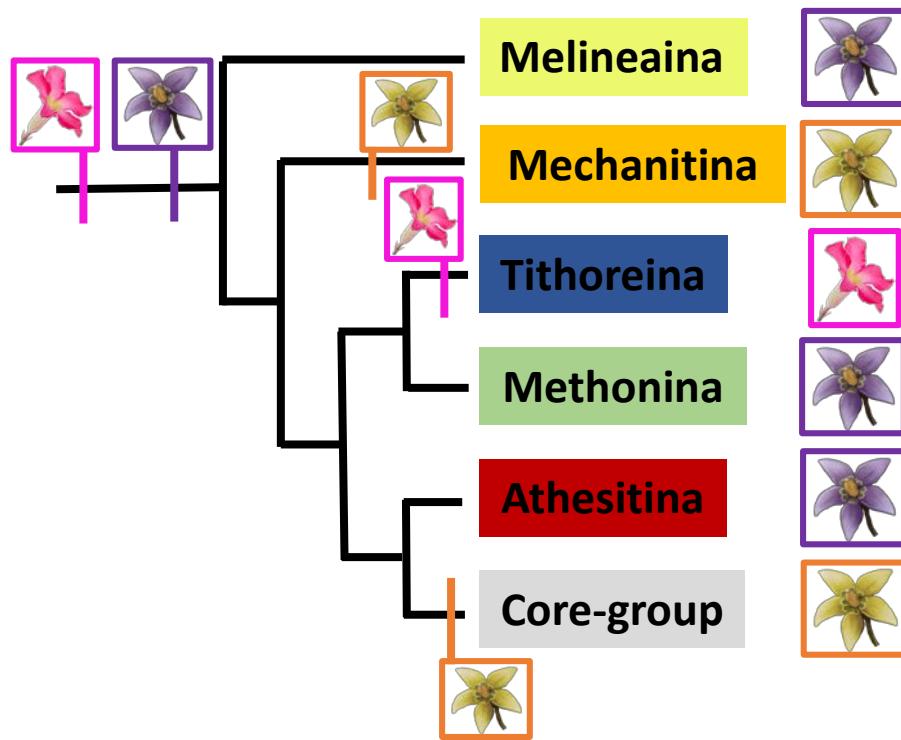
Doré *et al.*, *in prep.*



Phylogenomics resolving conflicts

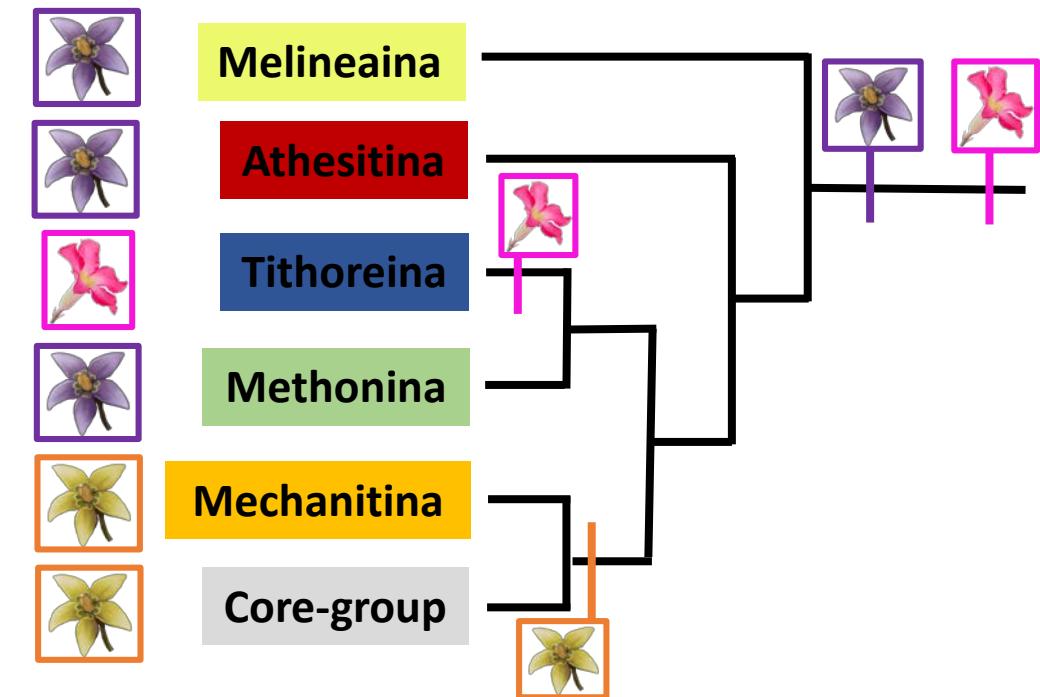
Sanger sequencing

Chazot *et al.*, 2019



Phylogenomics

Doré *et al.*, *in prep.*



Apocynaceae



Solanum

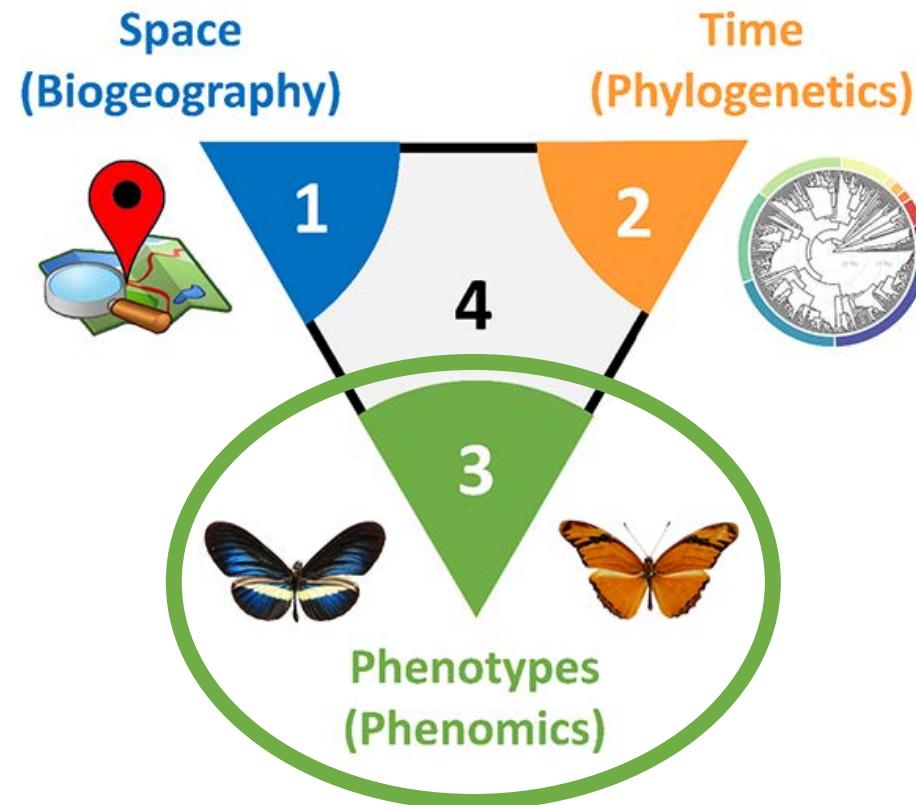


Other Solaneaceae

CHAPTER 3: Perceptual maps: a new tool to investigate mimicry patterns from Citizen Science to individual perception

CHAPTER 3

Quantify phenotypic similarity in wing patterns



Reference:

Doré, M., Pérochon, E., Aubier, T.G., Joron*, M. & Elias*, M. (2023). Perceptual maps: a new tool to investigate mimicry patterns from Citizen Science to individual perception. *In prep.*

The Heliconiini tribe as study model

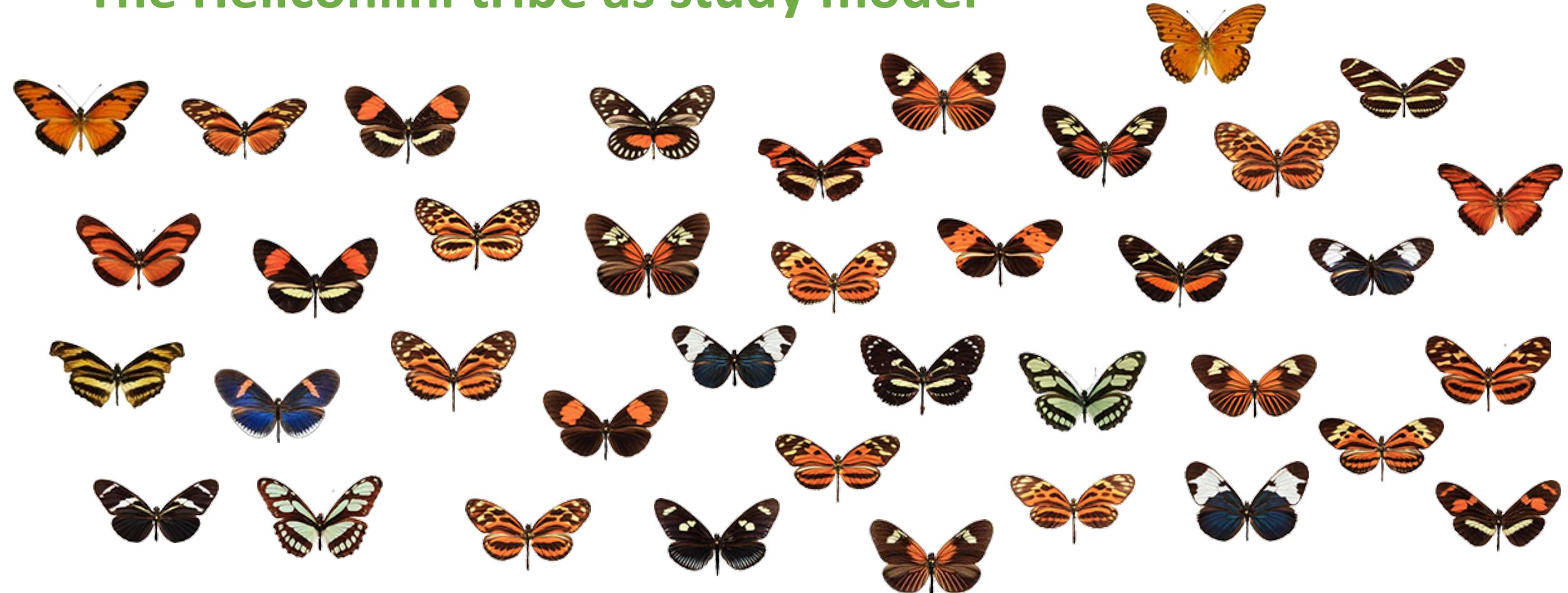


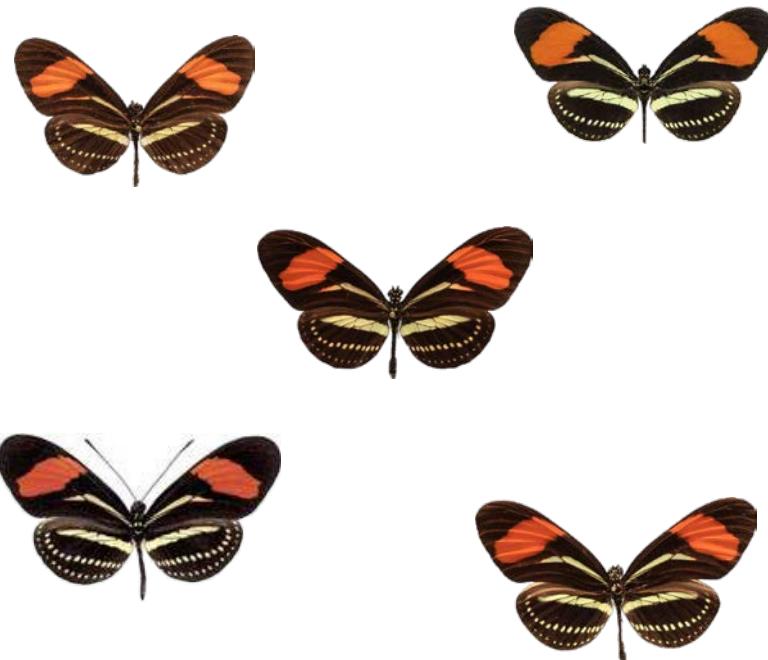
Photo credits: C. Jiggins

Model for the study of **genetics of color patterns** and **convergent evolution**

Yet... no standardized definition of **mimicry rings** in the group!

How to delineate mimicry rings?

POSTMAN ring



RED BAND ring

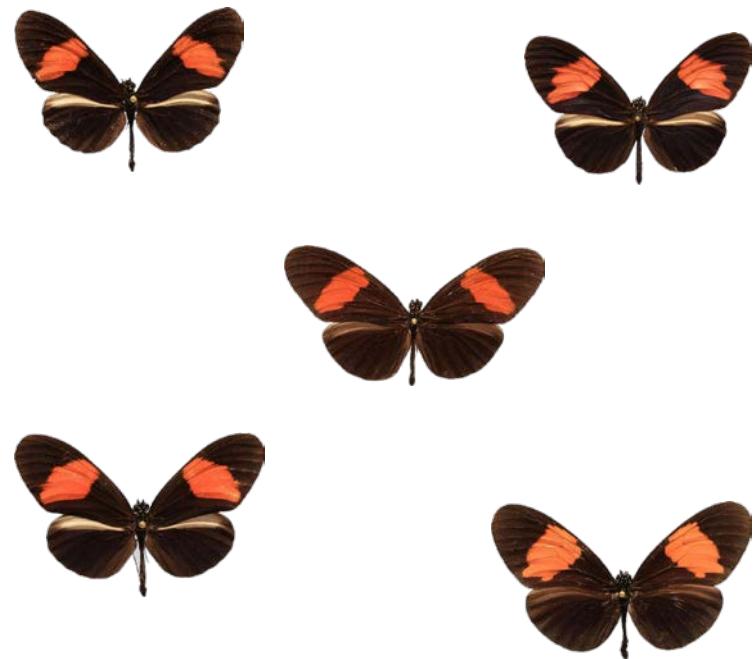
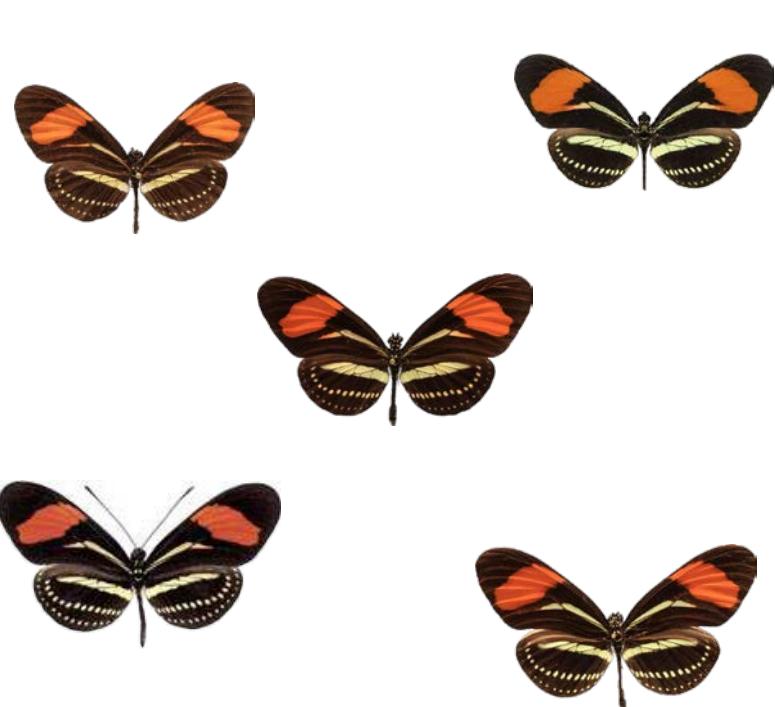


Photo credits: C. Jiggins

How to delineate mimicry rings?

POSTMAN ring



RED BAND ring

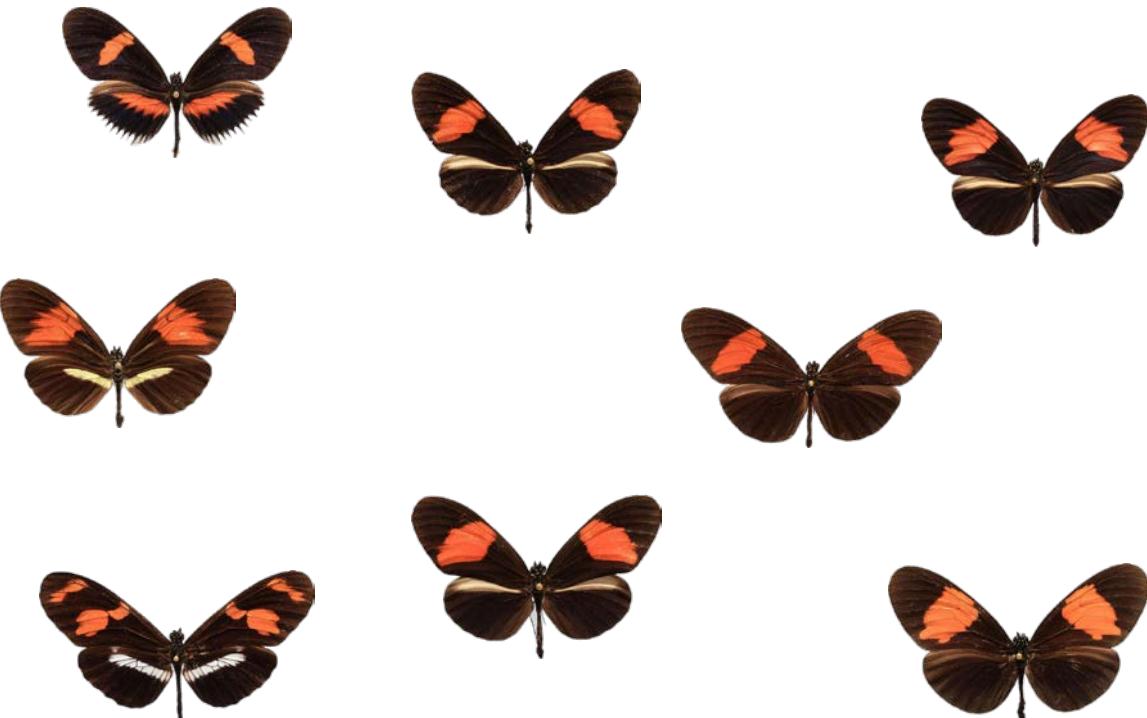
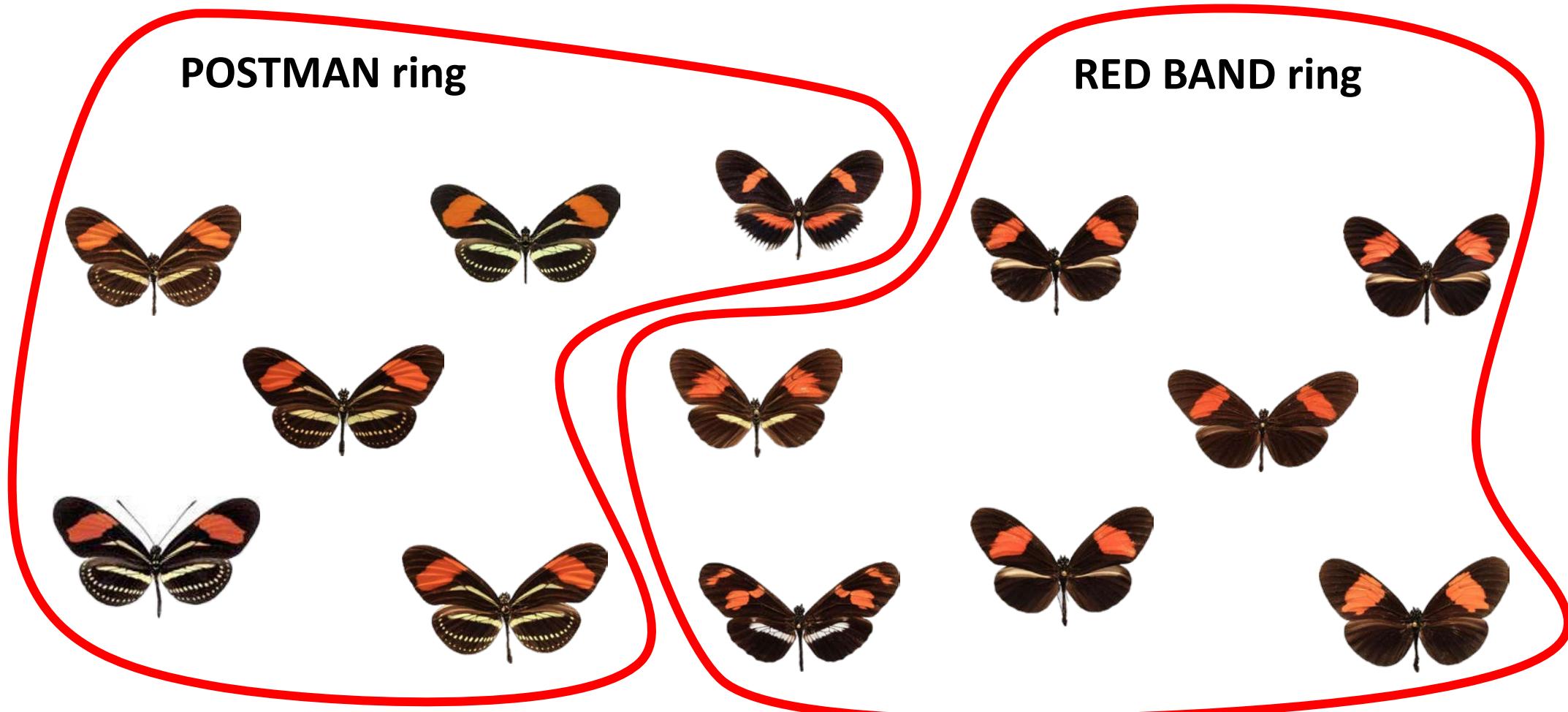


Photo credits: C. Jiggins

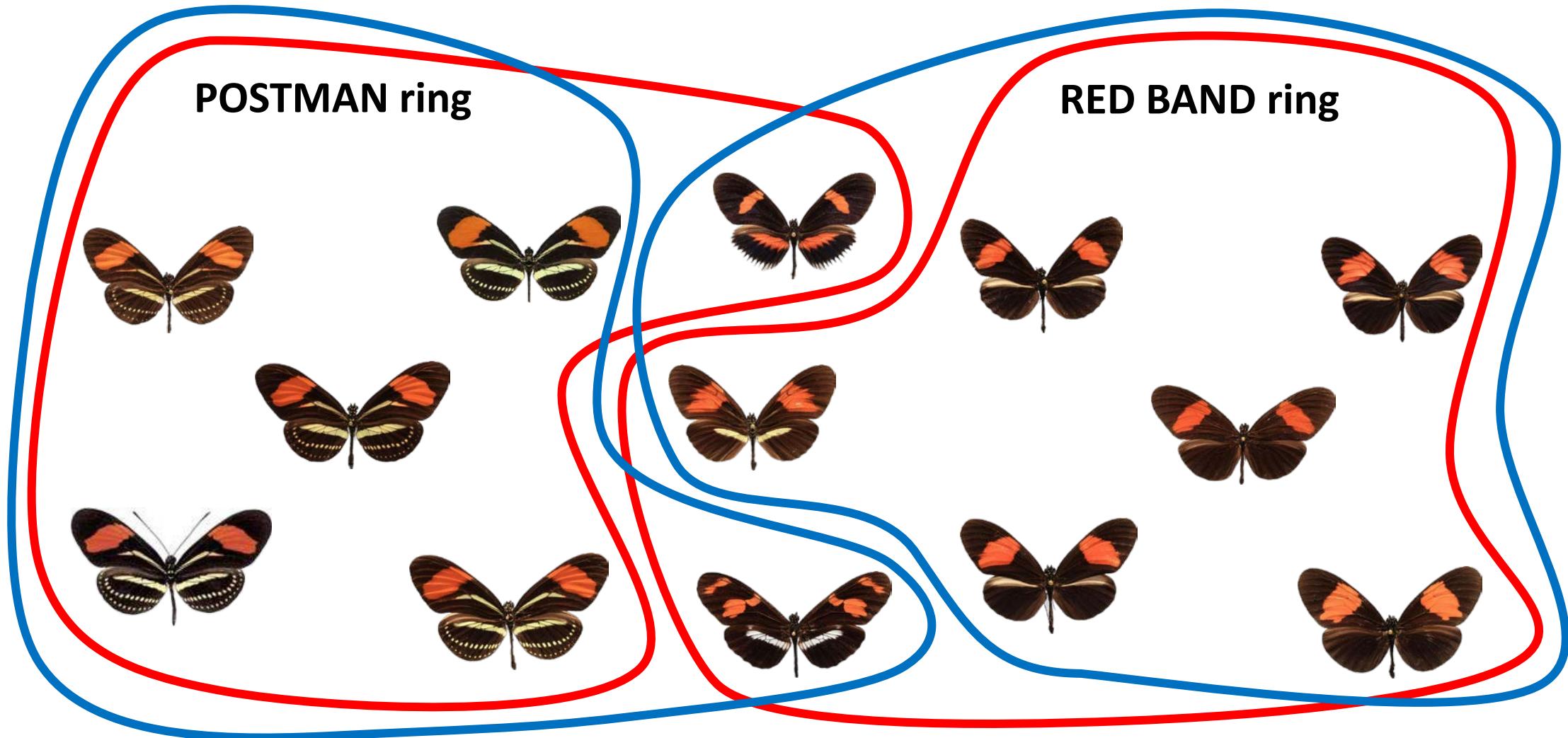
How to delineate mimicry rings?



Article A

Photo credits: C. Jiggins

How to delineate mimicry rings?



Article A

Article B

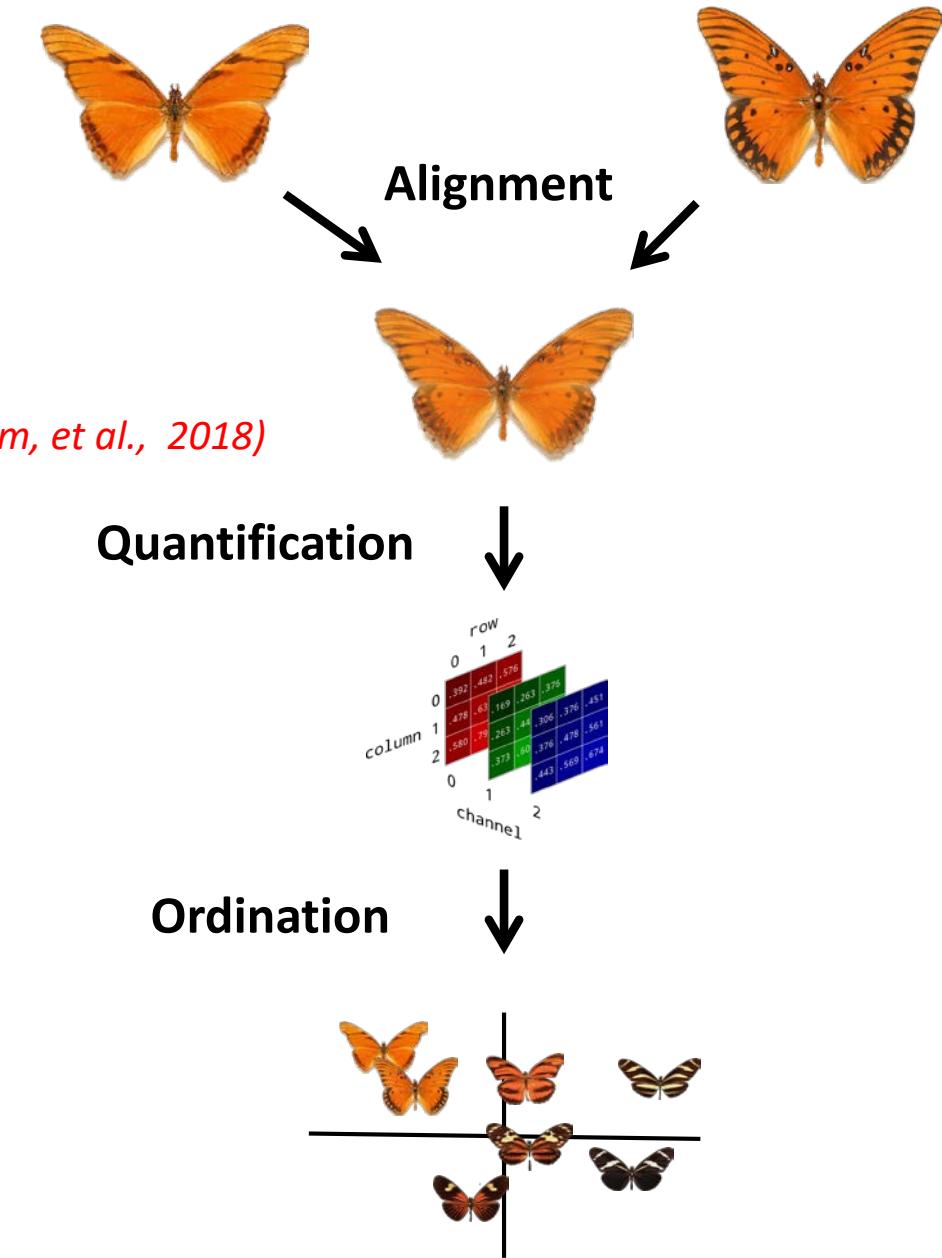
Photo credits: C. Jiggins

How to delineate mimicry rings?

Quantitative approach:

- Quantify visual signal => **patternize** *(Van Belleghem, et al., 2018)*
- Issues : alignment of pattern and outlines in case of high diversity

Photonic signal \neq Perception of this signal

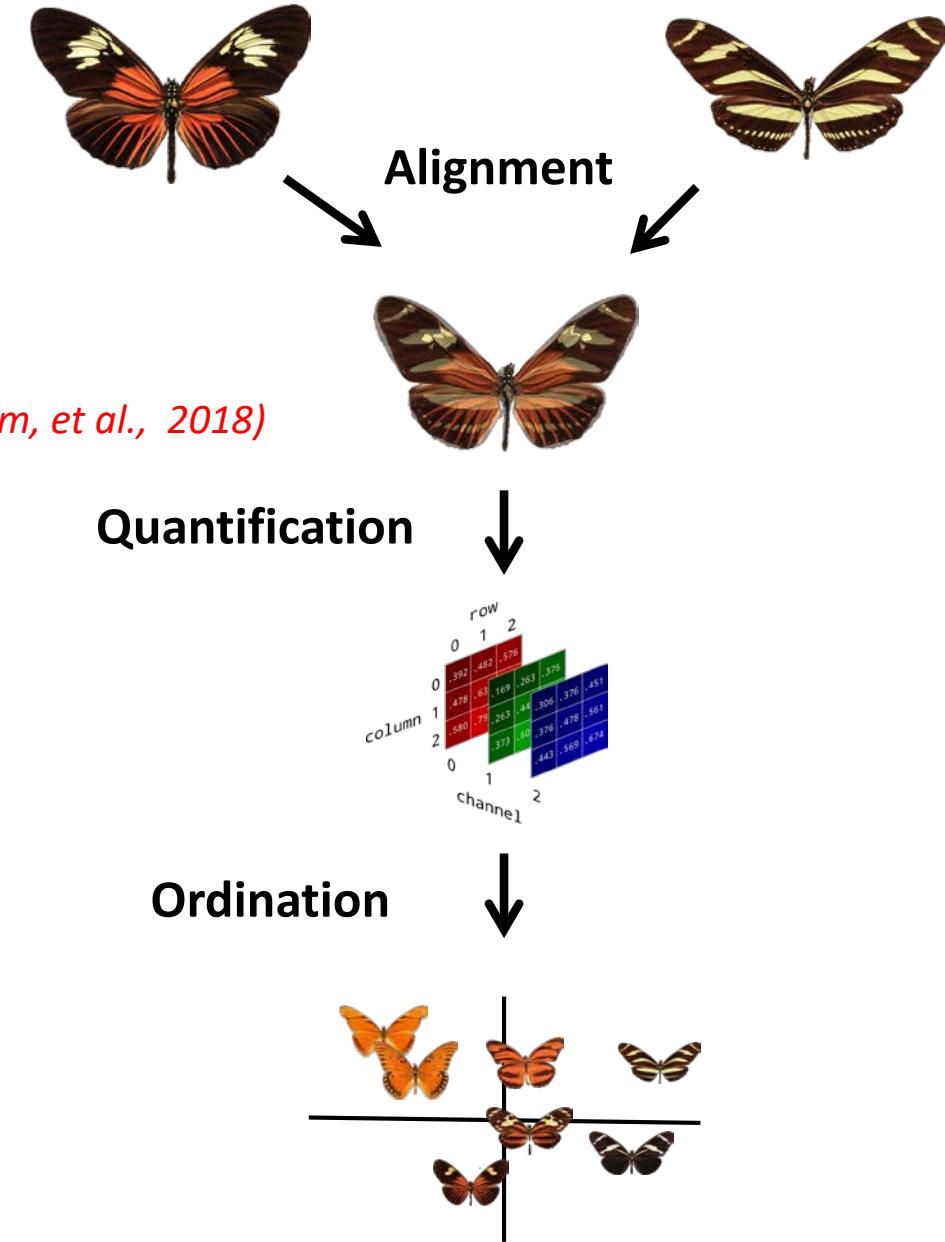


How to delineate mimicry rings?

Quantitative approach:

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How to delineate mimicry rings?

Quantitative approach:

- Quantify **visual signal** => **patternize**
- Issues : **alignment** of pattern and outlines in case of high diversity

Photonic signal ≠ Perception of this signal



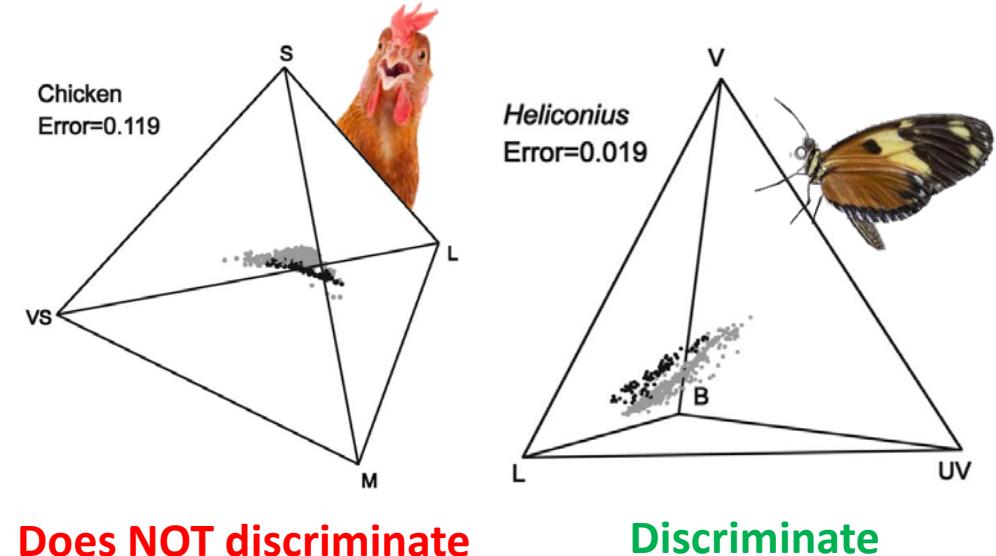
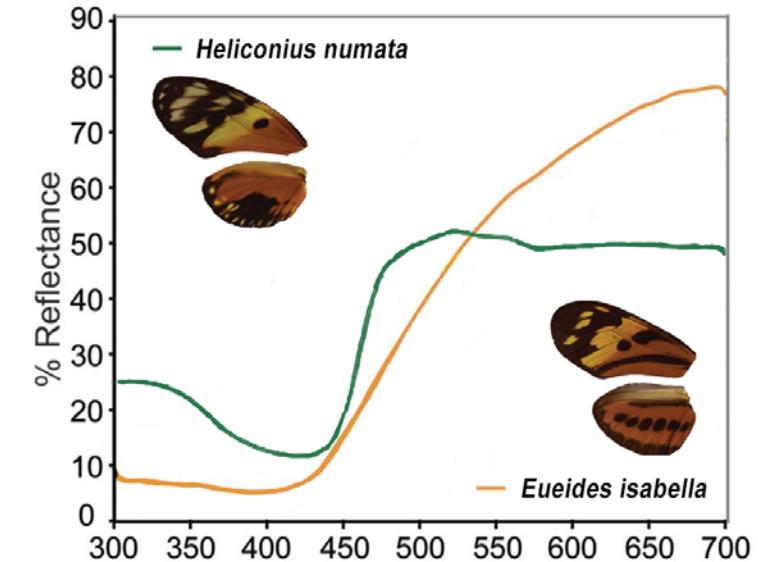
Photo credits: C. Jiggins

How to delineate mimicry rings?

Quantitative approach:

- Quantify **visual signal** => **patternize**
- Issues : **alignment** of pattern and outlines in case of high diversity

Photonic signal \neq Perception of this signal



Adapted from Bybee *et al.* 2012

Perceptual approach

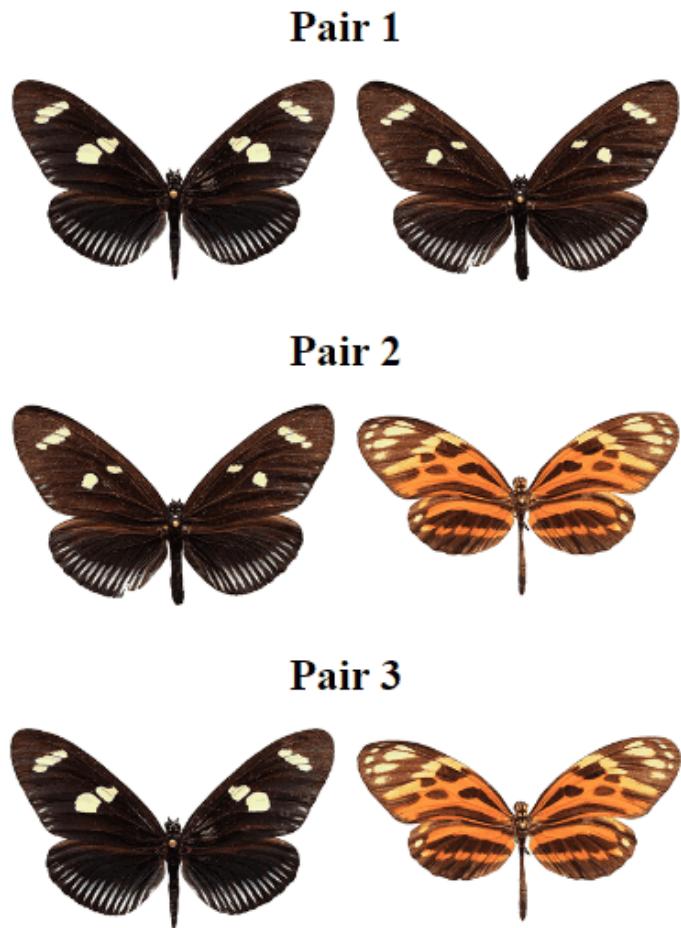
**Online survey to evaluate perception
of similarity across image triplets :**

<http://memometric.cleverapps.io/>

Input: triplet of pictures with a **reference**

Output: list of triplets with **relative distances**

Quality control: 3 control triplets for 30 items

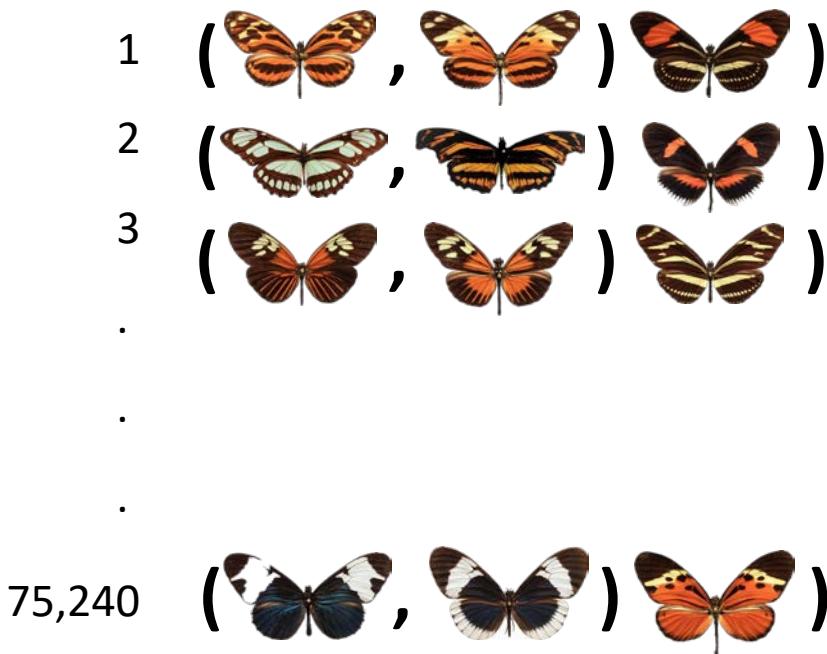


Skip
(I cannot decide)

Embedding method: t-STE

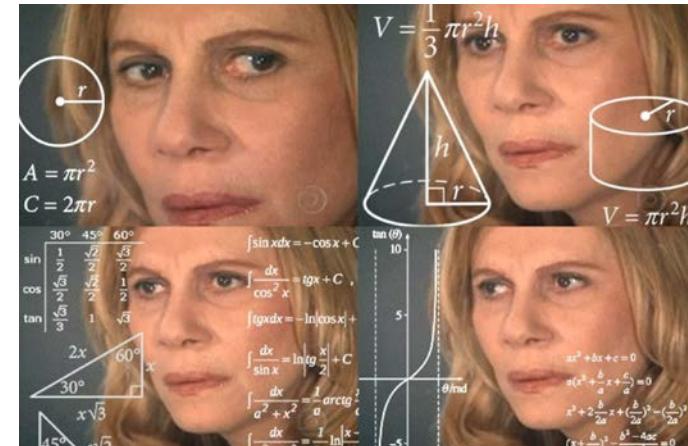
Goal = obtain a **perceptual map** that reflect the **perceptual distances** between images

Input = **relative triplet distances**



Embedding method:
Machine learning algorithm
(t-STE)

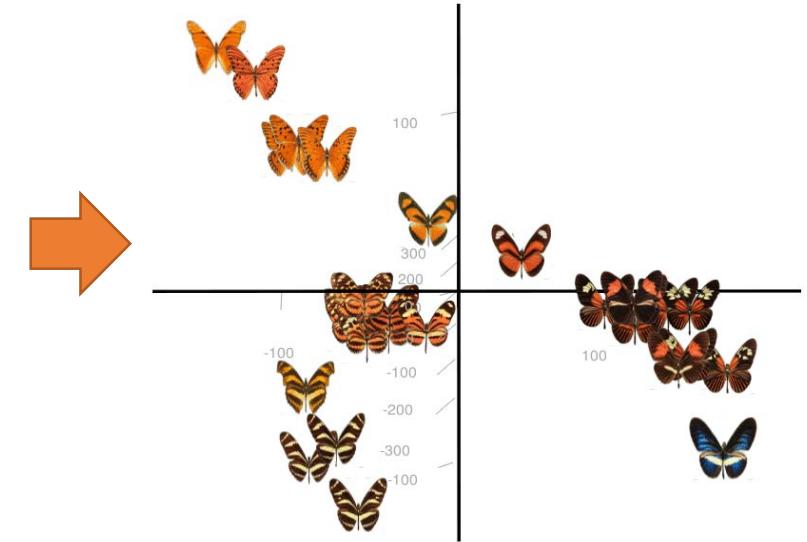
(van der Maaten & Weinberger, 2012)



Output = **coordinates** in space
of reduced dimensionality

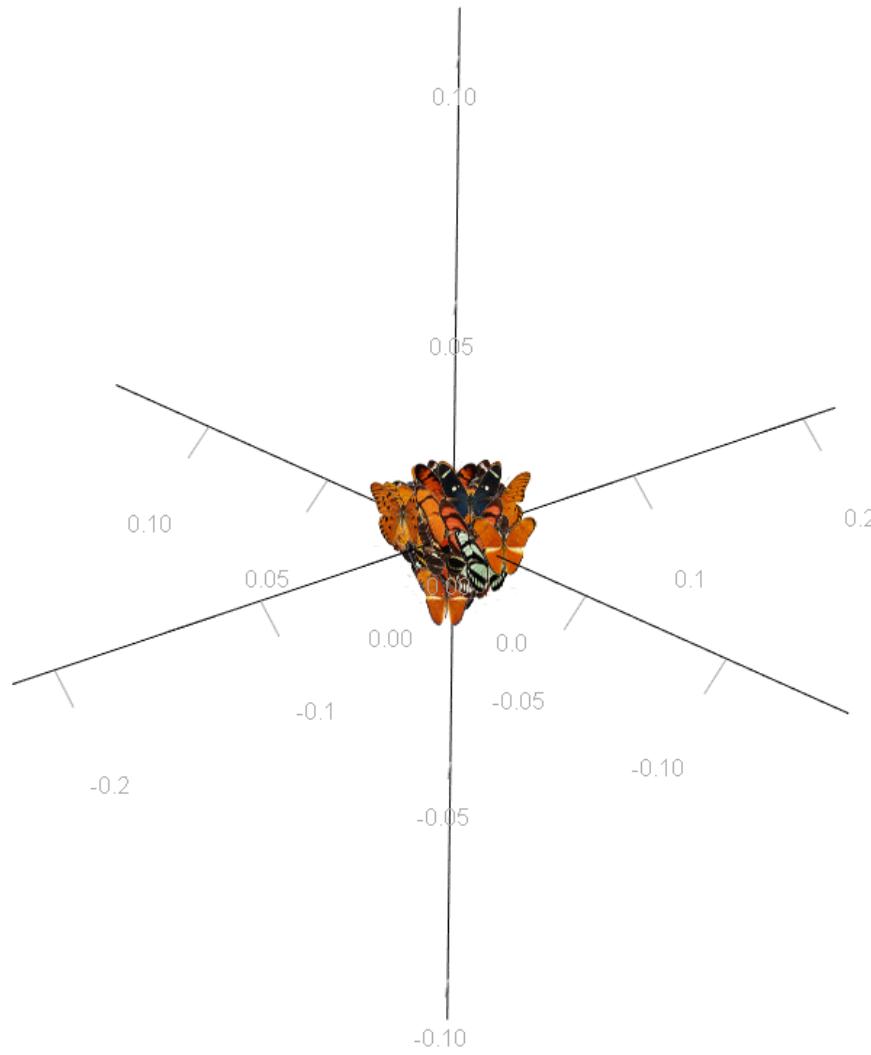
=

Perceptual map

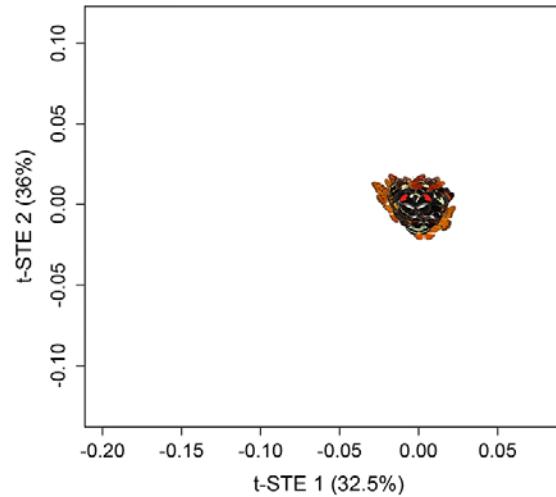


Embedding method: t-STE

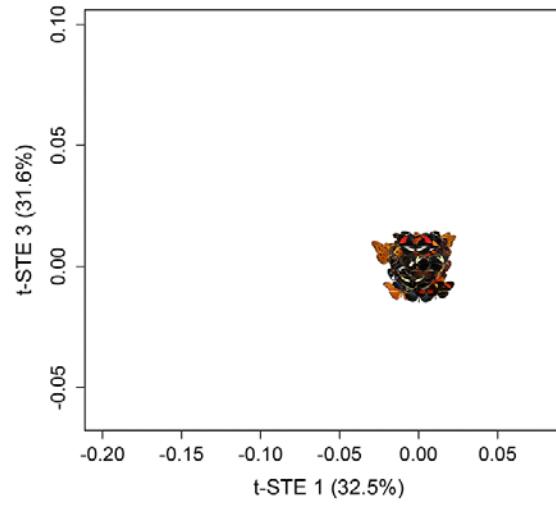
Iteration = 0



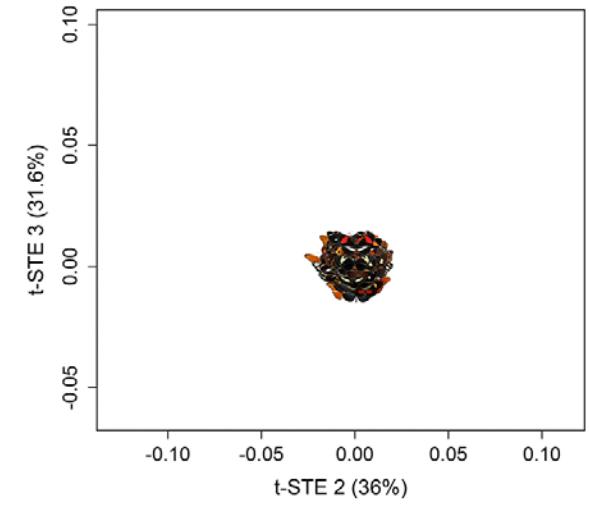
Map for 432 images
Learning step: 0
t-STE - X with Lambda = 0 in 3D



Map for 432 images
Learning step: 0
t-STE - X with Lambda = 0 in 3D



Map for 432 images
Learning step: 0
t-STE - X with Lambda = 0 in 3D



Perceptual space



75,240 triplets for 432 images

The spatial scale of mimicry rings

Why do we have a **continuum**?

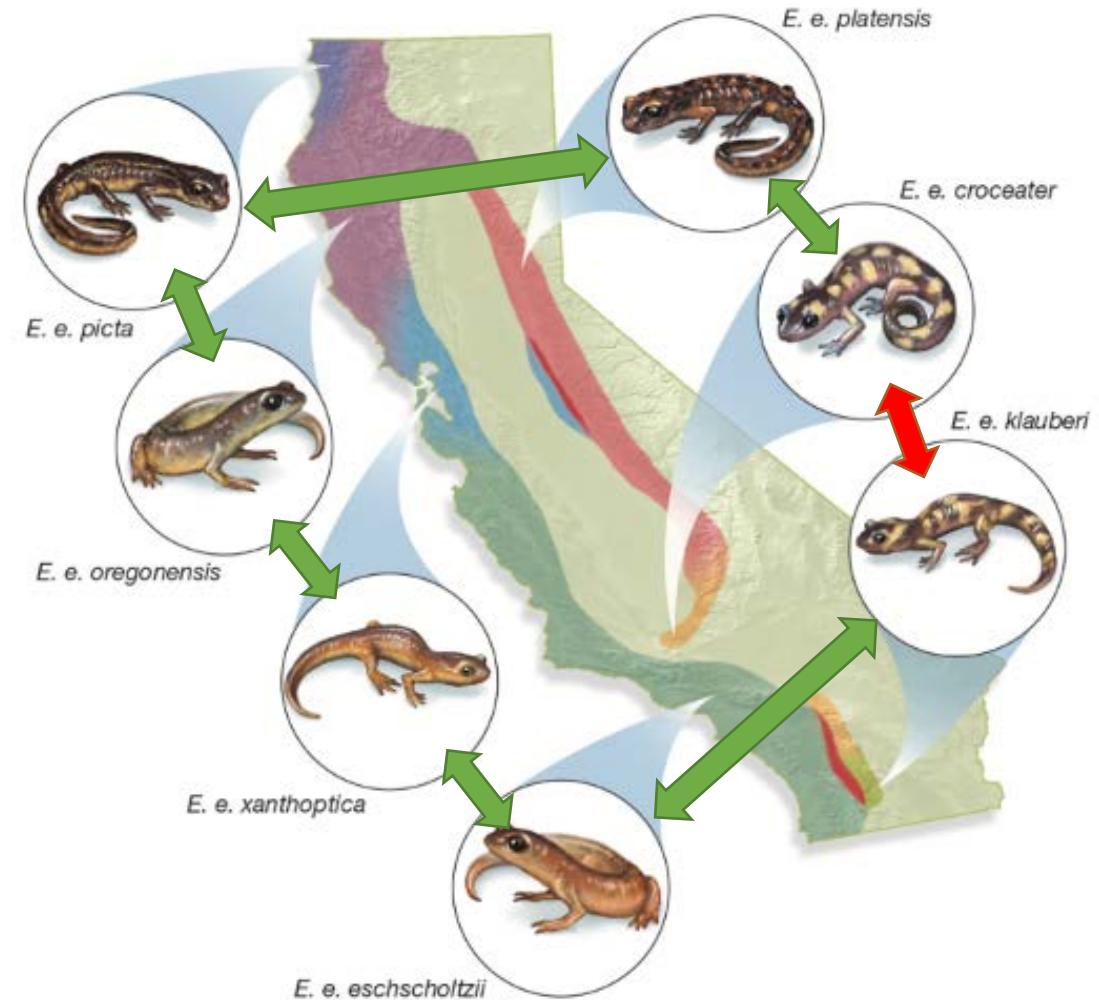
Analogy with **ring species**

- ↔ - **Hybridization** at local scale
- ↔ - **Incompatibility** at large distances
- **Continuum** at large scale

Conclusion:

- Expected to find a **continuum** at the **global scale**
- Mimicry rings should be defined at **community-level**?

Next step: Apply the perceptual map approach at **community-level**



Source: Chegg.com

The spatial scale of mimicry rings

Why do we have a **continuum**?

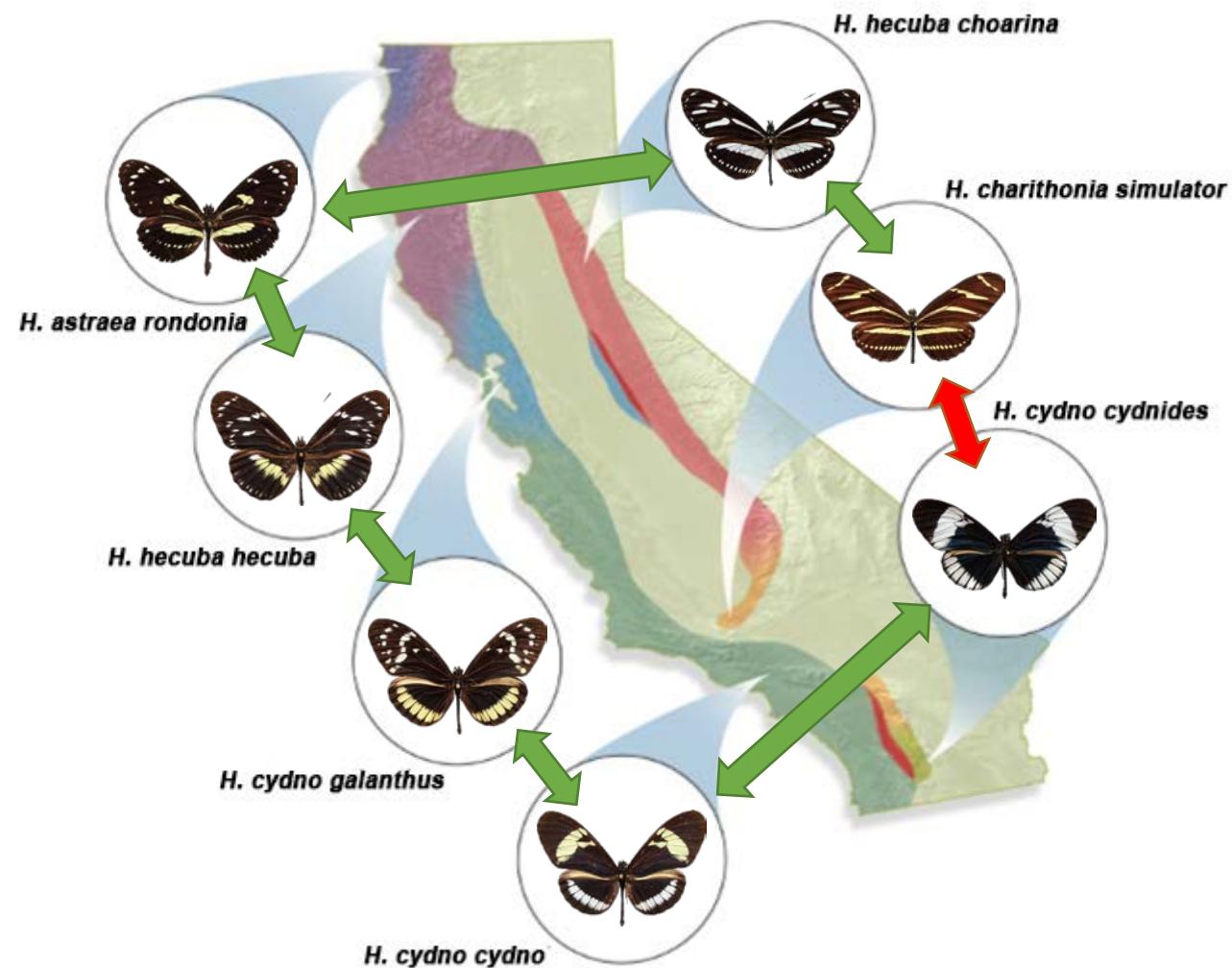
Analogy with **ring species**

- ↔ - **Mimicry** at local scale
- ↔ - **Dissimilarity** at large distances
- **Continuum** at large scale

Conclusion:

- Expected to find a **continuum** at the **global scale**
- Mimicry rings should be defined at **community-level**?

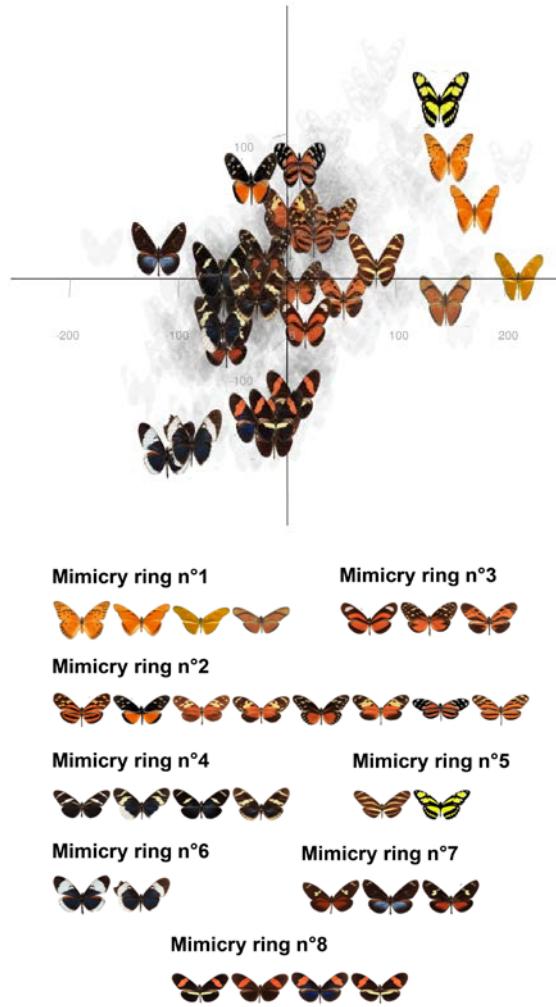
Next step: Apply the perceptual map approach at **community-level**



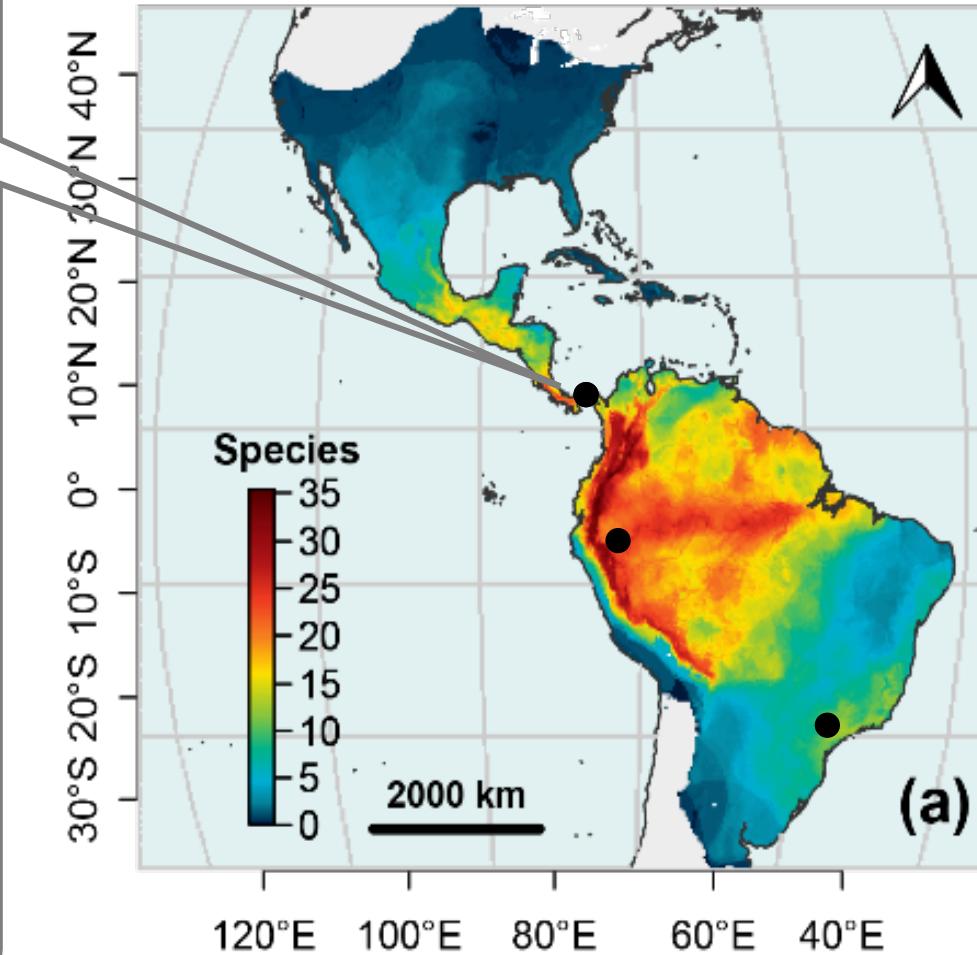
Source: Chegg.com

Interactive map of Müllerian mimicry communities

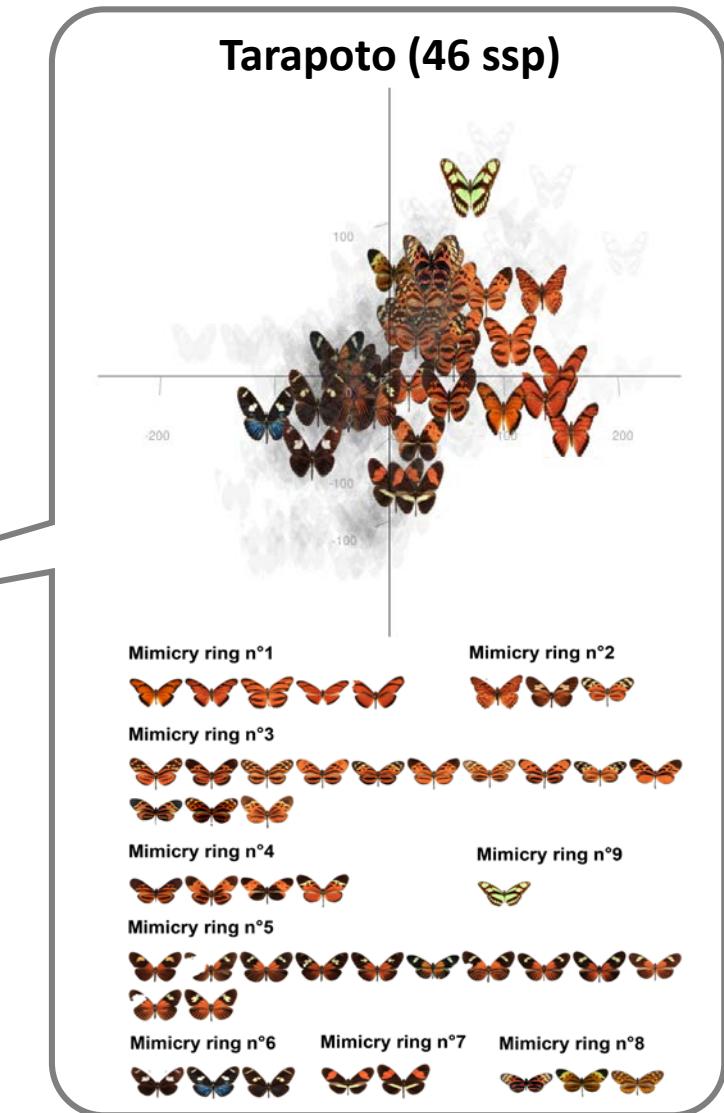
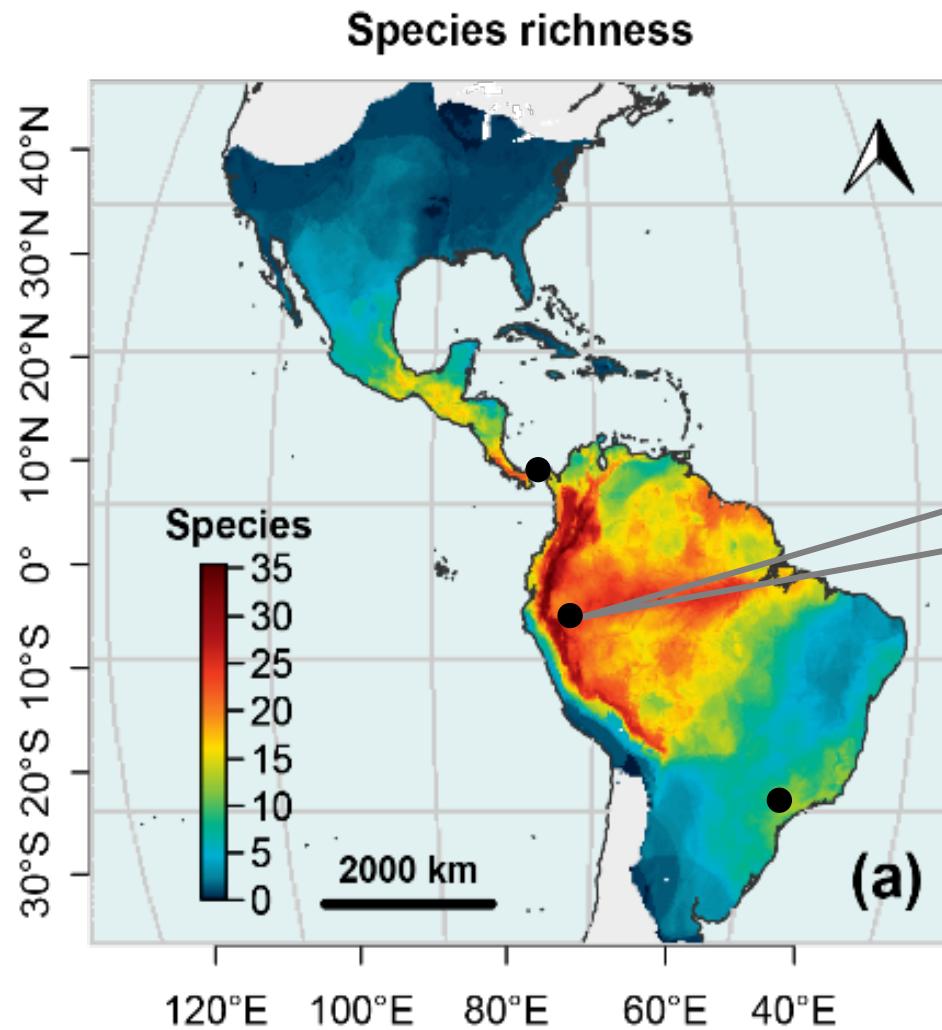
Gamboa (30 ssp)



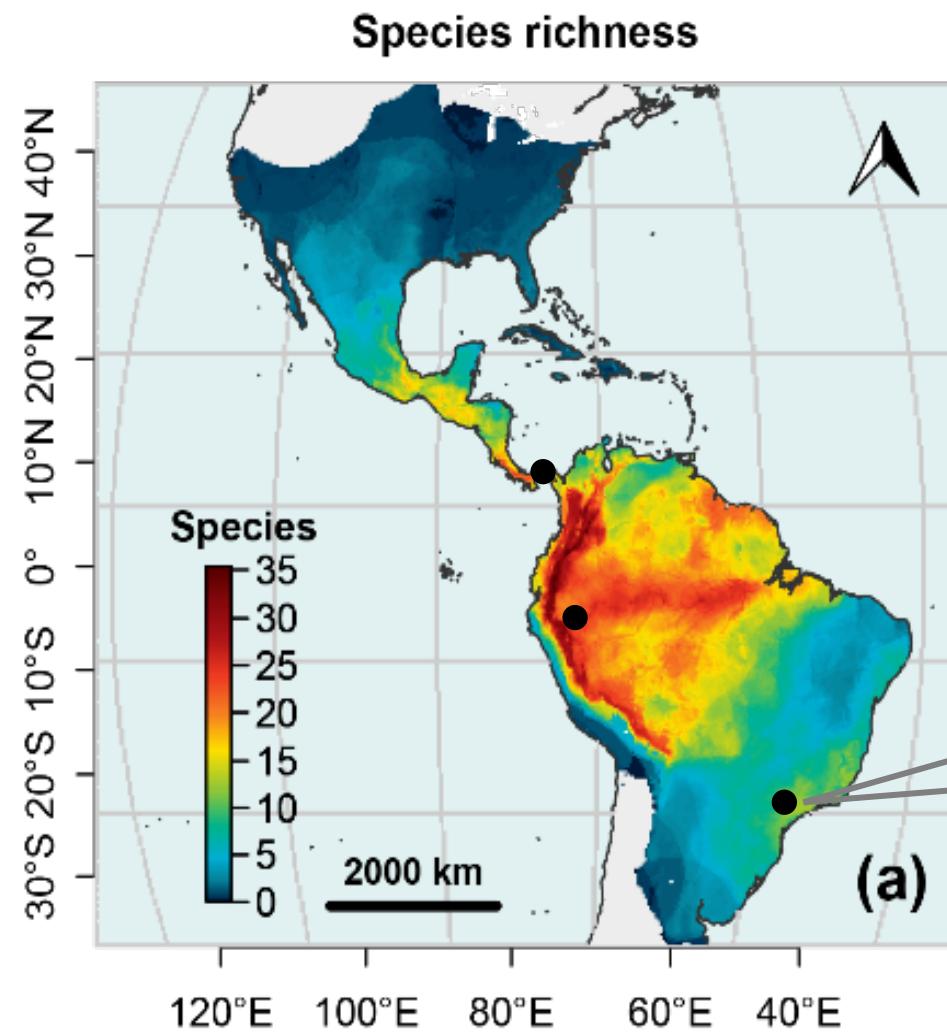
Species richness



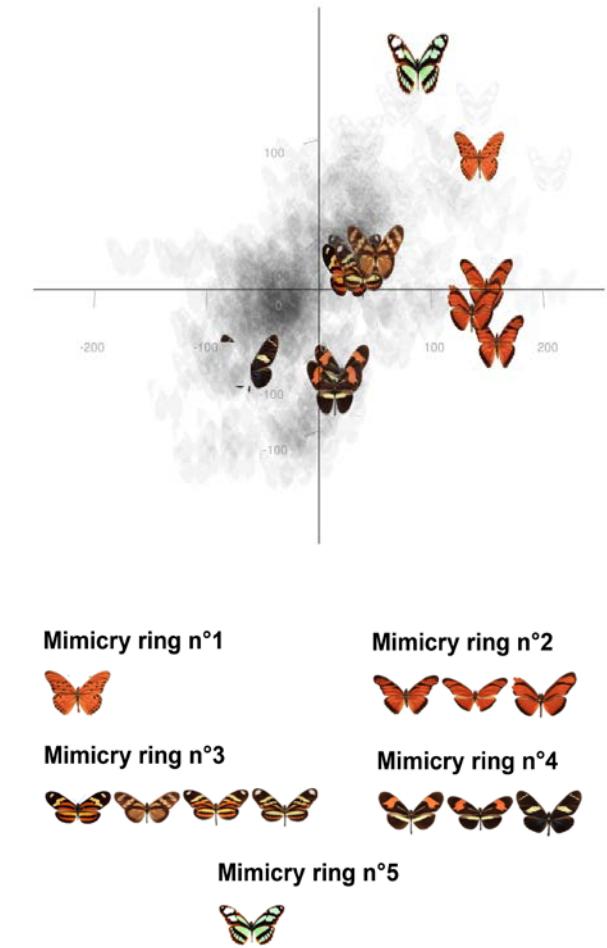
Interactive map of Müllerian mimicry communities



Interactive map of Müllerian mimicry communities



Campinas (12 ssp)



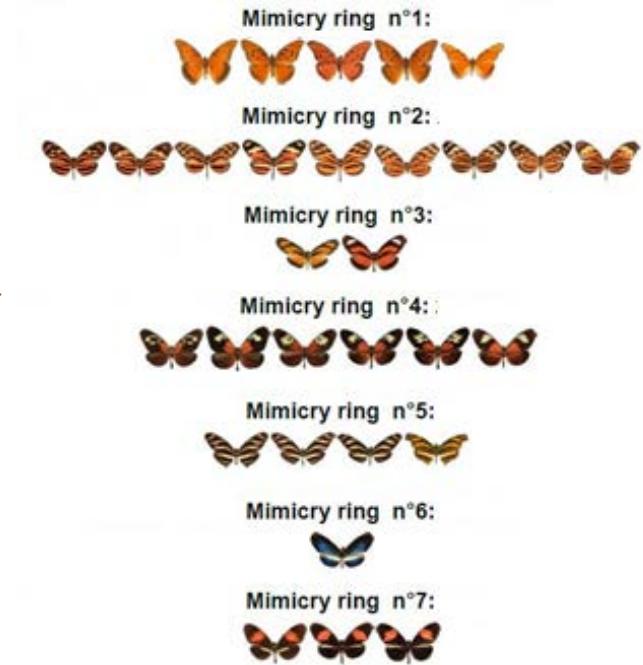
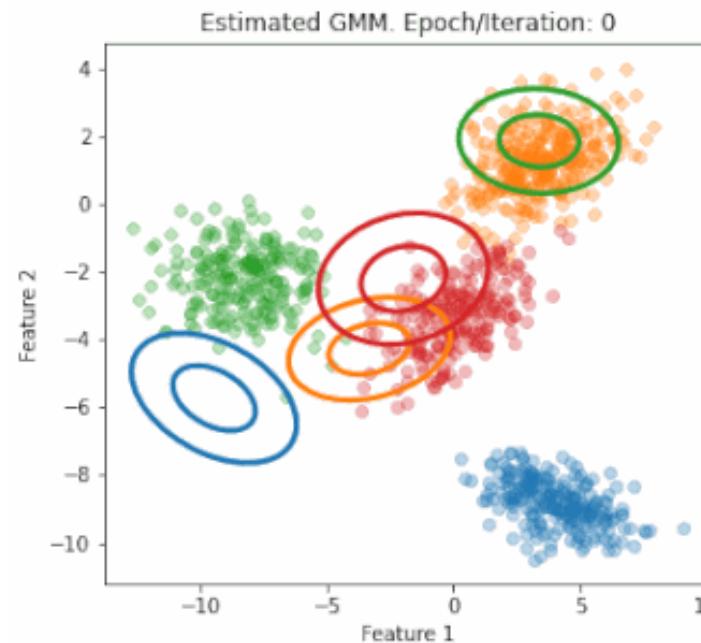
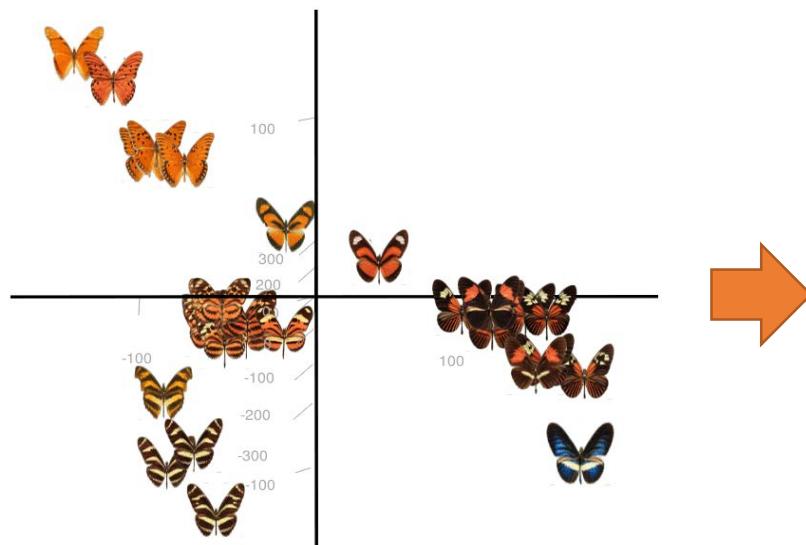
Clustering method

Goal = delineate groups of patterns (**mimicry rings**) in the **local perceptual space**

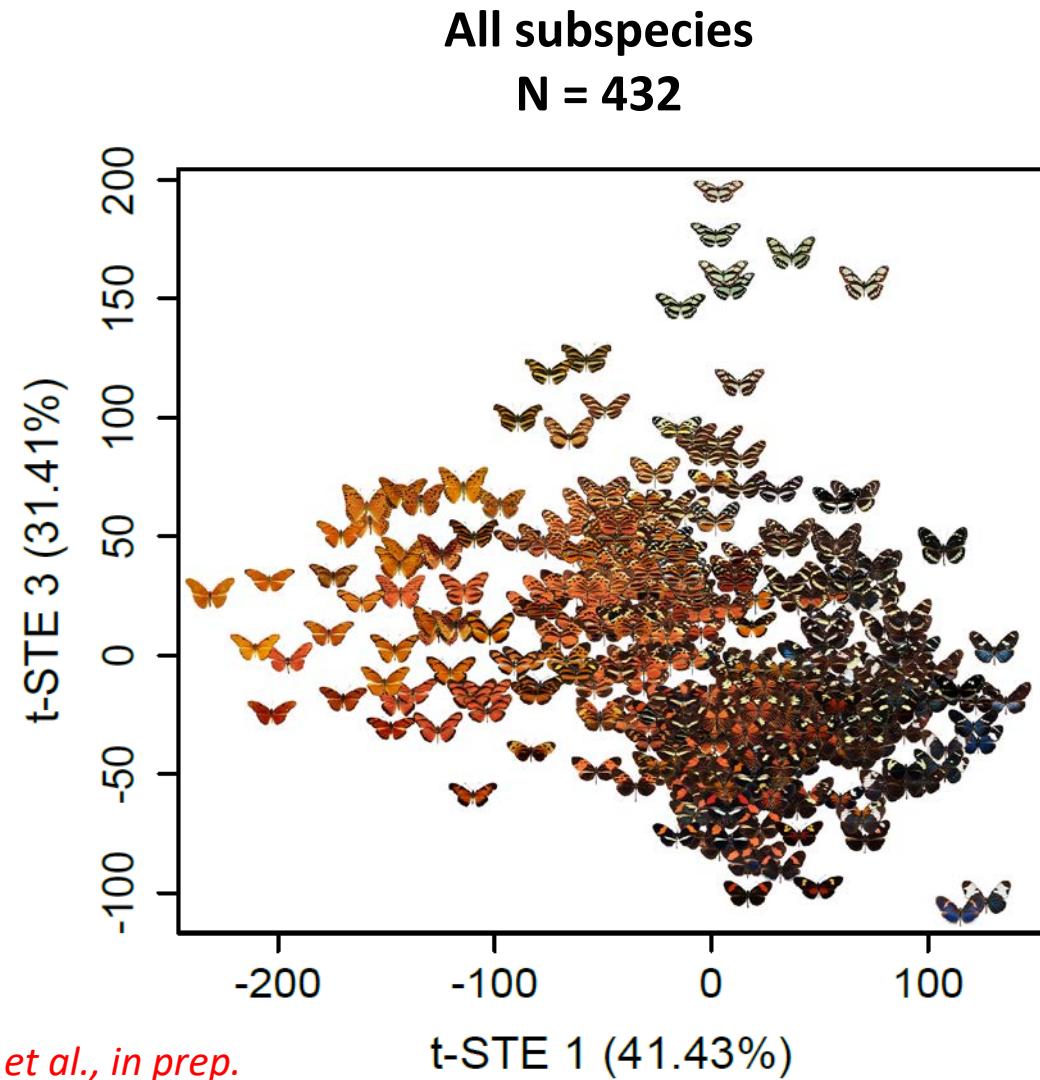
Input = **coordinates**
on the perceptual map

Clustering method:
Gaussian Mixture Models
(GMM)

Output = **clusters**
of wing patterns
=
Mimicry rings



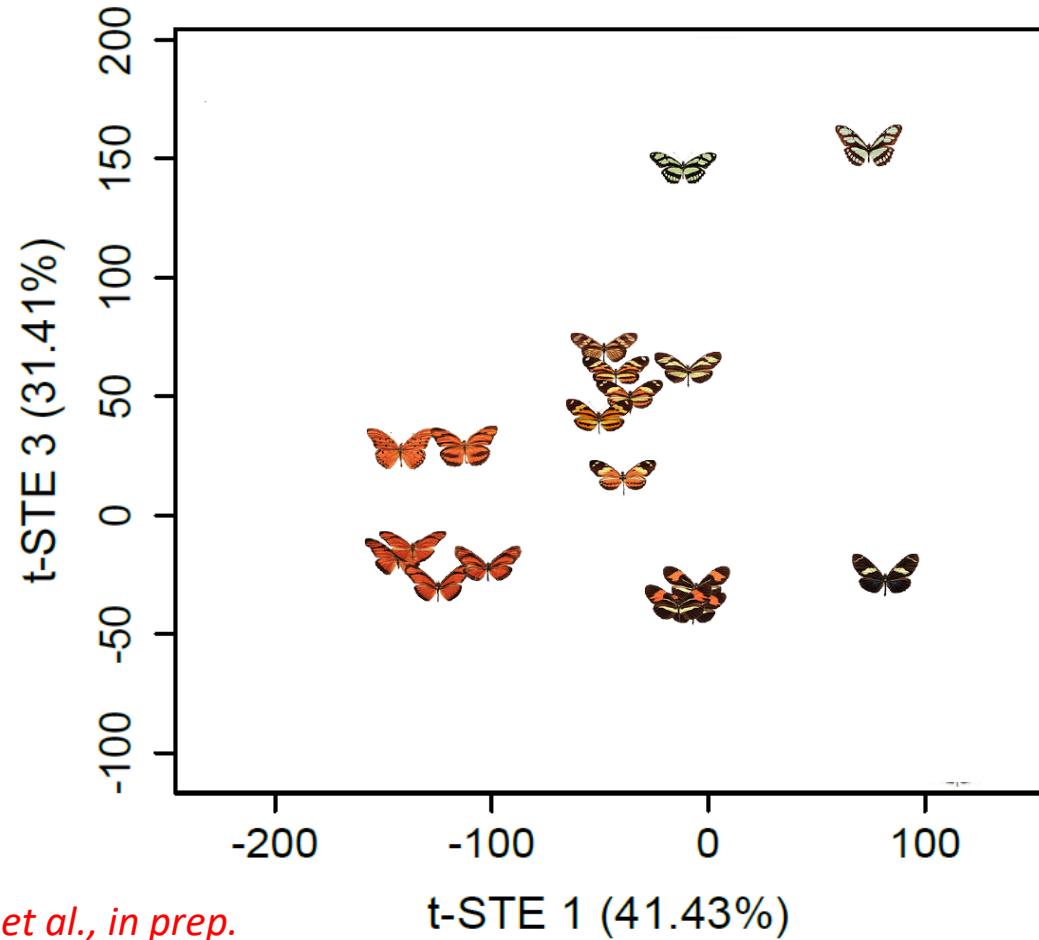
Local perceptual maps



Local perceptual maps

Santa Teresa, ES, Brazil

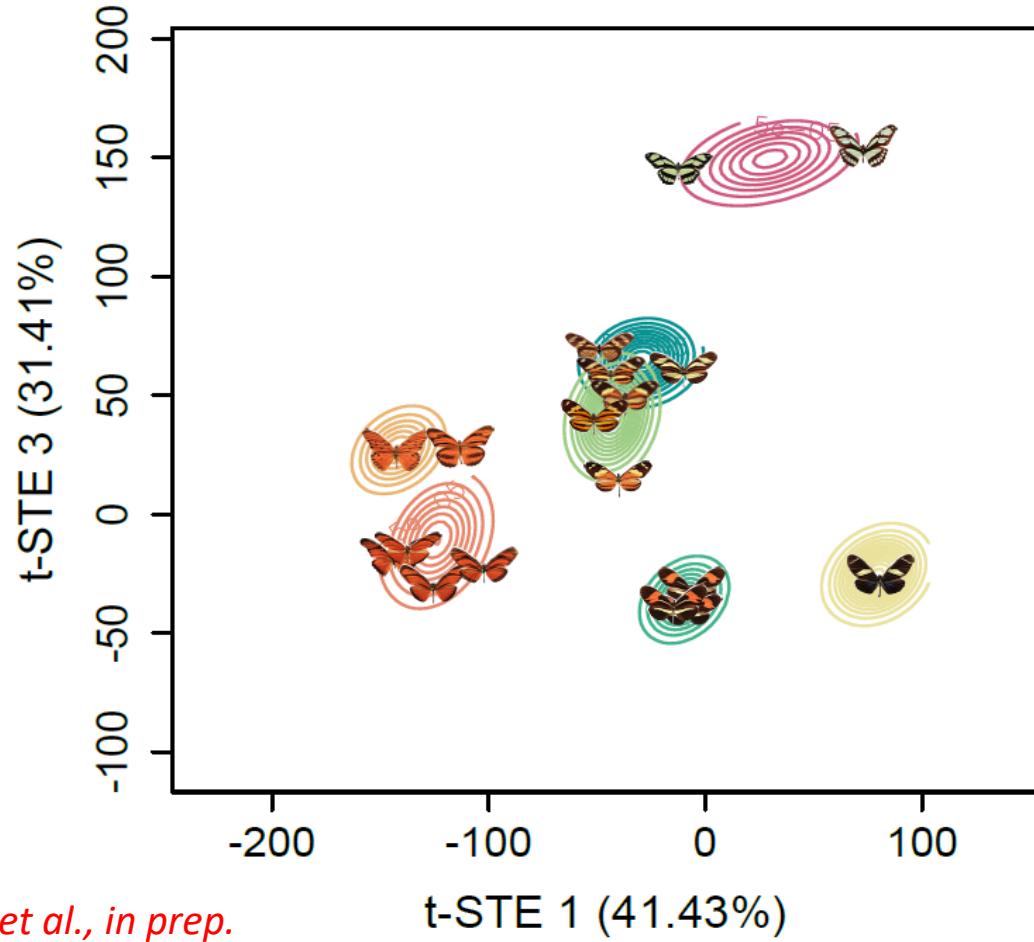
$N = 18 ; k = 7$



Doré et al., in prep.

Local perceptual maps

Santa Teresa, ES, Brazil
 $N = 18 ; k = 7$



Doré et al., in prep.

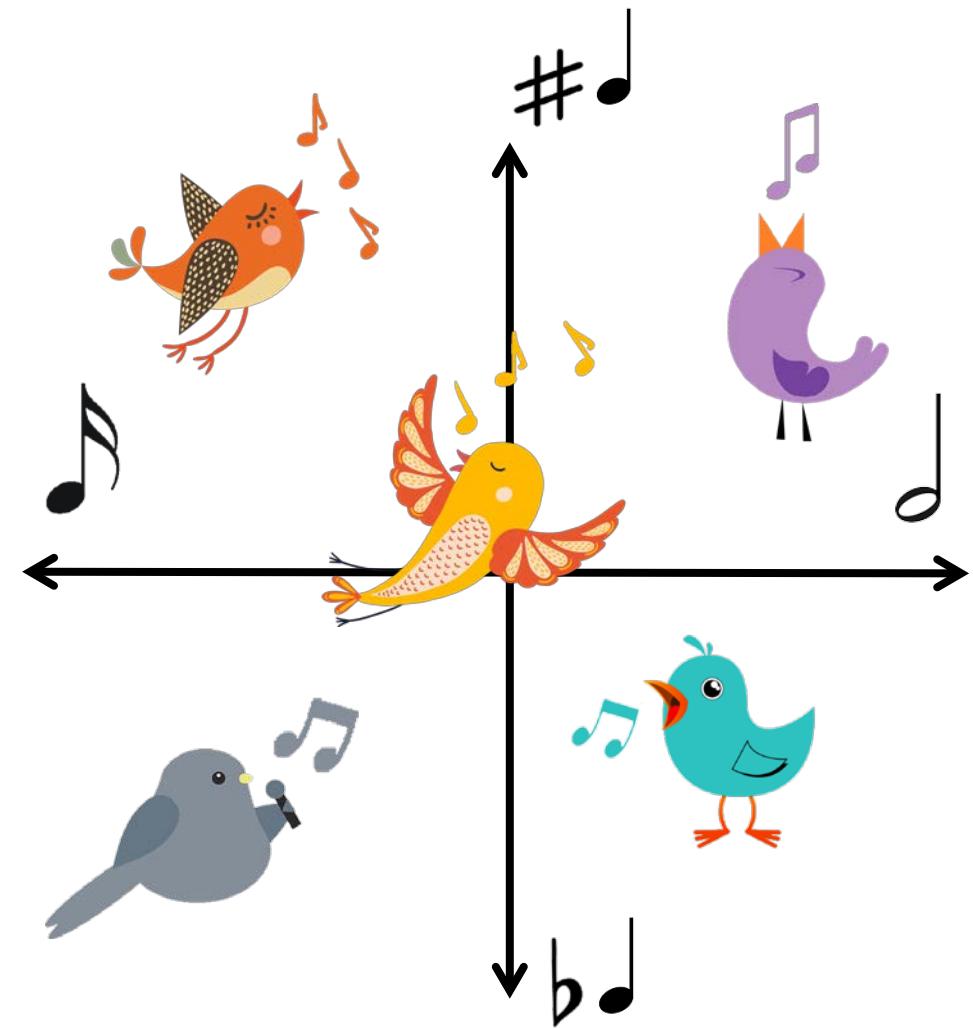


Conclusion

Perceptual space = a new tool to quantify any **ecological signals** through perception: colors, shapes, songs, smells, behaviors, etc.

Allow to explore phenotypic patterns at large spatial and taxonomic scale in context of high phenotypic diversity

Study differences in perception across individuals/social groups



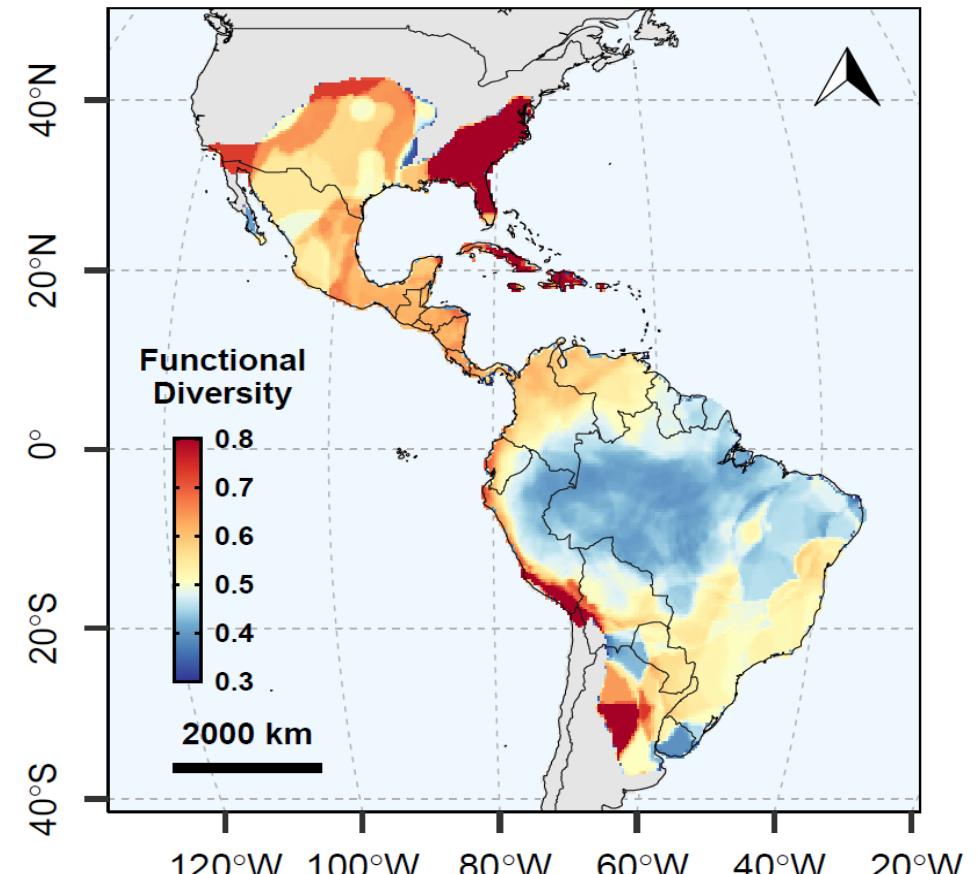
Perceptual space of bird songs

Conclusion

Perceptual space = a new tool to quantify any **ecological signals** through perception: colors, shapes, songs, smells, behaviors, etc.

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Doré et al., in prep.

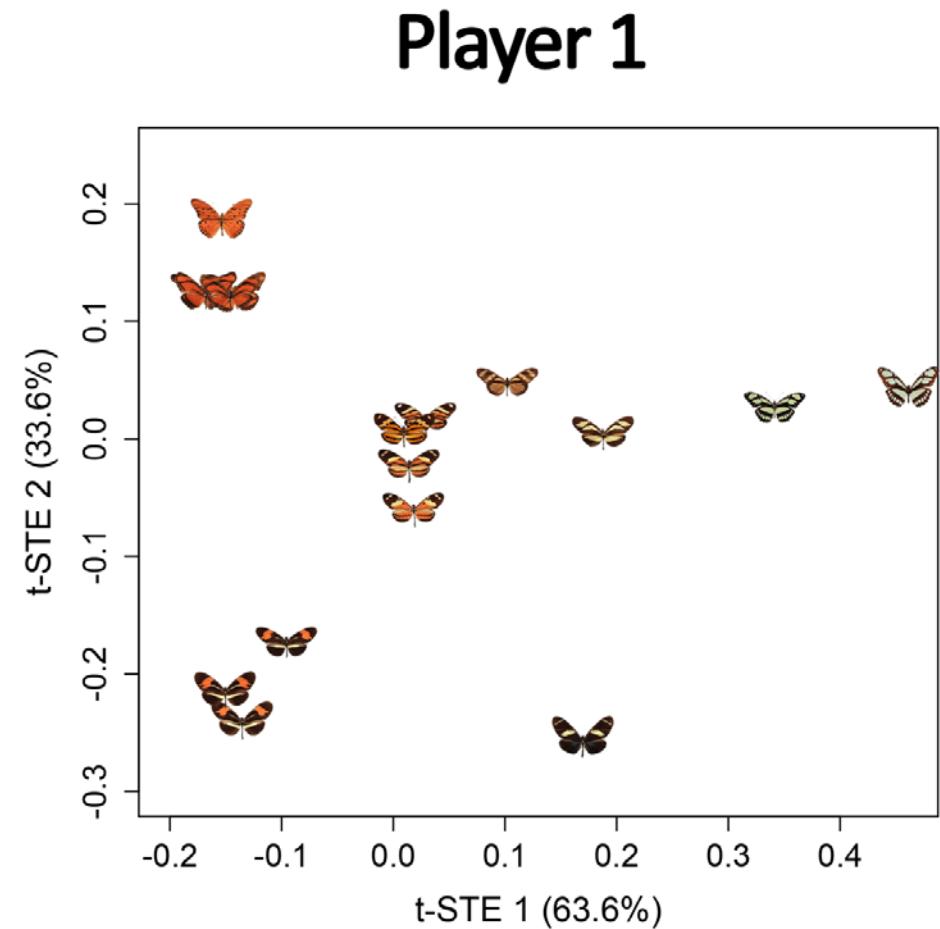
Mimetic diversity as clustering in the local perceptual space

Conclusion

Perceptual space = a new tool to quantify any **ecological signals** through perception: colors, shapes, songs, smells, behaviors, etc.

Allow to explore phenotypic patterns at **large spatial and taxonomic scale** in context of **high phenotypic diversity**

Study **differences in perception** across individuals/social groups



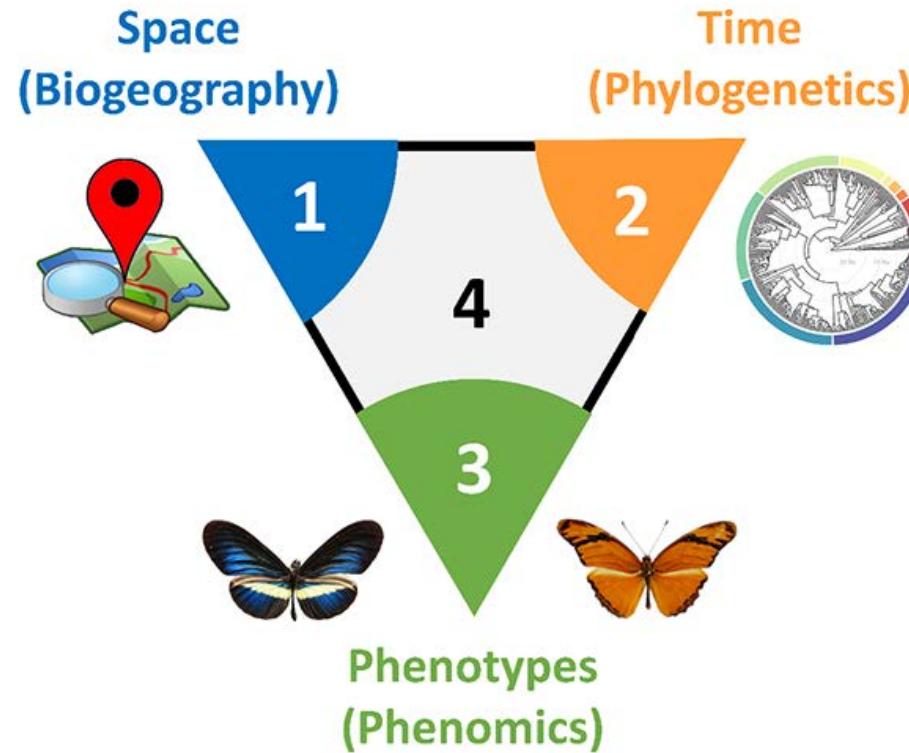
Outlines and objectives

CHAPTER 1

Map biodiversity patterns

CHAPTER 3

Quantify phenotypic similarity in wing patterns

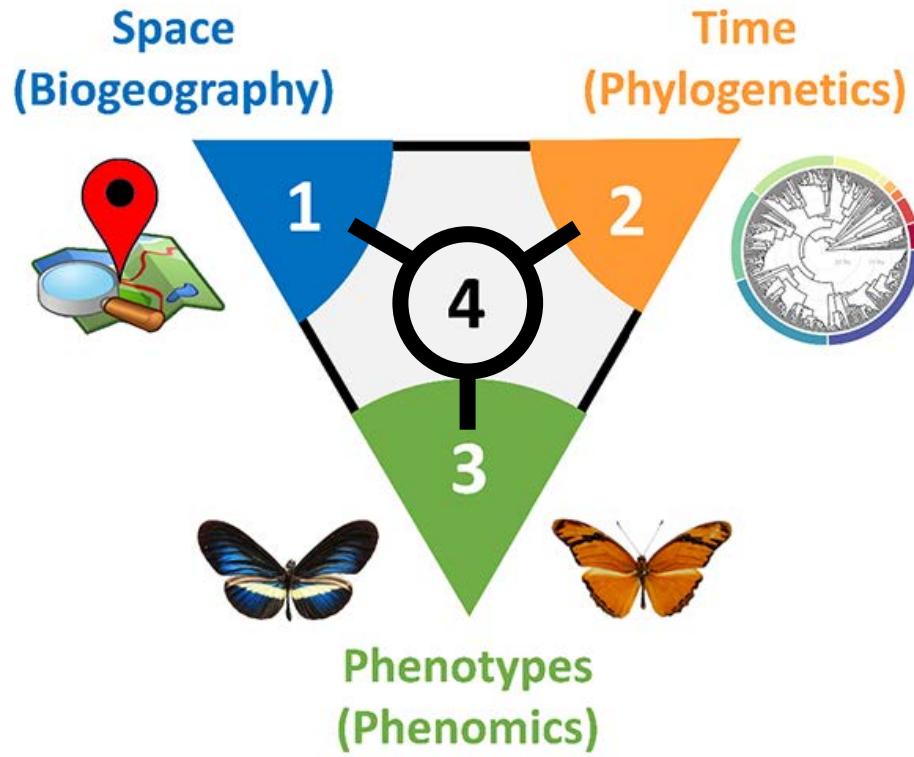


CHAPTER 2

Resolve deep evolutionary relationships

How mutualistic interactions affect the structure and evolution of biodiversity at the macroecological scale?

CHAPTER 4: Mutualistic interactions shape global spatial congruence and climatic niche evolution in Neotropical mimetic butterflies



Reference:

Doré, M., Willmott, K., Lavergne, S., Chazot, N., Freitas, A. V. L., Fontaine, C. & Elias, M. (2023). Mutualistic interactions shape global spatial congruence and climatic niche evolution in Neotropical mimetic butterflies. *Ecology Letters*. In production. <https://doi.org/10.1111/ele.14198>

Questions & Hypotheses

How **mutualistic interactions** affect the **structure** and **evolution** of biodiversity at the **macroecological scale**?

Spatial pattern

Promote the large-scale **cooccurrence** of mutualistic species

Niche evolution

Drive the **convergence** of the niche of mutualistic species

Study system: Neotropical butterflies

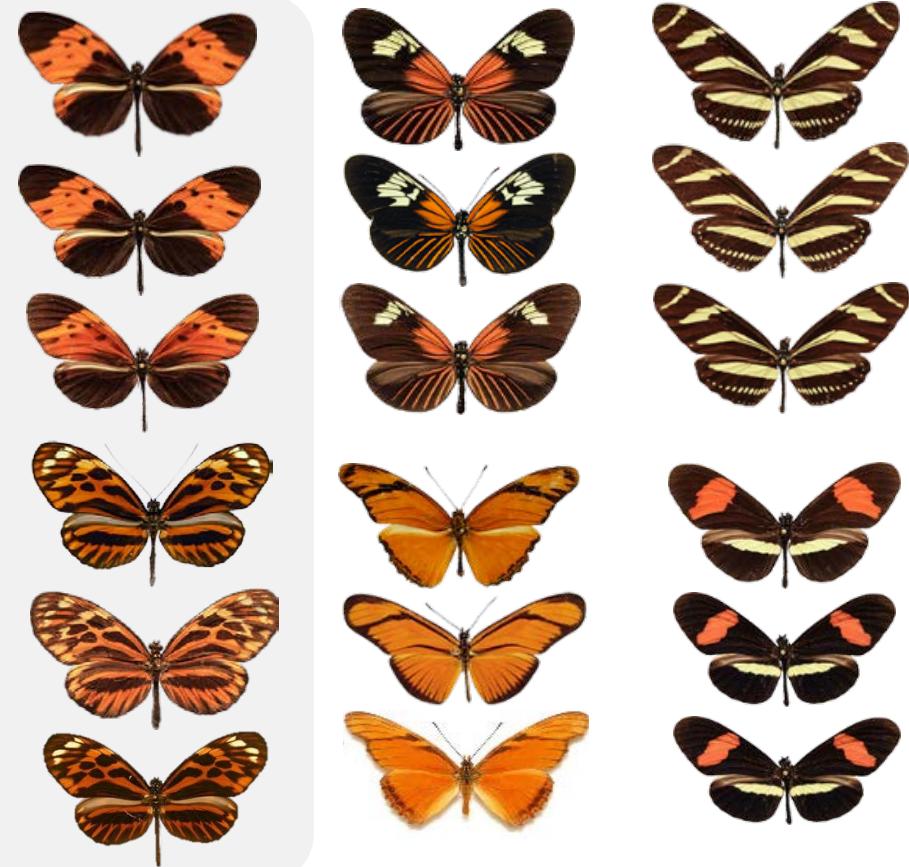
Credits Photo: N. Chazot

Ithomiini tribe



44 phenotypic groups / putative mimicry rings

Heliconiini tribe



39 phenotypic groups / putative mimicry rings

Credit photos: C. Jiggins

8 shared
patterns

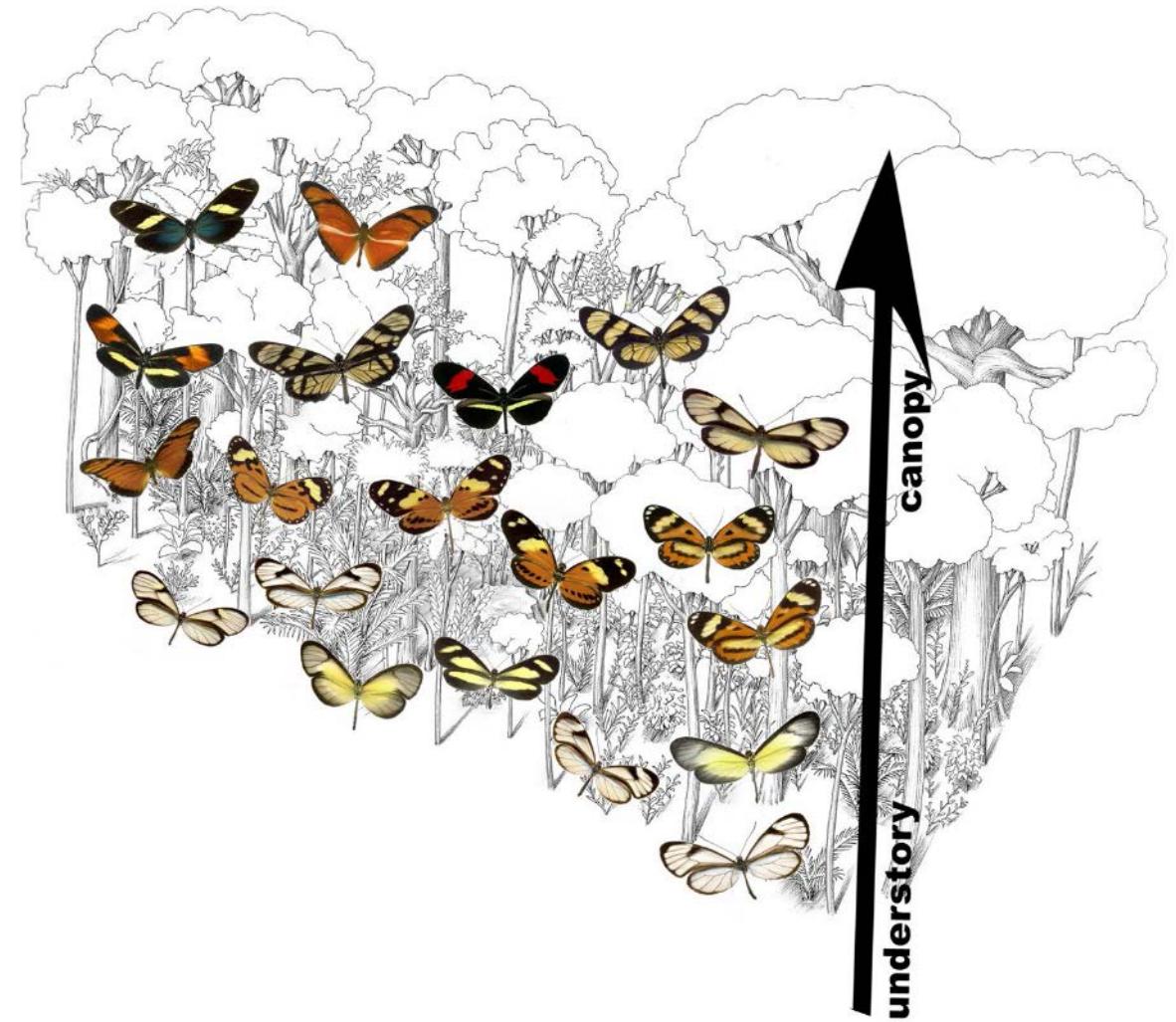
State-of-the-art

Structuration of mimetic communities by:

- **flight height** (*Beccaloni, 1997*)
- **microhabitats** (*Elias et al., 2008*)
- **host plants** (*Willmott & Mallet, 2004*)
- **altitude** (*Chazot et al., 2014*)

Limits :

- **Spatial:** local to regional
- **Taxonomic:** few genera



Adapted from Birskis-Baros et al., 2021

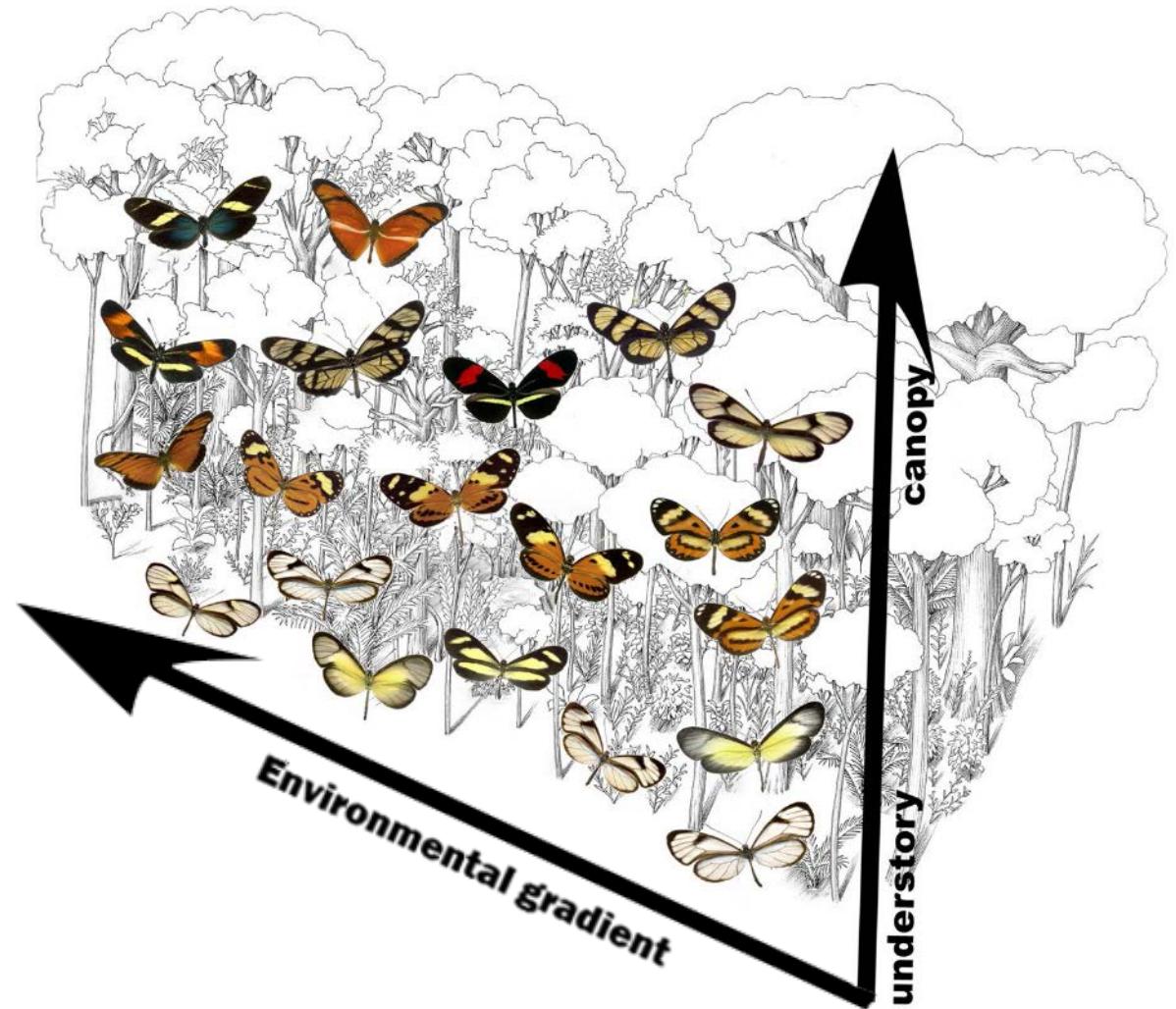
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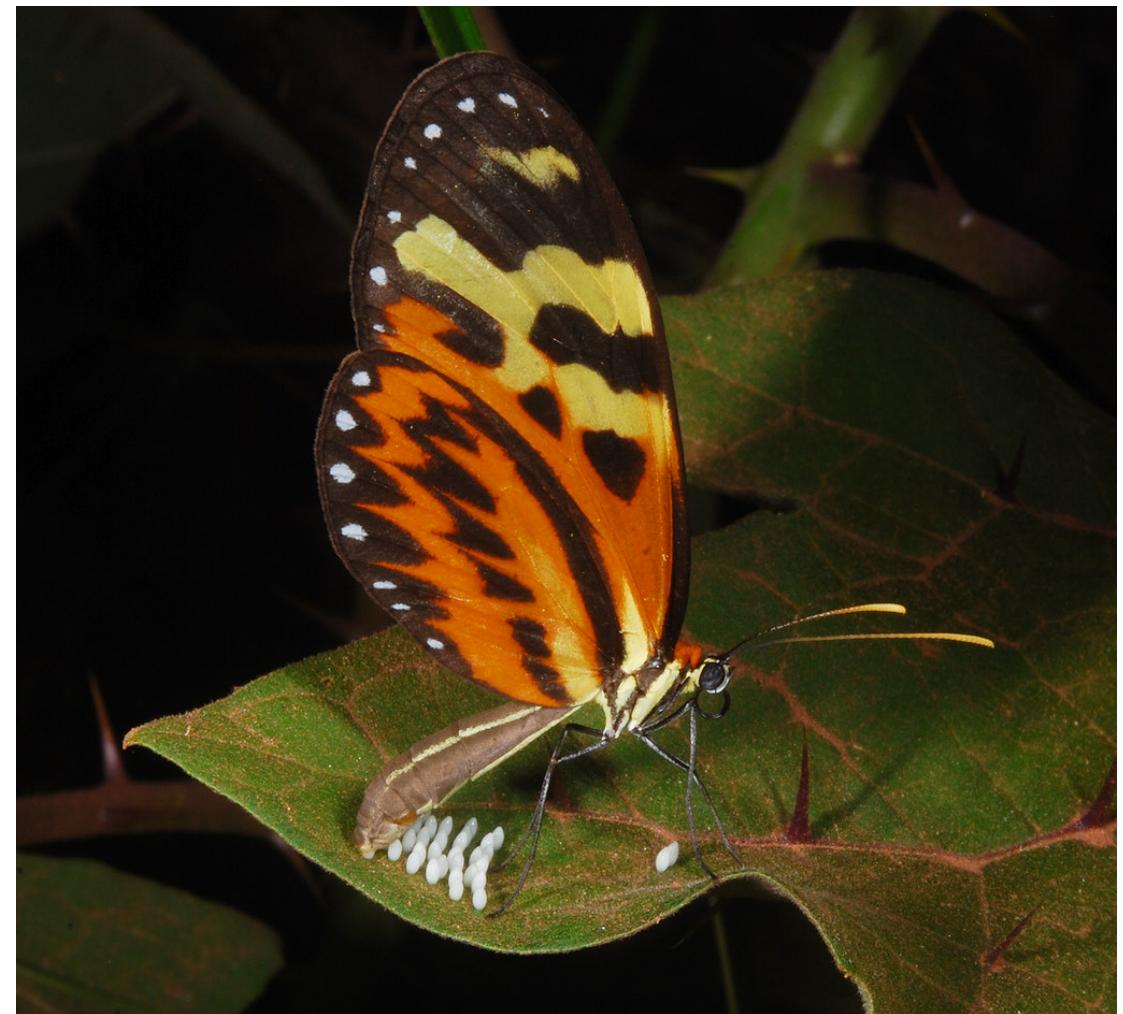
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Credits: Arthur Anker

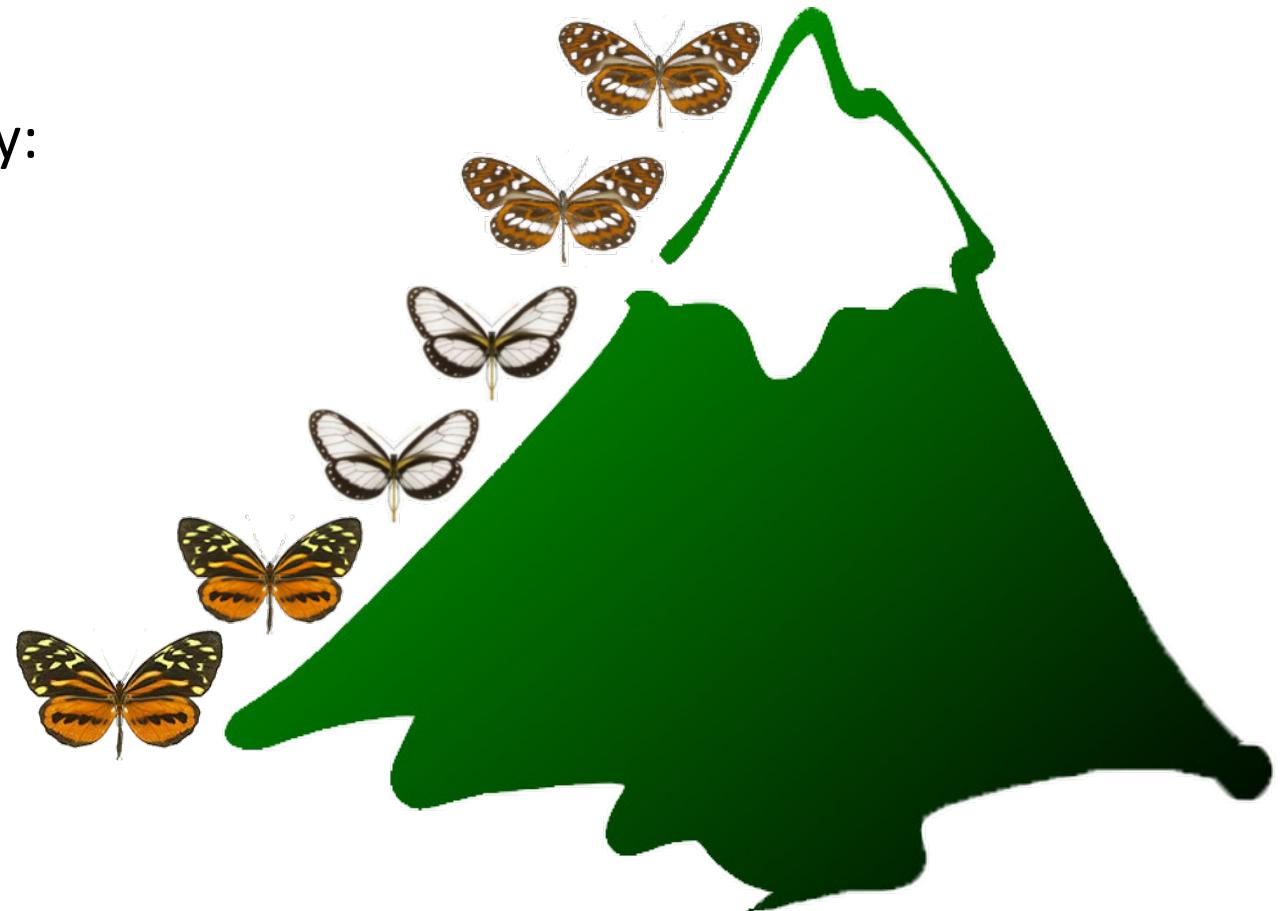
State-of-the-art

Structuration of mimetic communities by:

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- **altitude** (*Chazot et al., 2014*)

Limits :

- **Spatial**: local to regional
- **Taxonomic**: few genera



This study: **macroecological scale for the whole tribes**
Dimensions = climatic niche

Objectives

How **mutualistic interactions** affect the **structure** and **evolution** of biodiversity at the **macroecological scale**?



Spatial congruence: Do phenotypically similar species **cooccur** more than expected at random?



Niche similarity: Do phenotypically similar species have similar climatic niche?



Niche convergence: Is the climatic niche of phenotypically similar species **more similar** than expected from **shared ancestry**?



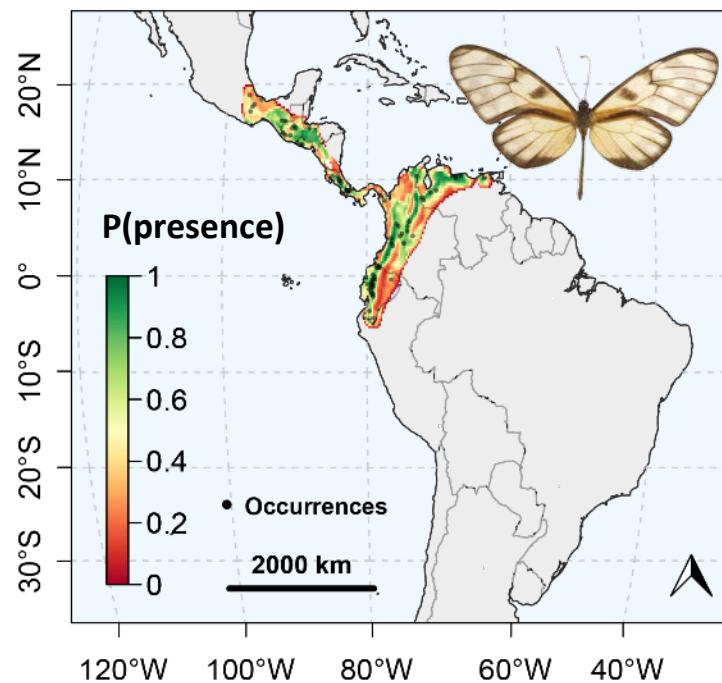
Spatial congruence

Question: Do phenotypically similar species **cooccur** more than expected at **random**?

Hypothesis: Lower **spatial dissimilarity** for comimetic species

$$BC_{ij} = 1 - \frac{2 \sum \min(P_i, P_j)}{\sum P_i + P_j}$$

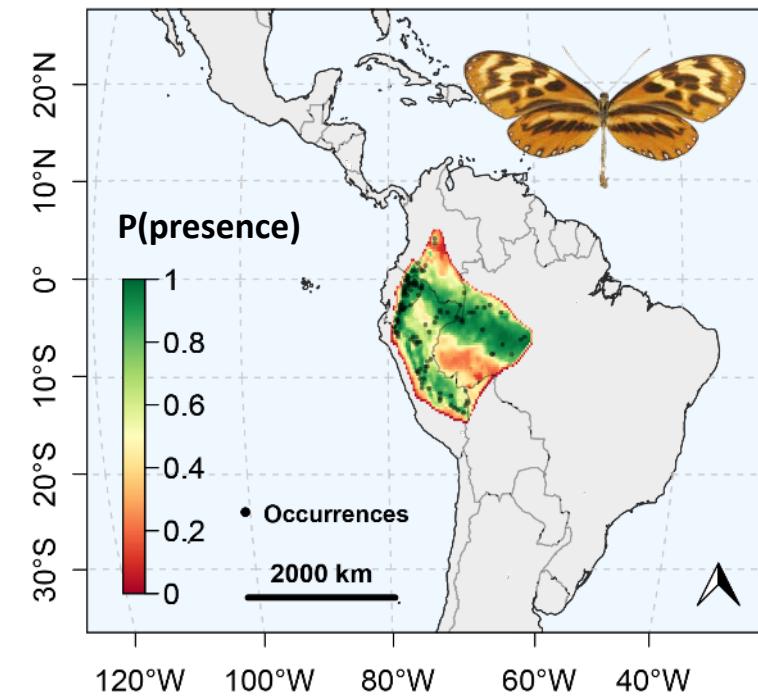
Dircenna jemina (DILUCIDA)



No mimicry

BC = 0.9

Mechanitis mazaeus (MAELUS)



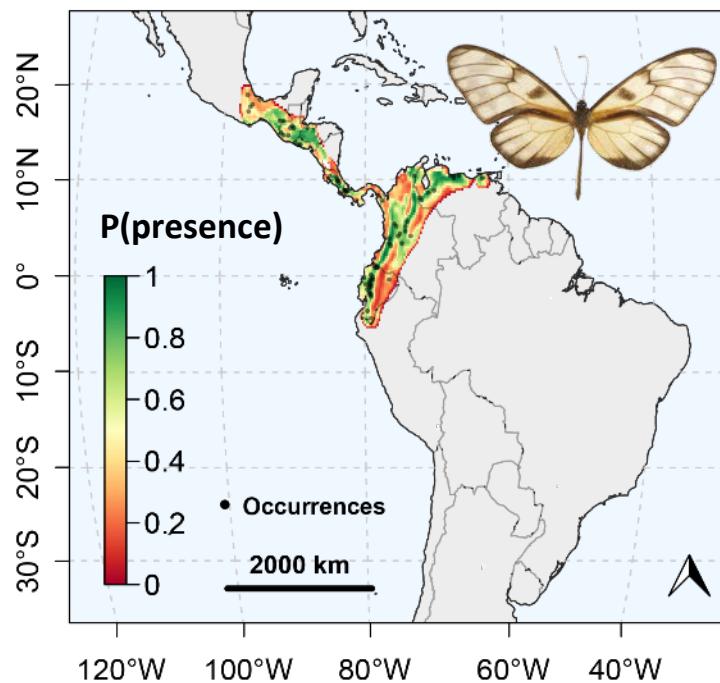
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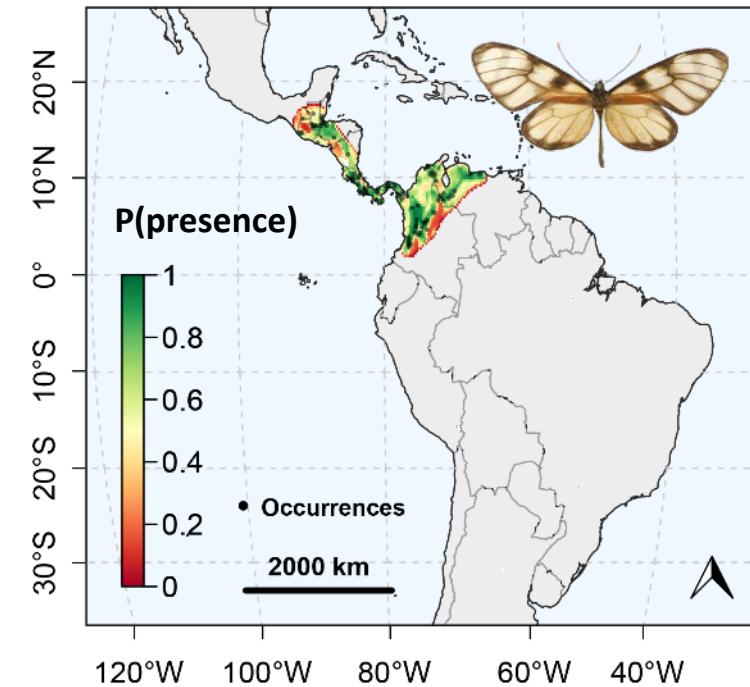
Dircenna jemina (DILUCIDA)



Mimicry

BC = 0.2

Dircenna dero (DILUCIDA)



Spatial congruence

Question: Do phenotypically similar species **cooccur** more than expected at **random**?

Global: Mean $BC_{obs} << \text{Mean } BC_{perm}$

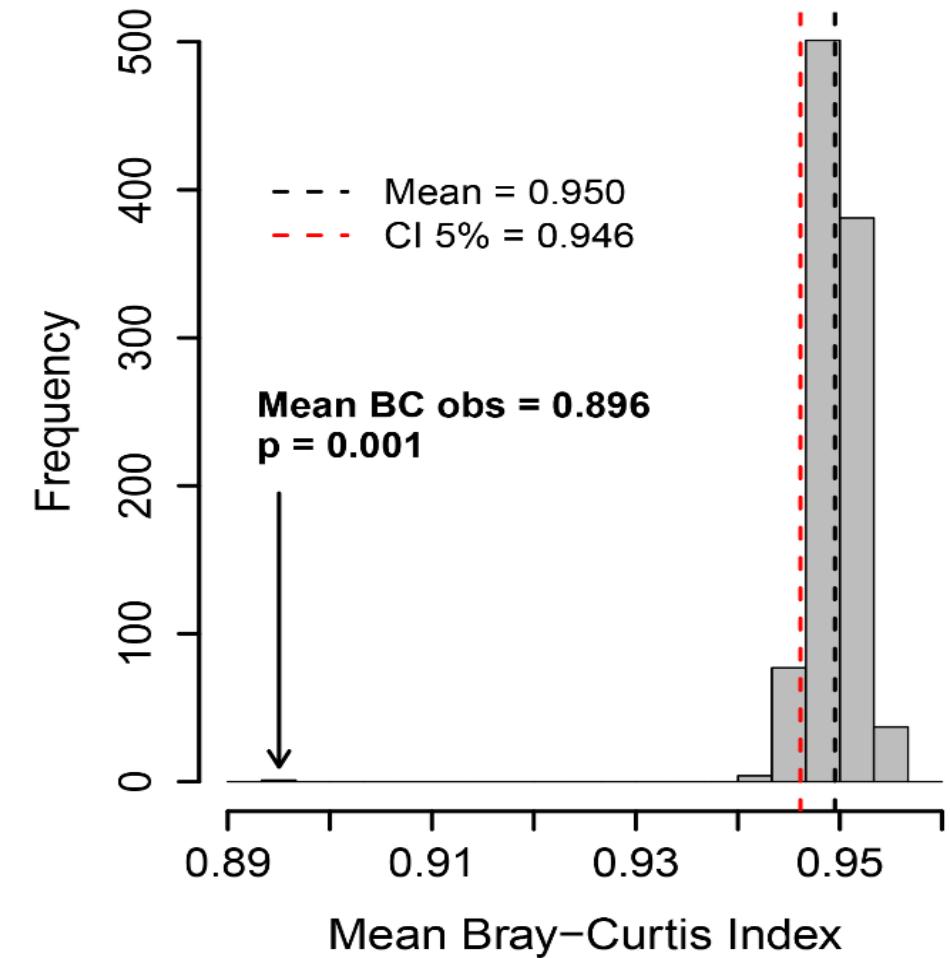


Per ring: Significant for 85% of **putative** rings

- Non-significant rings = low N

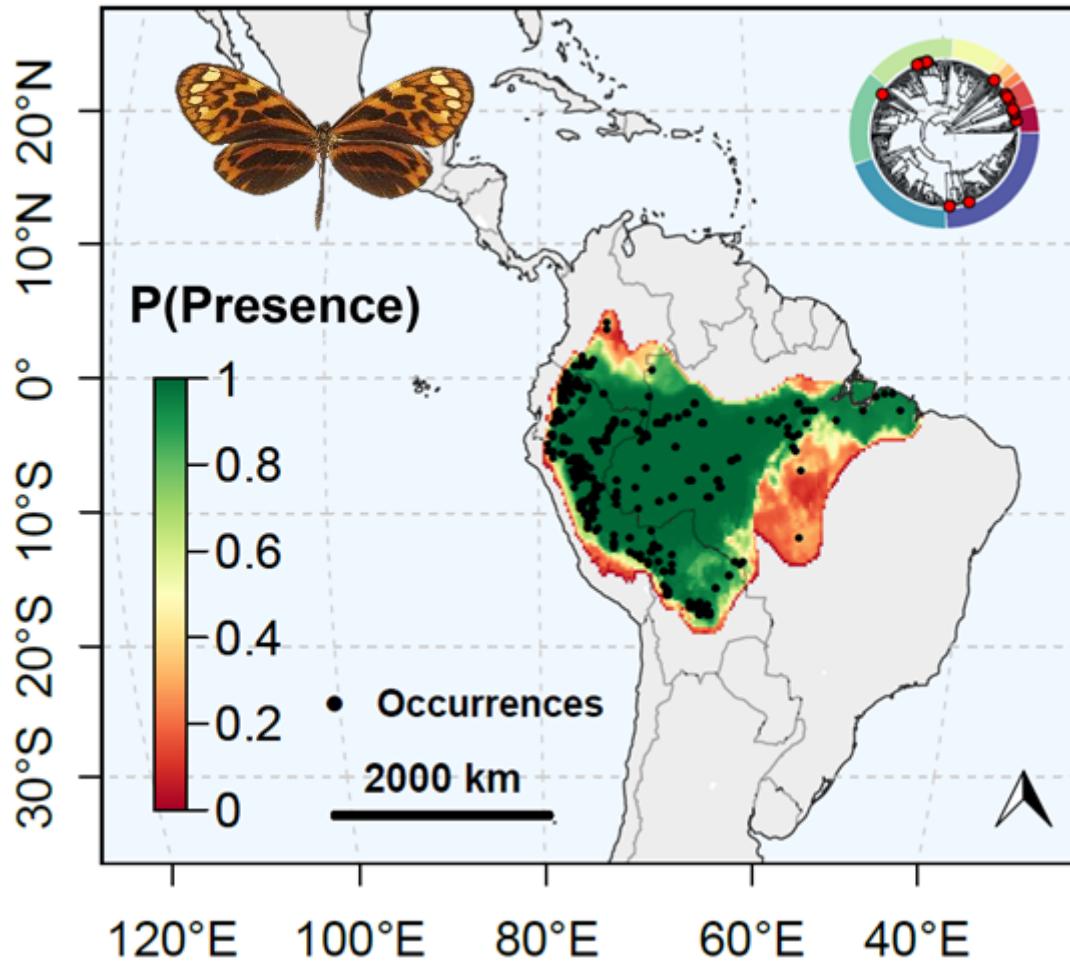
Results: Mimicry promotes the **spatial congruence** of phenotypically similar species at **large-scale**

Next: What happens between tribes?

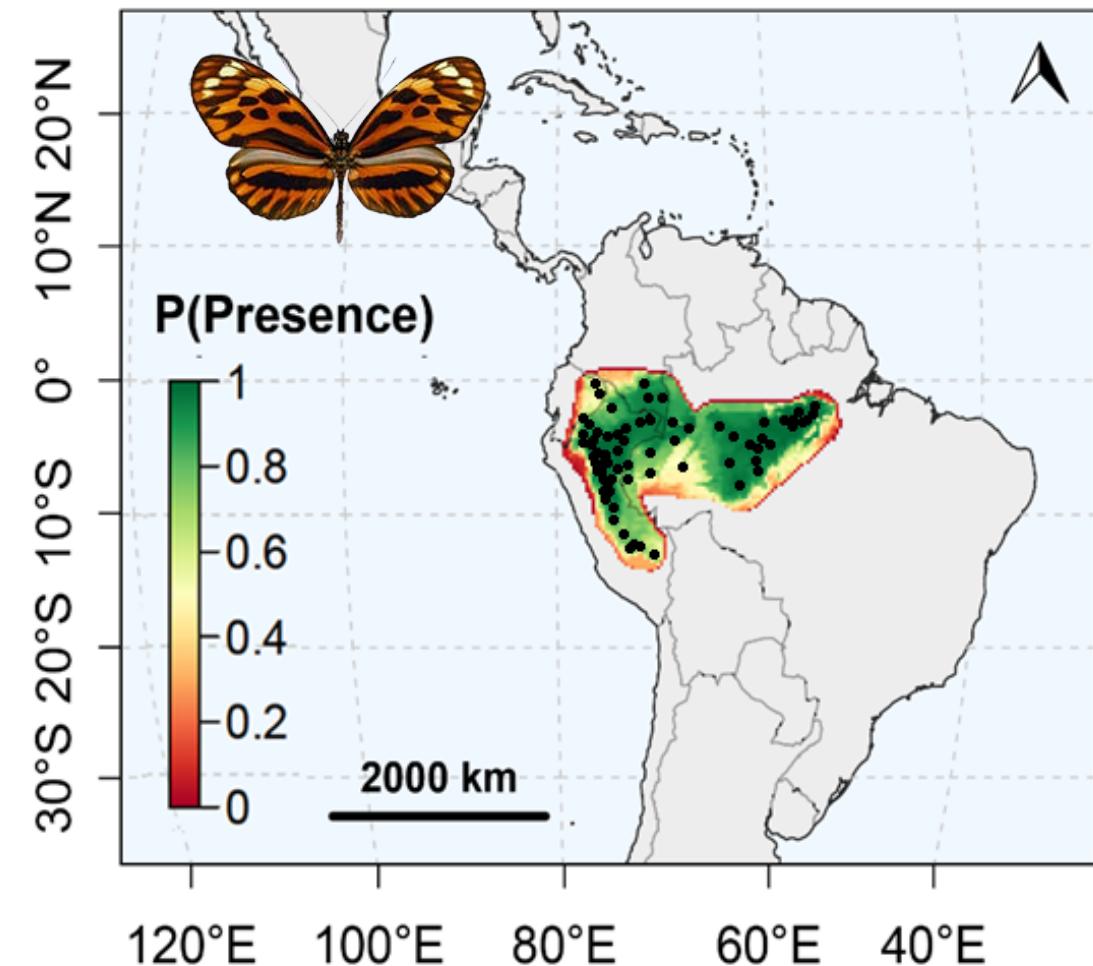


Spatial congruence

Ithomiini: pattern MAELUS
(16 species)

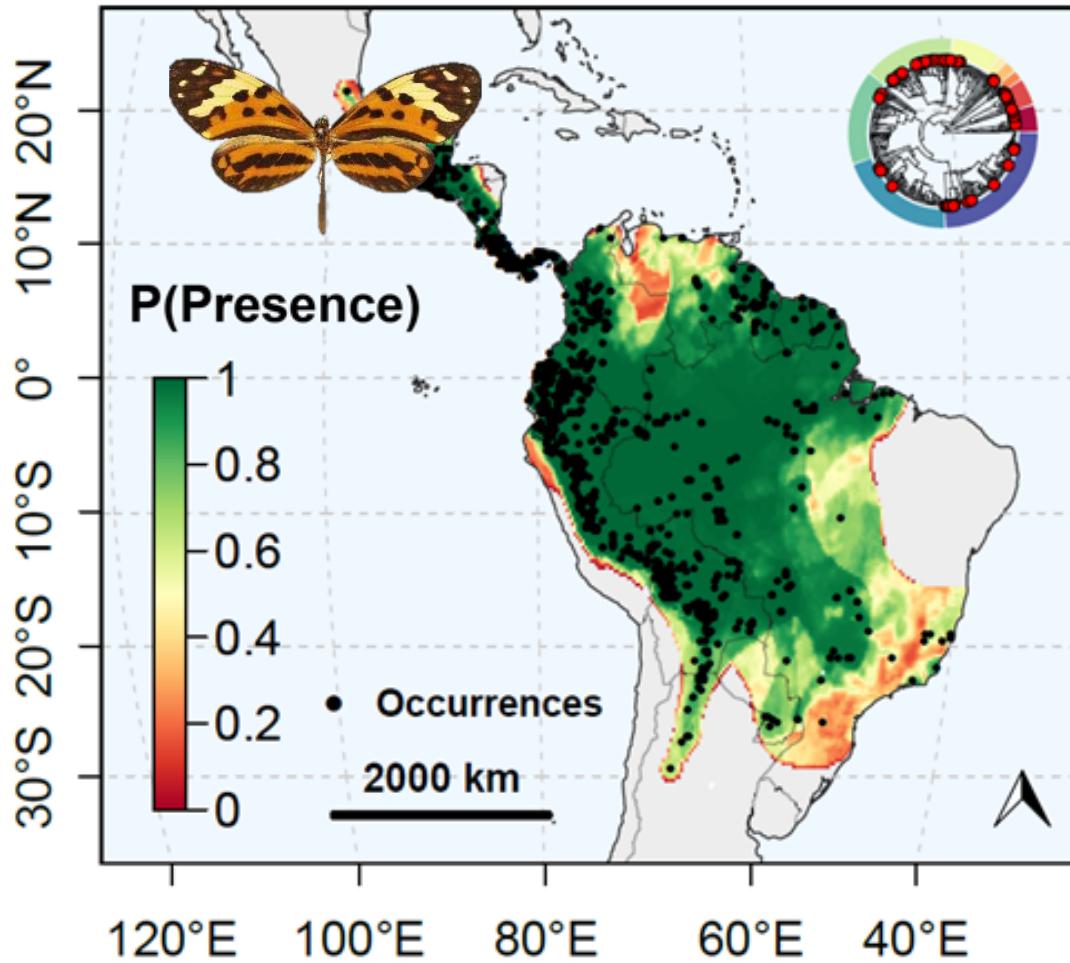


Heliconiini: pattern MAELUS
(5 species)

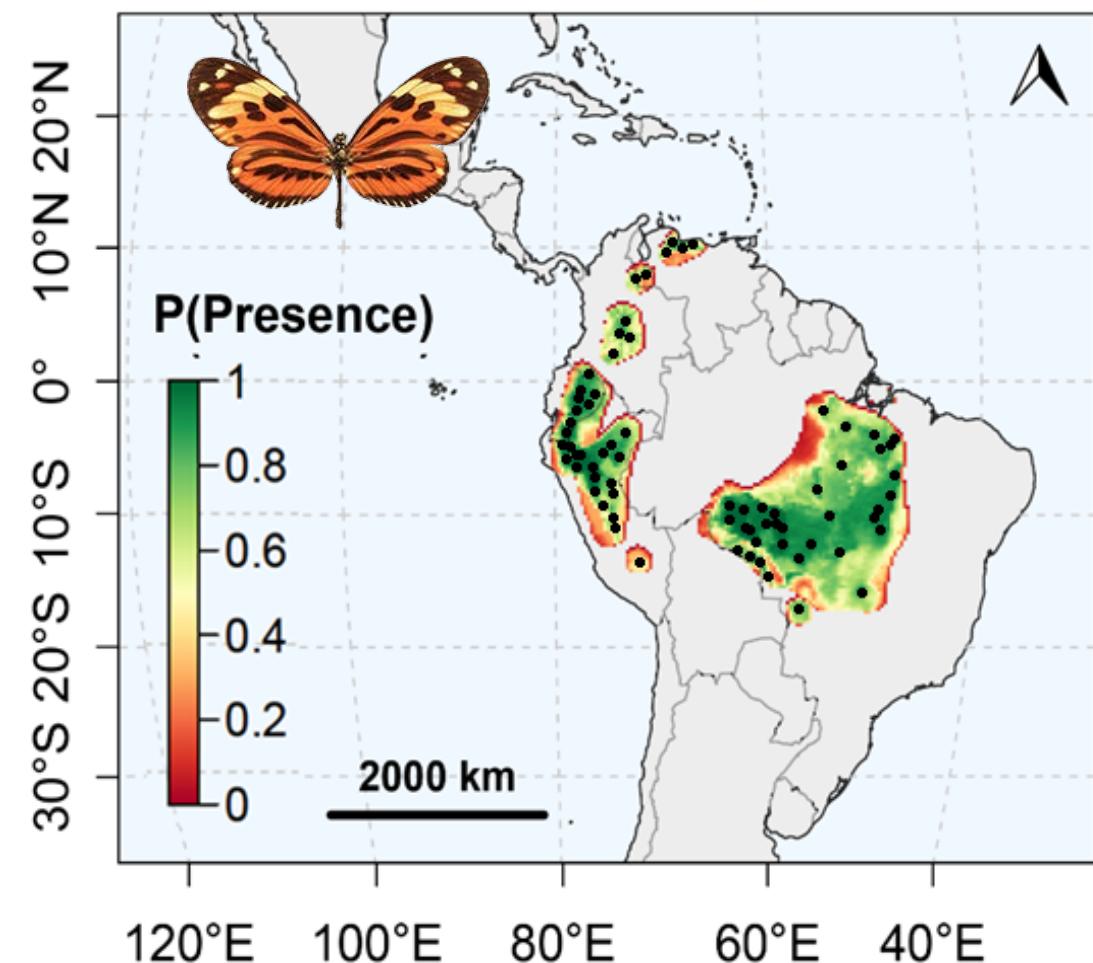


Spatial congruence

Ithomiini: pattern MAMERCUS
(64 species)



Heliconiini: pattern MAMERCUS
(10 species)



Objectives

How **mutualistic interactions** affect the **structure** and **evolution** of biodiversity at the **macroecological scale**?



Spatial congruence: Do phenotypically similar species **cooccur** more than expected at random?



Niche similarity: Do phenotypically similar species have **similar climatic niche**?



Niche convergence: Is the climatic niche of phenotypically similar species **more similar** than expected from **shared ancestry**?



Climatic niche similarity

Question: Do phenotypic groups occupy different climatic niche?

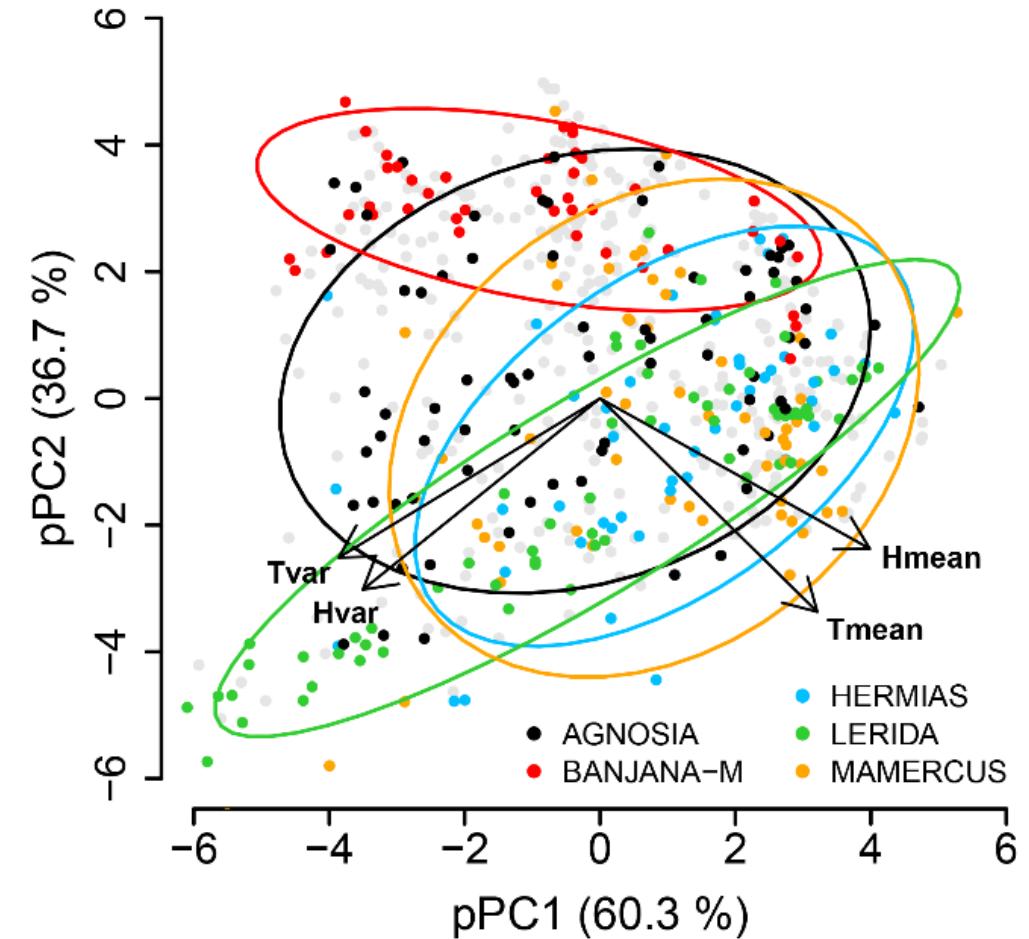
Global: perMANOVA, $R^2 = 0.41$, $p = 0.001$



Per phenotypic group:

- 81.0% pairs with $p < 0.05$
- 66.4% pairs with $p < 0.001$

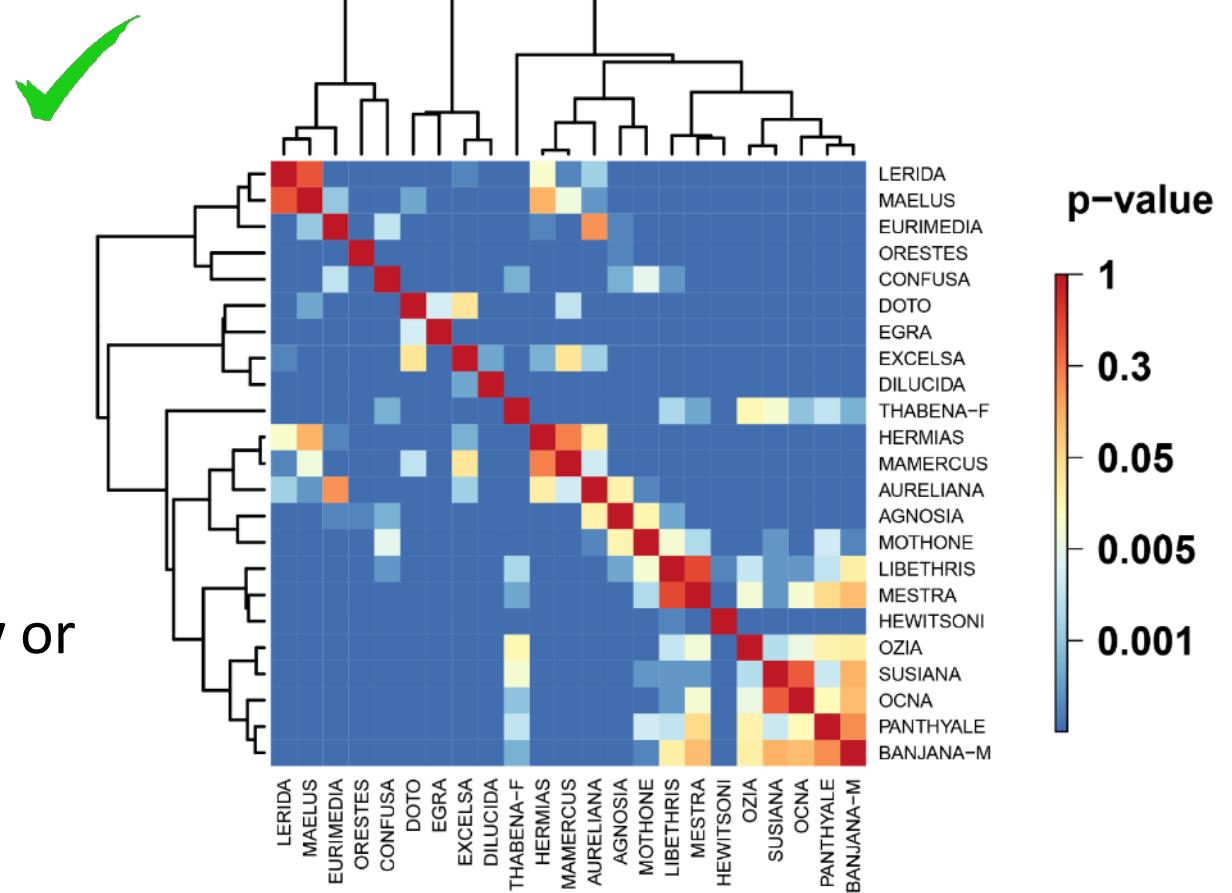
Next: Is this pattern due to shared ancestry or evolutionary convergence?



Climatic niche similarity

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Global: perMANOVA, $R^2 = 0.41$, $p = 0.001$



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Objectives

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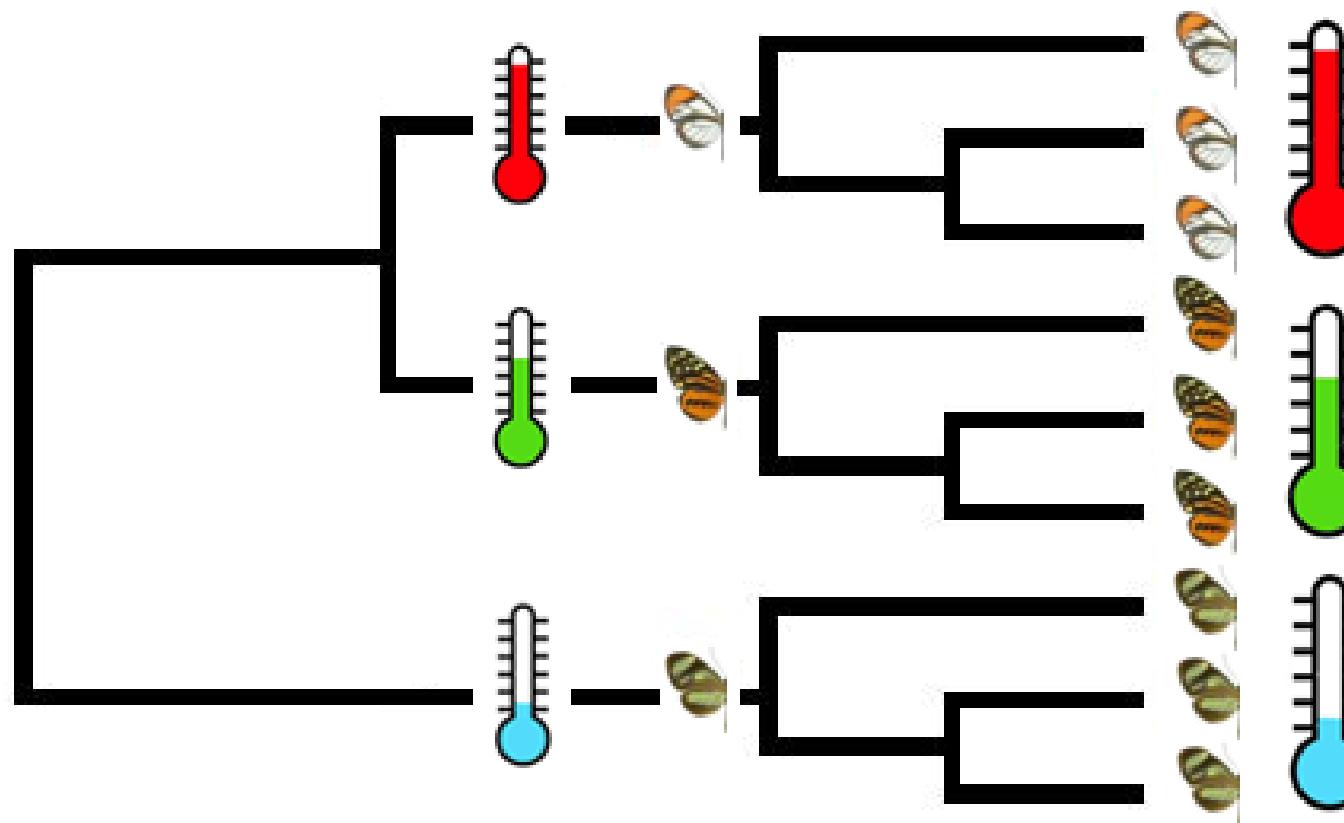


Niche convergence: Is the climatic niche of phenotypically similar species **more similar** than expected from **shared ancestry**?



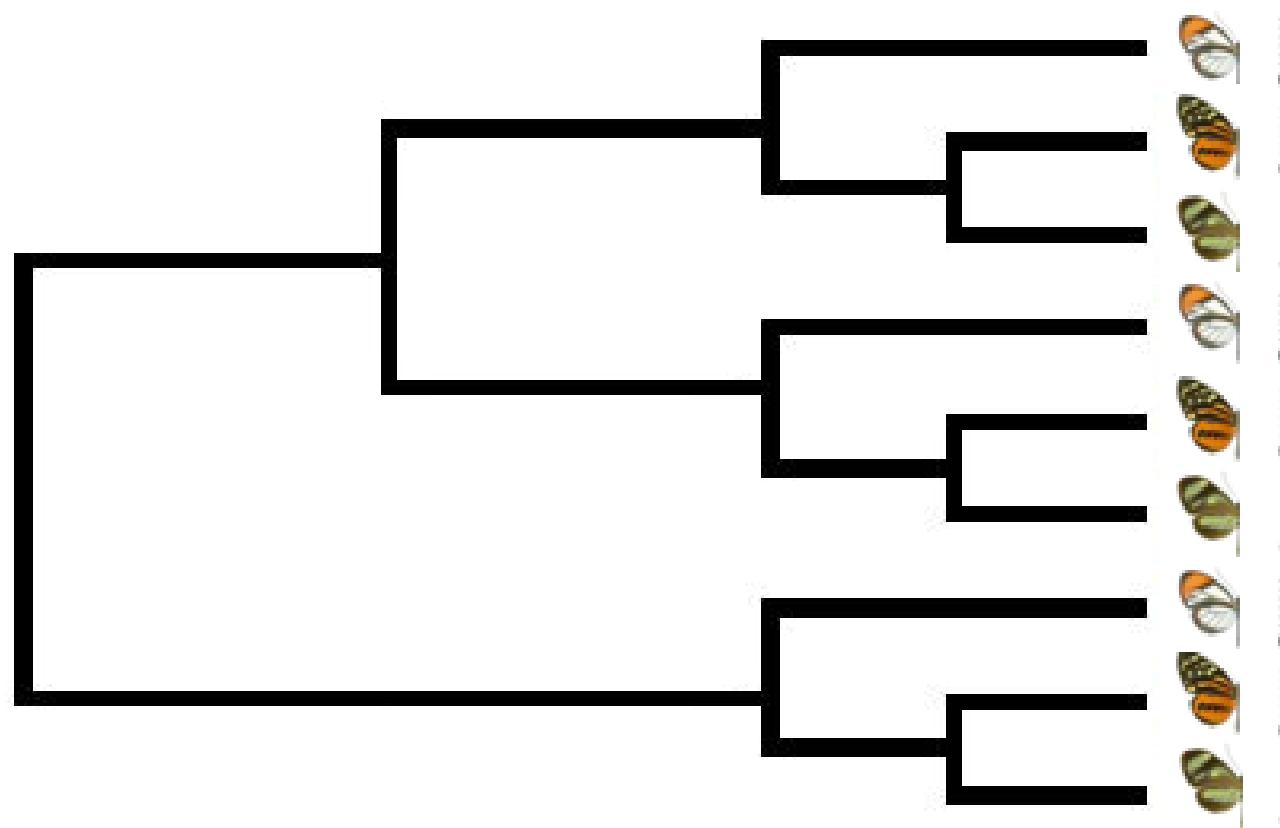
Climatic niche evolution

Question: Is this pattern due to **shared ancestry** or **evolutionary convergence**?



Climatic niche evolution

Question: Is this pattern due to **shared ancestry** or **evolutionary convergence**?



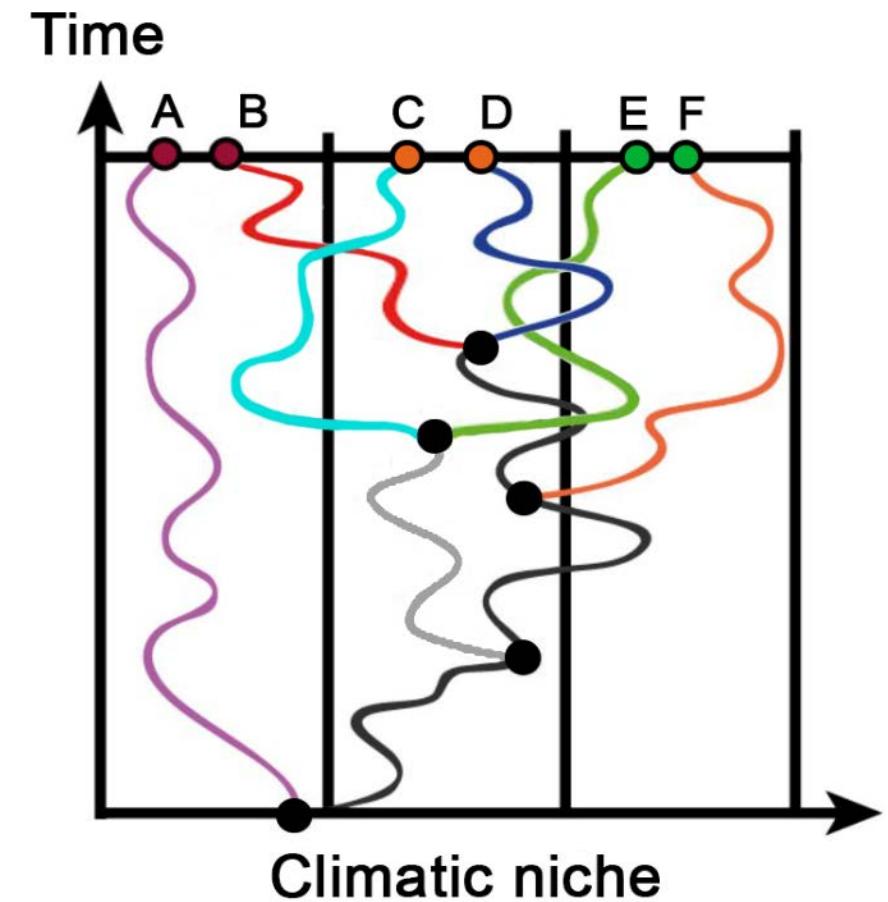
Climatic niche evolution

Question: Is the **climatic niche** of phenotypically similar species more similar than expected from the **phylogeny**?

Simulate the evolution of climatic niche under multivariate **neutral evolutionary model**

phyloMANOVA: $\lambda_{\text{obs}} << \lambda_{\text{simul}}$

Results: Evolutionary association between climatic niche and color patterns



Climatic niche evolution

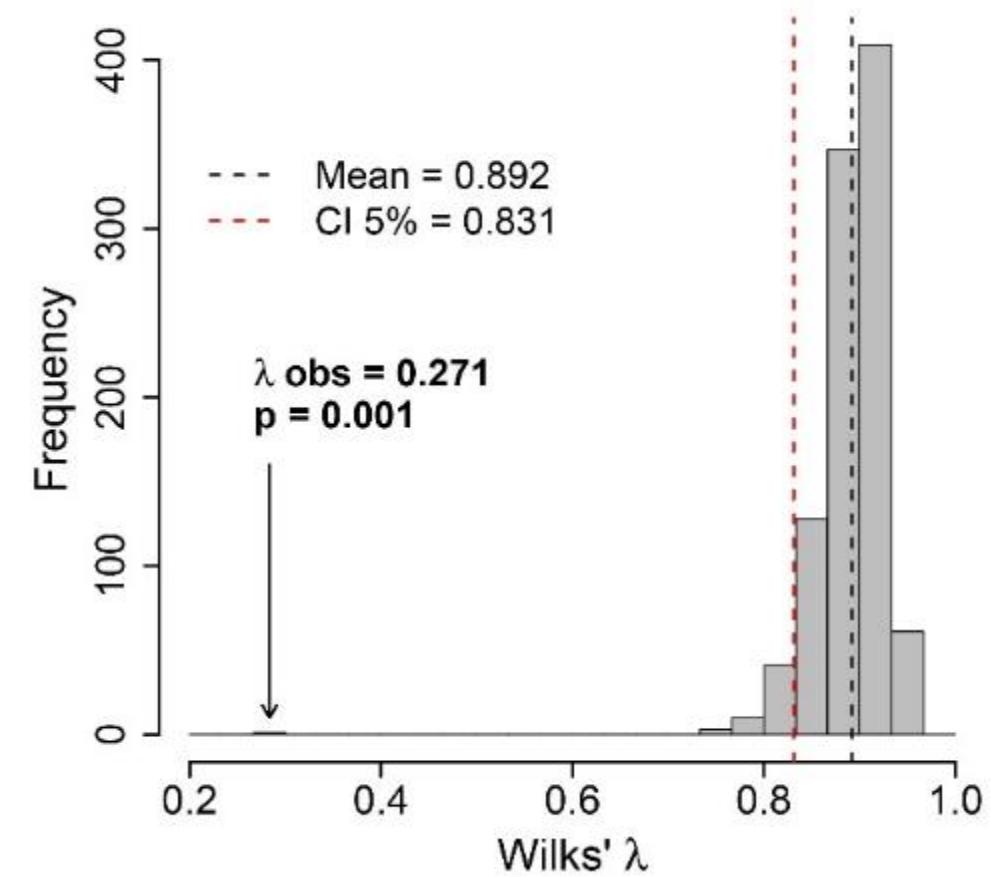
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Climatic niche evolution

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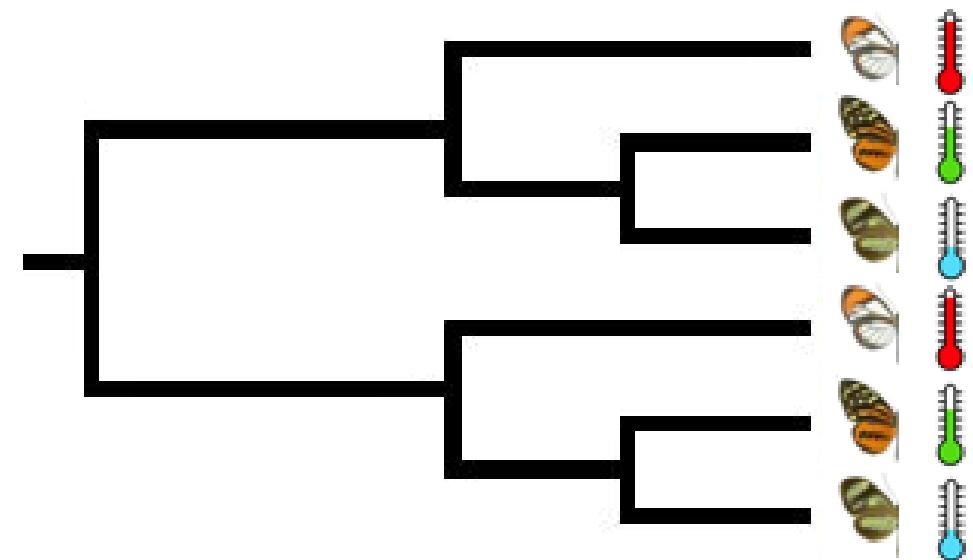
Simulate the evolution of climatic niche under
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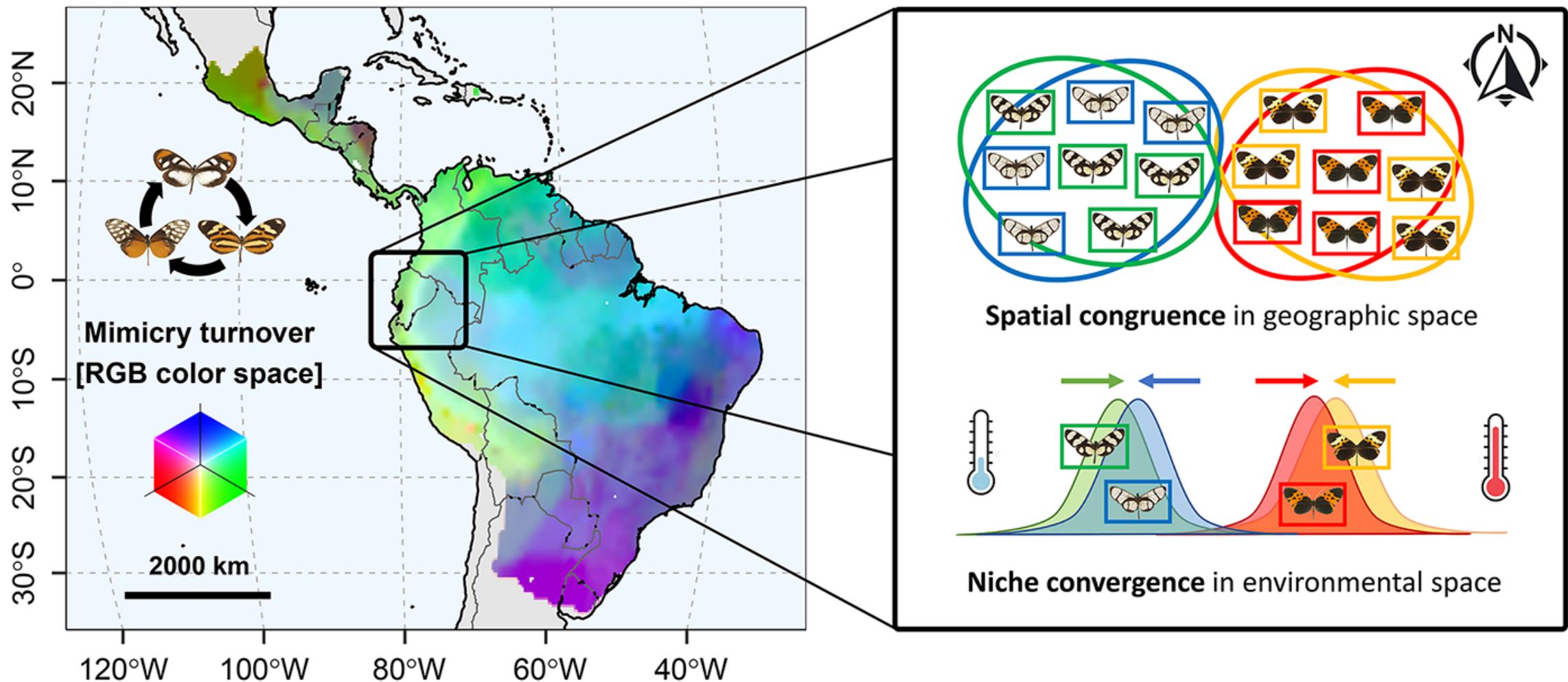


Results: **Evolutionary convergence** between
climatic niche and **color patterns**

Evolutionary convergence



Conclusion



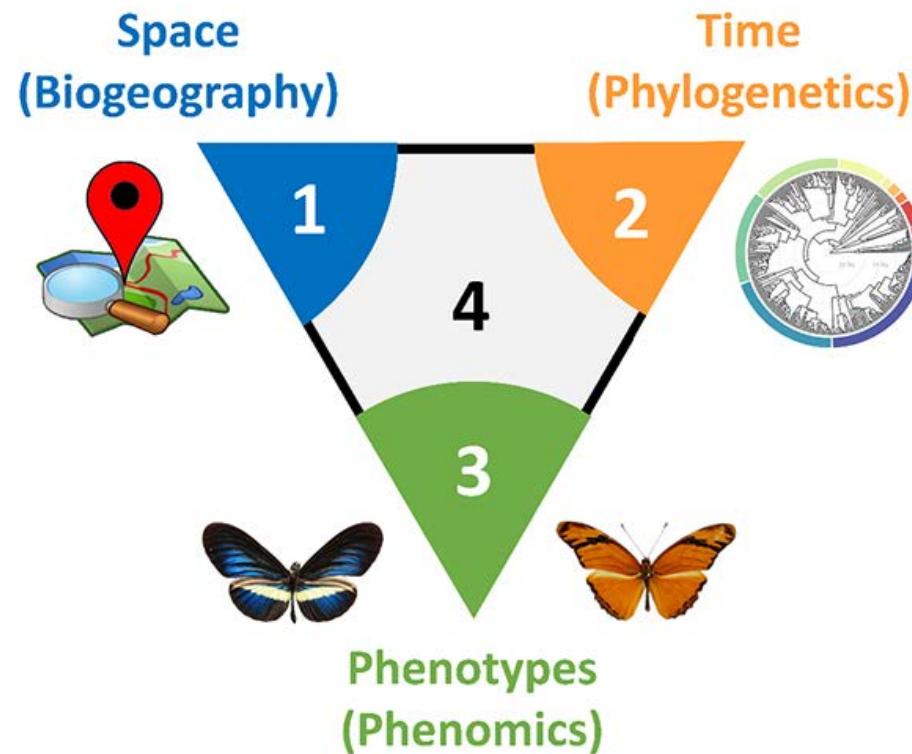
SUMMARY AND PERSPECTIVES

Map biodiversity patterns for an indicator group in the Neotropics

Address the Wallacean shortfall

New method to quantify phenotypic similarity in any ecological signals

Enable multi-scale analyses



Resolve deep evolutionary relationships

New robust phylogeny to test for macroevolutionary hypotheses

New empirical evidence for Müller's model at a macroecological scale

Importance of mutualistic interactions in shaping species distributions and niche evolution

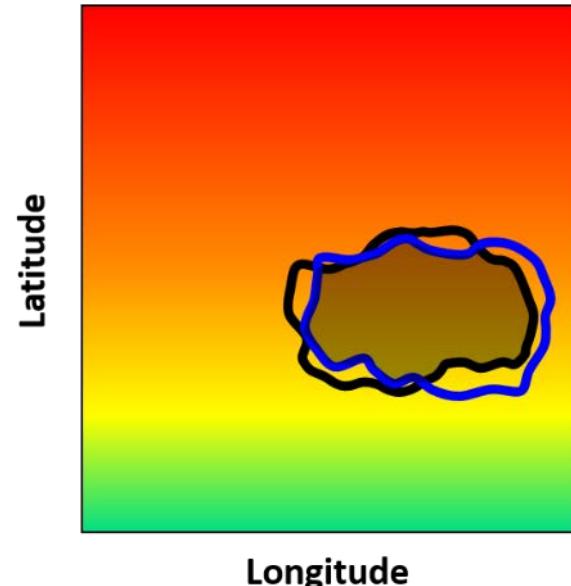
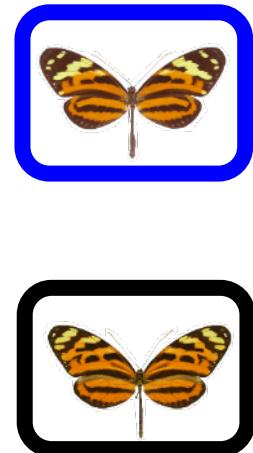
Inclusion in conservation assessments and global change studies

Neotropical mimetic butterflies facing climate changes

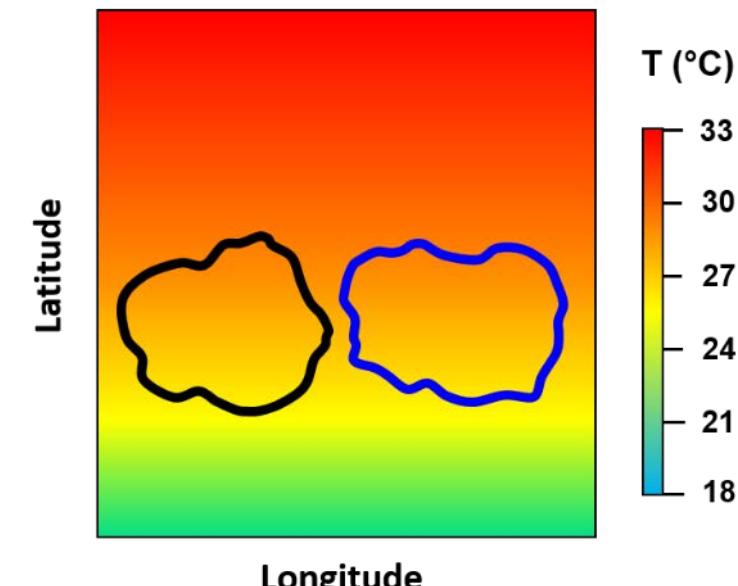
Mutualistic communities = **adaptive assemblage** of species (patterns + niche)
- Sensitivity to interaction disruption and **cascade of extinctions**

Which response to **climate change**?

Cohesion?

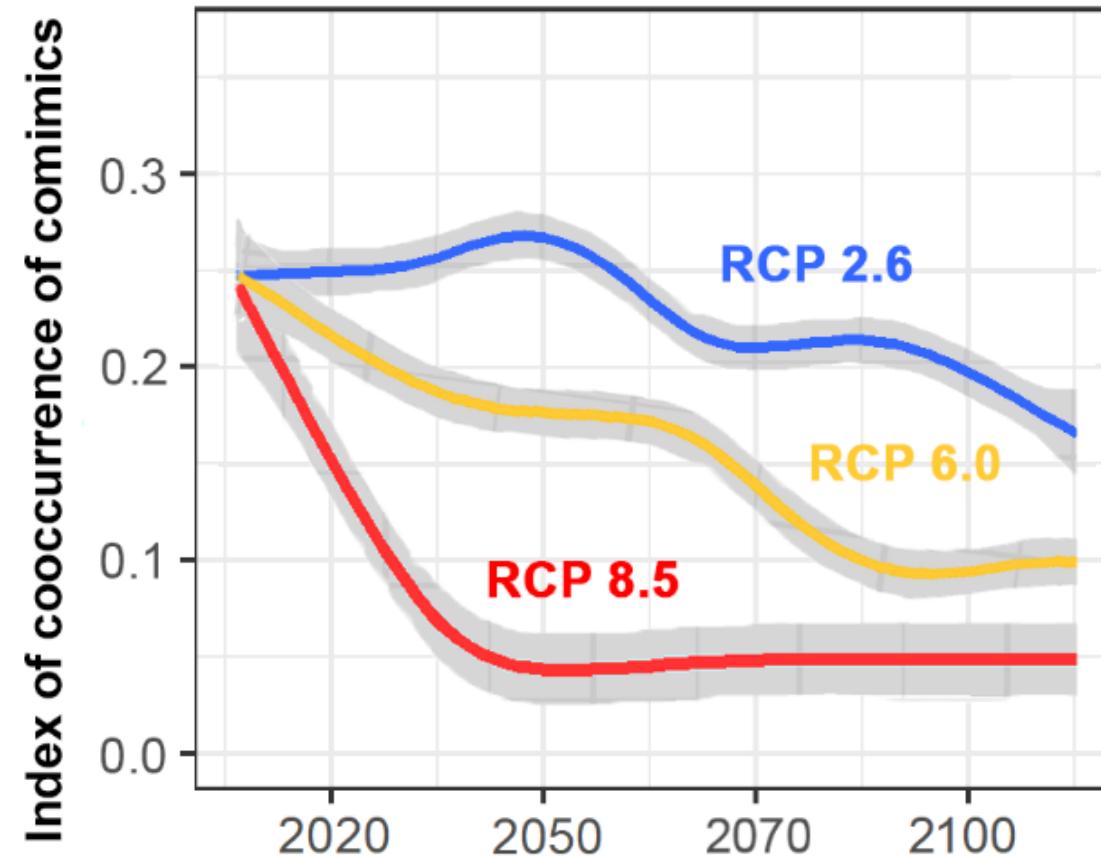


Disassembly?



Neotropical mimetic butterflies facing climate changes

Model future distributions with J-SDM: changes in community composition



Acknowledgments

Advisors: Marianne Elias (ISYEB) & Colin Fontaine (CESCO)

Funding: PhD Grant (French MESR) & Marianne Elias (HFSP Grant)



Main collaborators:

- Eddie Pérochon
- Keith Willmott
- Boris Leroy
- Jérémie Gauthier
- Rémi Allio
- Thomas Aubier
- Mathieu Joron
- André VL Freitas
- James Mallet
- Nicolas Chazot
- Nick Grishin
- Joana Meier
- Sébastien Lavergne
- Neil Rosser
- Krzysztof Kozak





Thanks for your
attention

Hypomenitis enigma (Ithomiini). Credits: M. Elias

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