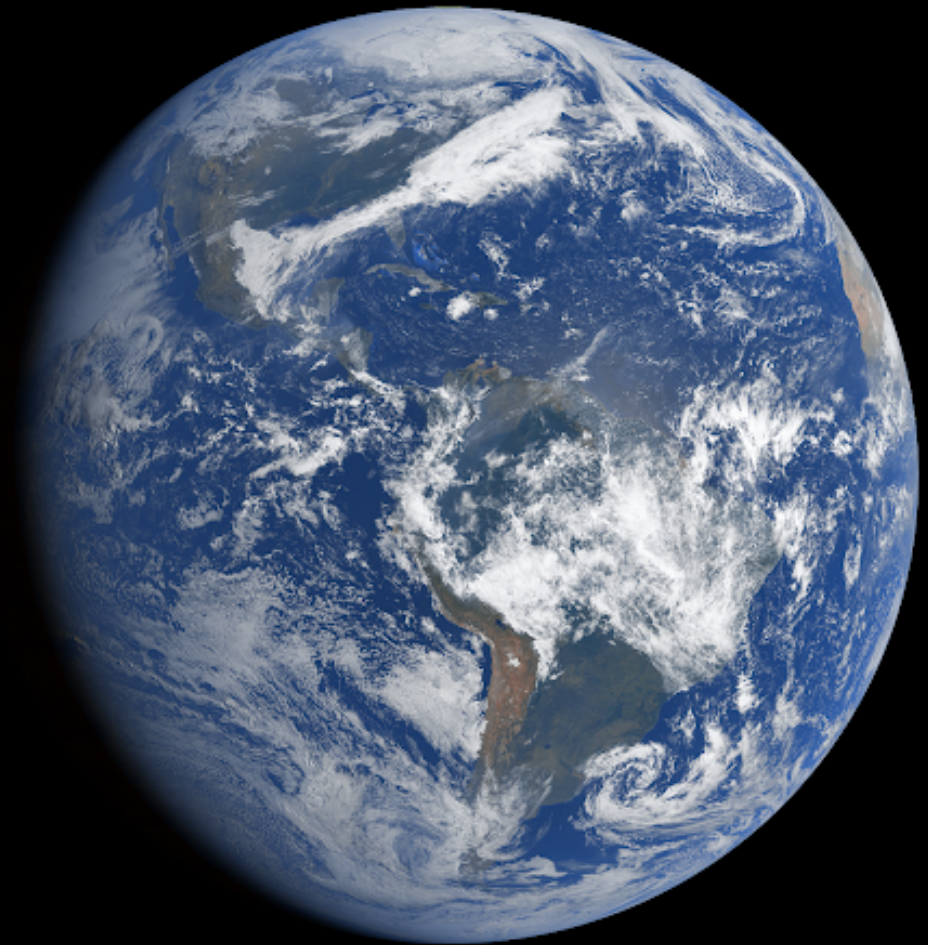


Phylogeography of Arachnids



Věra Opatová
Dept. of Zoology
Faculty of Sciences -Charles University

Biogeography

Processes responsible for current and past distributions of the biota

Ecological Biogeography – species/intraspecific level

- limiting characteristics of current distribution
- ecological preferences, environmental factors, competition, host distribution....

Historical Biogeography – related taxa, family level

- processes that shaped the distributions into the patterns we observe today
- geological history, climate

...our resulting hypotheses are only as good as our input data and our own biases...

...our resulting hypotheses are only as good as our input data and our own biases...
[particularly in case of Historical Biogeography]

Evolutionary theory + Biogeography - Ch. Darwin (1859), AR Wallace (1869, 1878)

Plate tectonics theory (Continental drift)

formulated 1912 by Wegener not accepted until 1960s

→ **Dispersal** is responsible for today's distribution patterns, same geography

X

Vicariance – Croizat 1950

The organisms had the same distribution in the past (always!)

- slow steady spread across continuous land, barriers appeared later

What is the contribution of each process?

Fossil record and the lack of thereof

- not very rich in case of Arachnids, extinct lineages - difficult to assign
- modern lineages in amber: Burmese (~100 Ma), Baltic Amber (~44 Ma)
Dominican Amber (~30 Ma)
- more resources: <https://wsc.nmbe.ch/resources/fossils/Fossils20.5.pdf>

...our resulting hypotheses are only as good as our input data and our own biases...
[both Ecological and Historical Biogeography]

Taxonomy/understanding of Biodiversity

- what is a species, how many species there are, how are they related?

Group	Number of described species	Likely total	%
Insects	950 000	8 000 000	12
Fungi	70 000	1 000 000	7
Arachnids	75 000	750 000	10
Viruses	5000	500 000	5
Nematodes	15 000	500 000	3
Bacteria	4000	400 000	1
Vascular plants	250 000	300 000	83
Protozoans	40 000	200 000	20
Algae	40 000	200 000	20
Molluscs	70 000	200 000	35
Crustaceans	40 000	150 000	27
Vertebrates	45 000	50 000	90

Cox et al 2010

Resources/Interest and the lack of thereof

Phylogeography

Phylogenetics + Biogeography (Avice 2000)

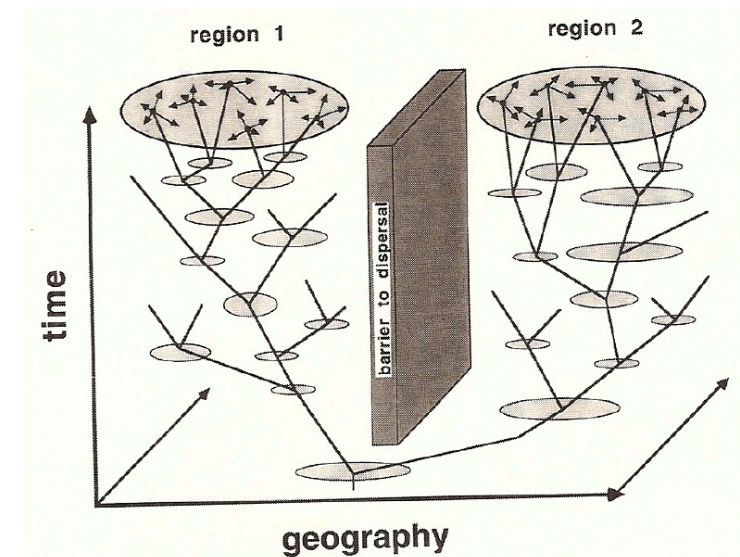
Geographic distribution of genetic lineages (traditionally - closely related)

The question remains:

- Which processes shaped the current and past spatial distributions of these lineages?

Implementation of molecular methods – new perspective

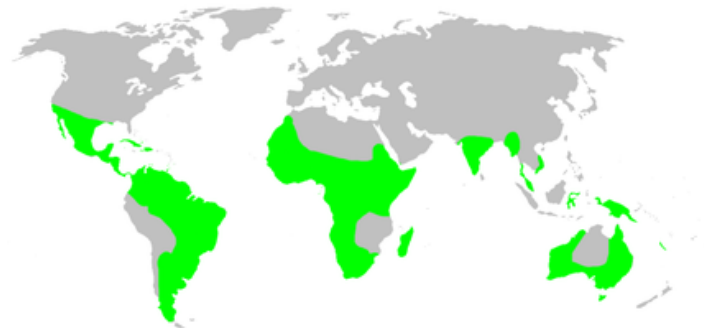
- geographic structure in the populations
 - geographic history, dispersal routes & barriers
 - concordant patterns among different species
 - conservation purposes
- potential existence of cryptic species
 - taxonomy
- molecular dating



Dispersal? Introduction? Vicariance? Extinction?



Archaeidae



Deinopidae



Salticidae

usually a combination of more than one factor...

Dispersal in Arachnids

The capability to overcome barriers differs significantly

Key role in colonizing new habitats

- weak population structure in highly mobile groups
- deep population structuring in sedentary groups

Passive dispersal:

Phoresy: pseudoscorpions, mites, *Attacobius attarum*

Rafting – short/long distance dispersal

Accidental introduction: synanthropes in advantage

Airborne/wind: mites

Host mediated - ticks

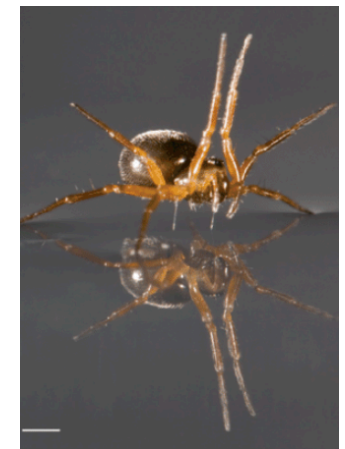
Active dispersal:

Ballooning: spiders

Walking

Sailing

Tumbling - *Cebrennus rechenbergi*



Phoresy

Attaching of non-vagile individual to a different species "carrier"

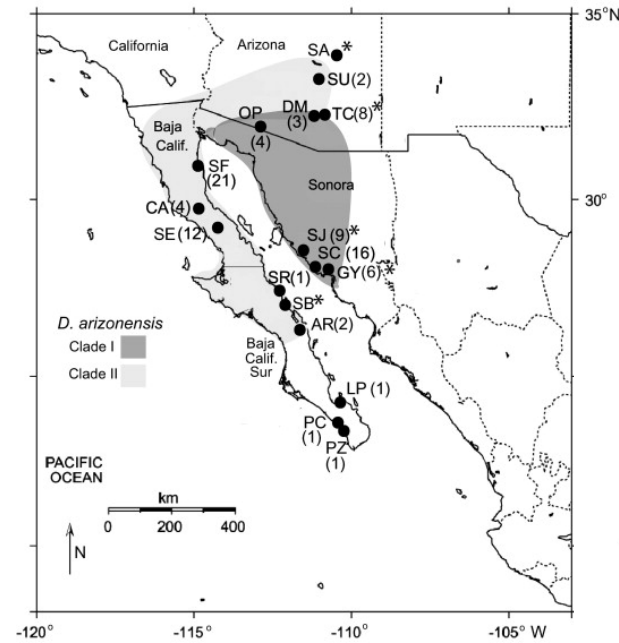
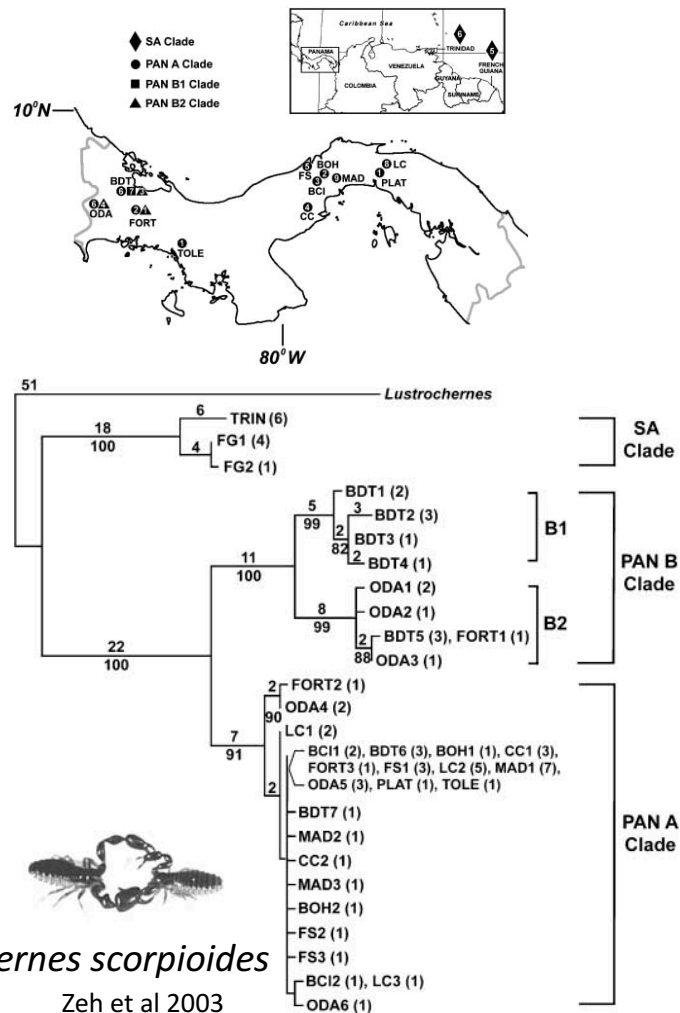
Colonization of temporary habitats

Little information about the genetic structure of phoretic species

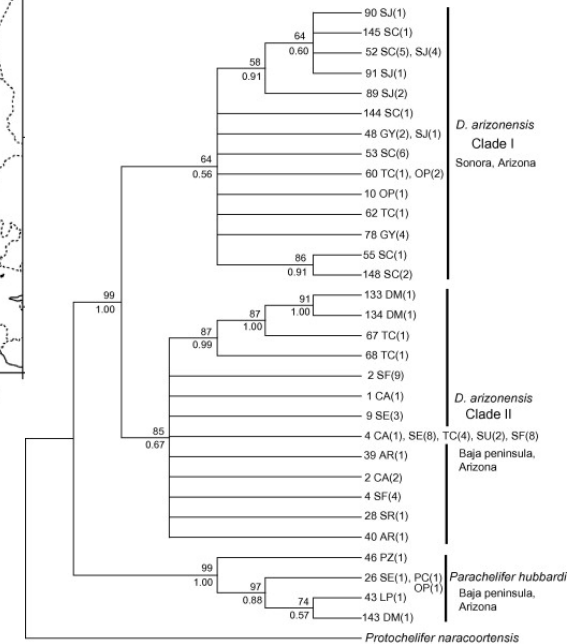


44 Ma

Dunlop & Penney 2012

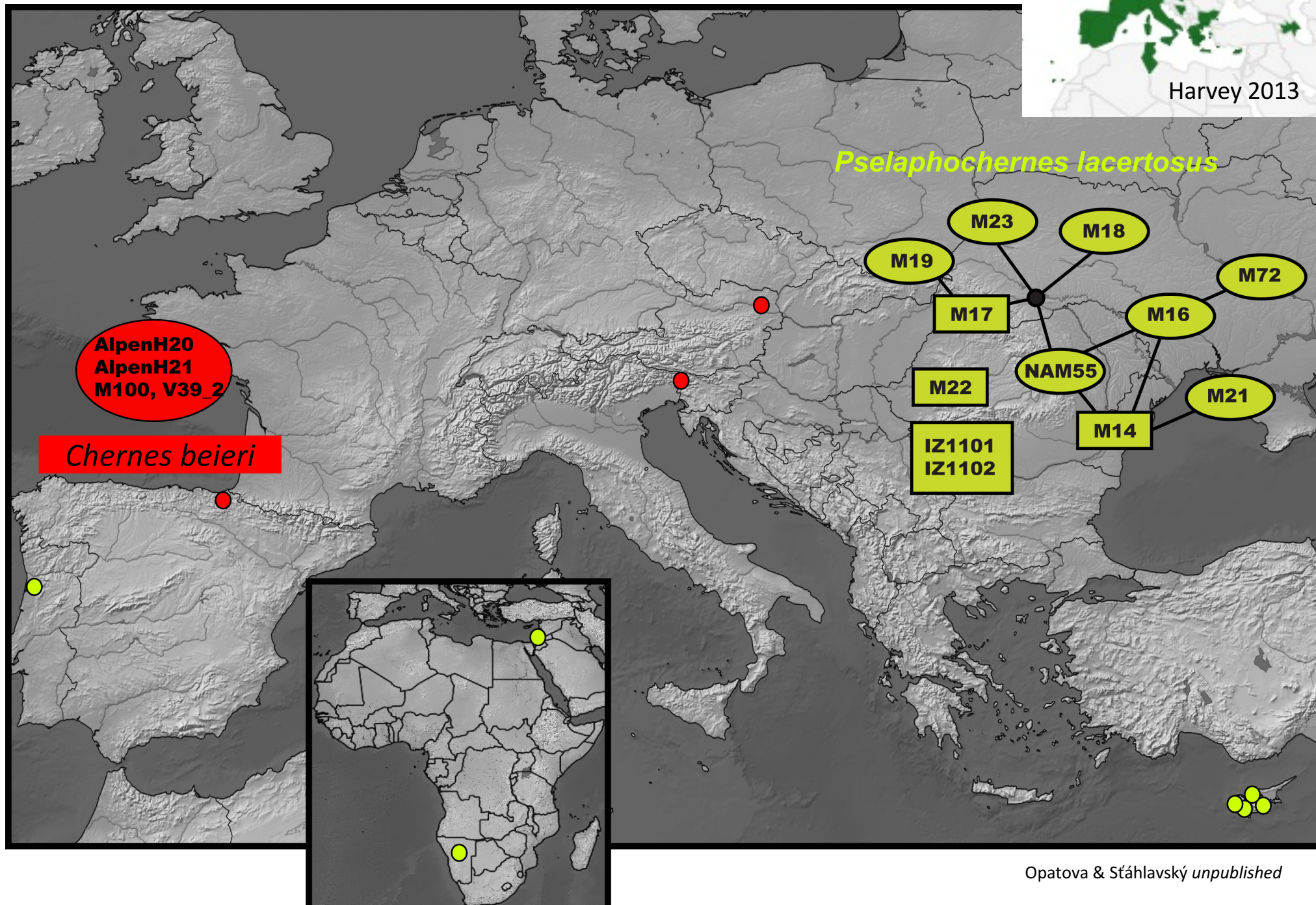


Dinocheirus arizonensis

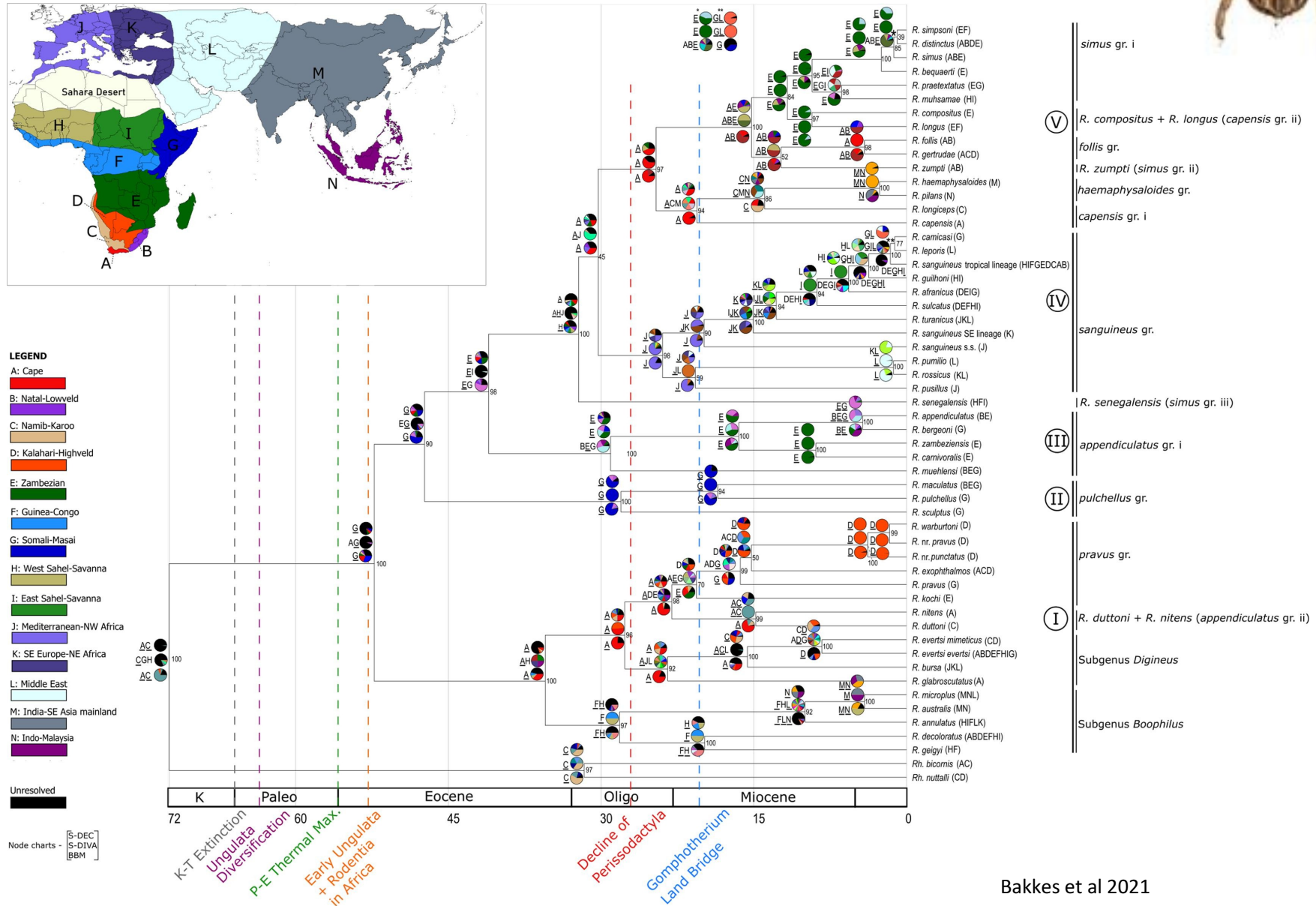


Pfeiler et al 2009

P. lacertosus, *C. beieri* – likely phoresy on birds



Host mediated dispersal and radiation - *Rhipicephalus* ticks



Host mediated dispersal and radiation - *Rhipicephalus* ticks

Host-enabled dispersal events to new environments

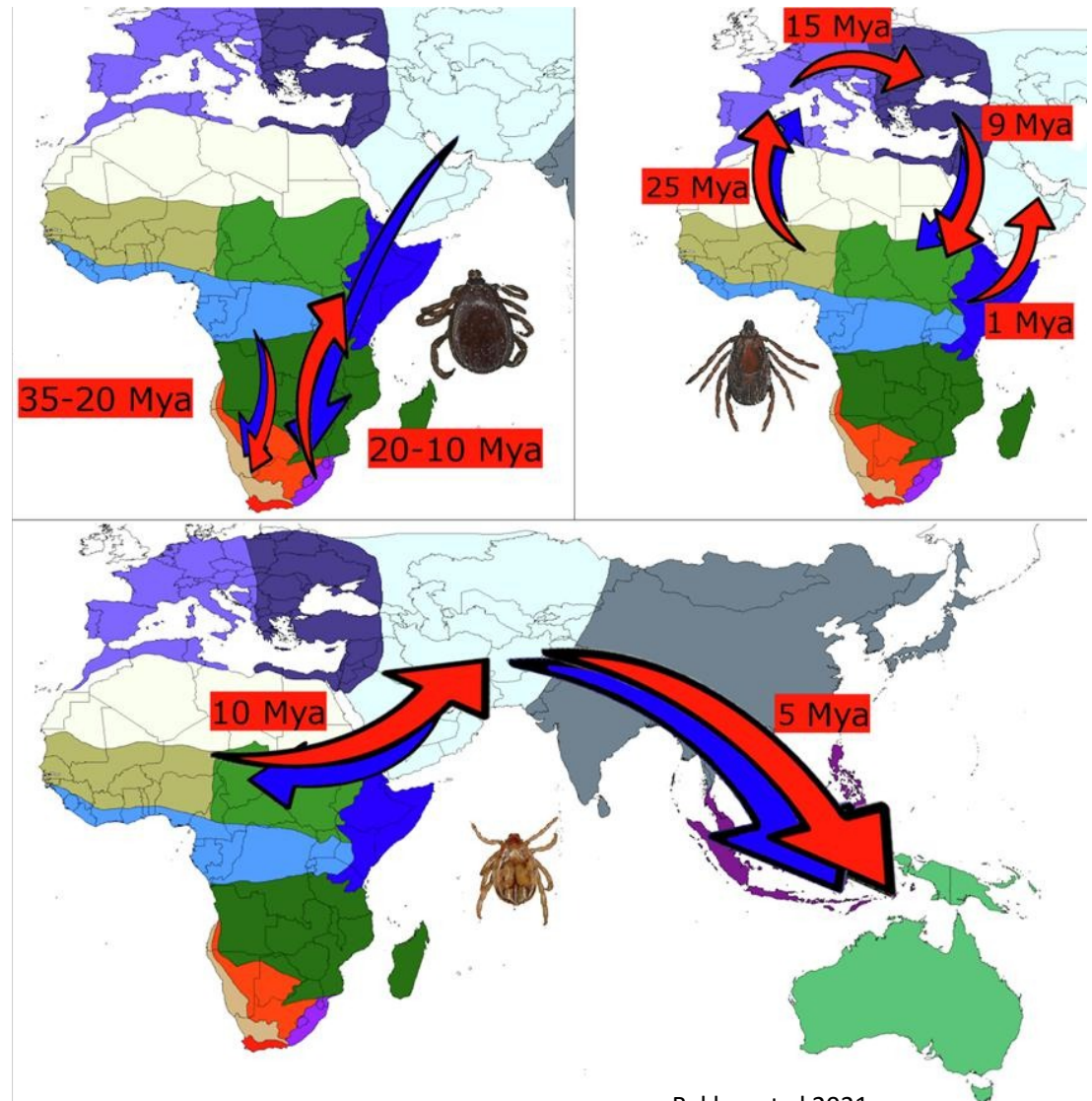
followed by local adaptations

larvae on large and mobile hosts

→ larger ranges, slower rates

larvae on small and less mobile hosts

→ smaller ranges, faster rates



Ballooning

aerial dispersal

Long/ short distance dispersal

juveniles of large species/ adults of small species

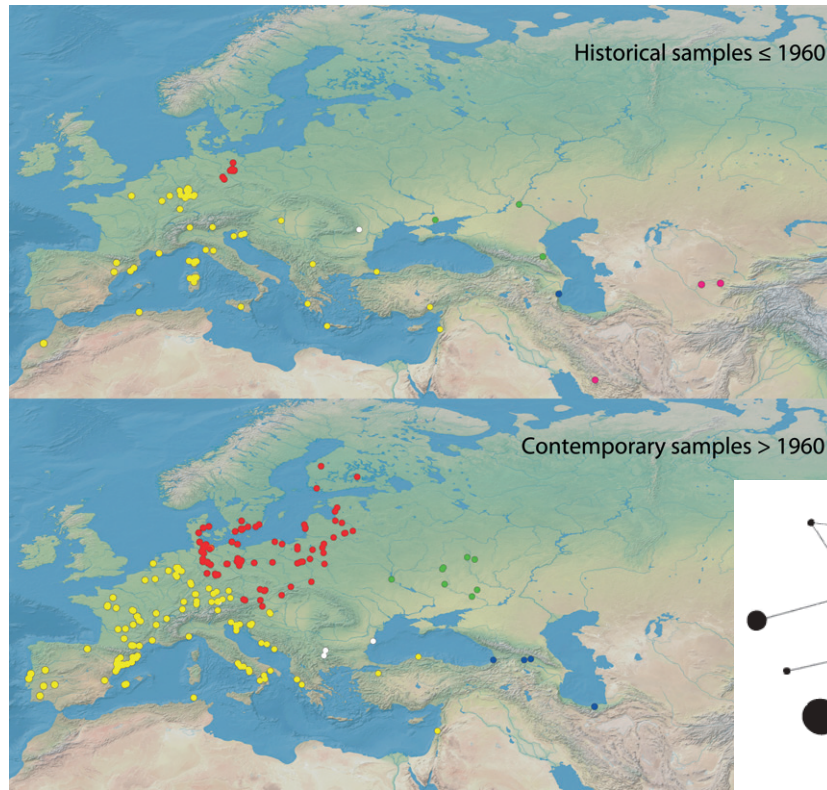
very common in Araneomorphae
- may differ within a family

uncommon in Mygalomorphae
- not as effective



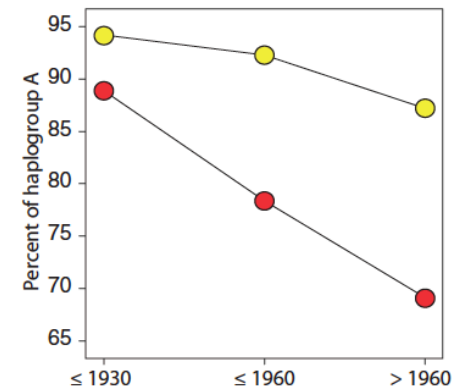
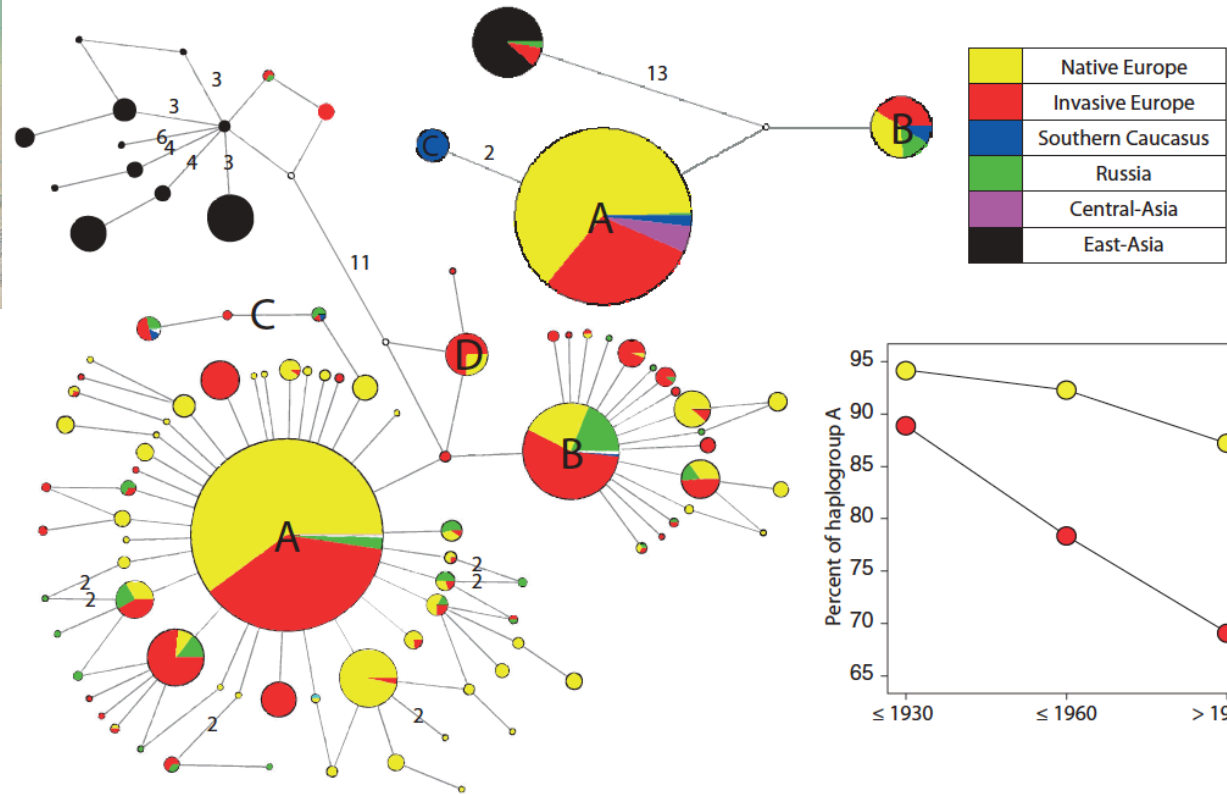
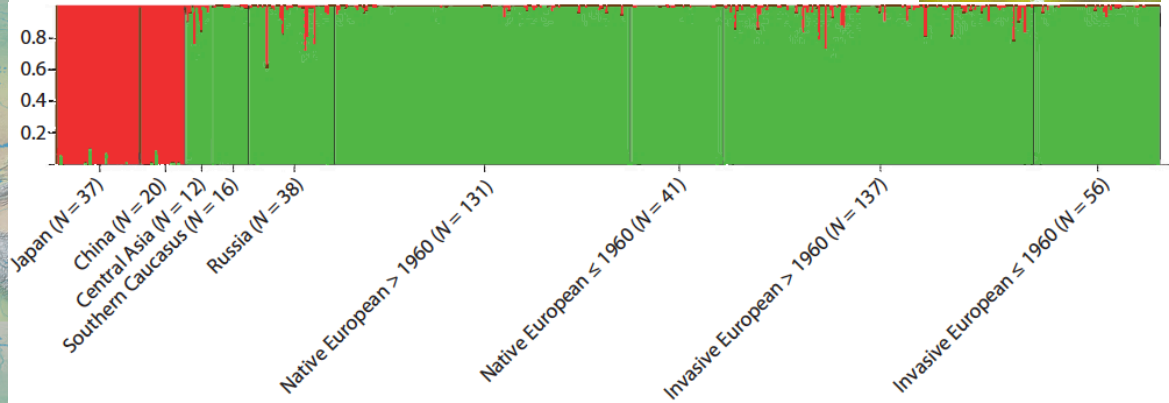
In all cases the ballooning individuals must land in an area with favorable conditions
or be preadapted to novel conditions

Argiope bruennichi – range expansion



SNP

Krehenwinkel & Tautz (2013), Mol Ecol
Krehenwinkel et al (2015), Glob Change biol



Mt diversity

- higher in invasive populations

Body size

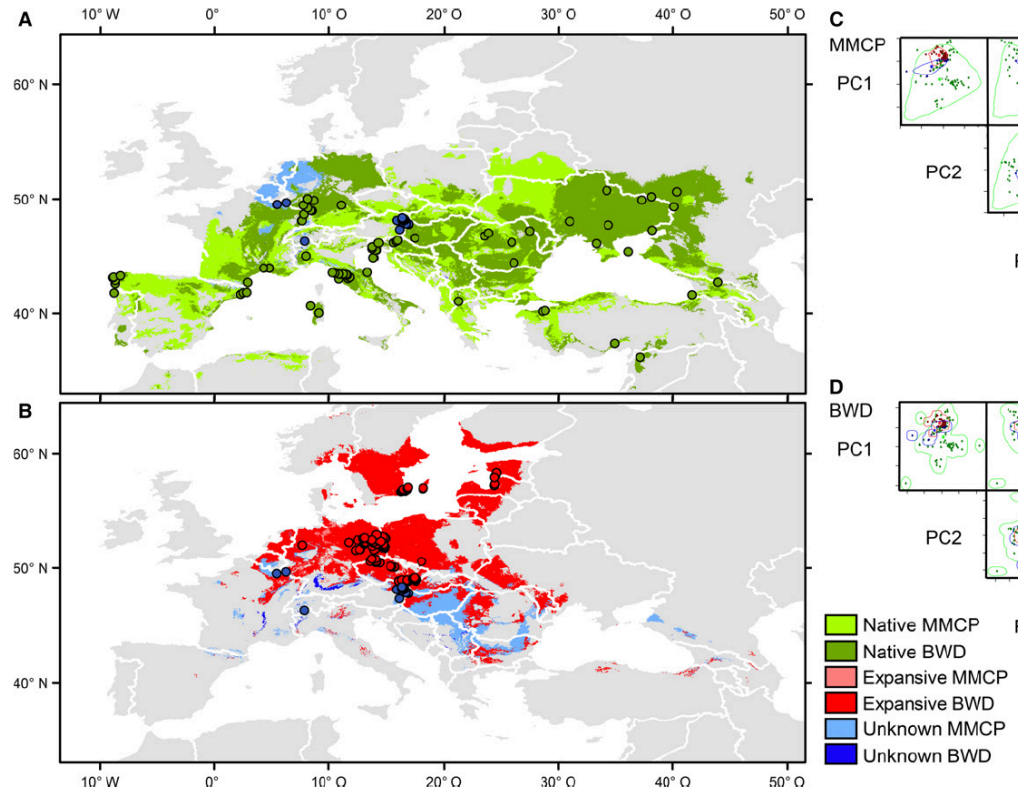
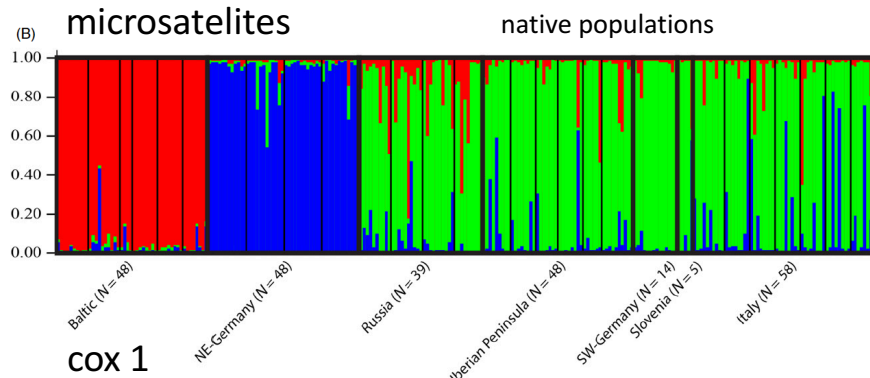
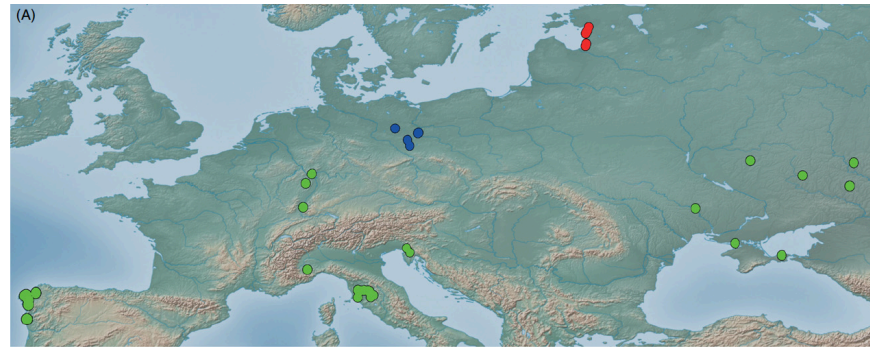
- smaller in invasive populations

Cold tolerance

- differences in gene expression

Cheiracanthium punctorium – range expansion

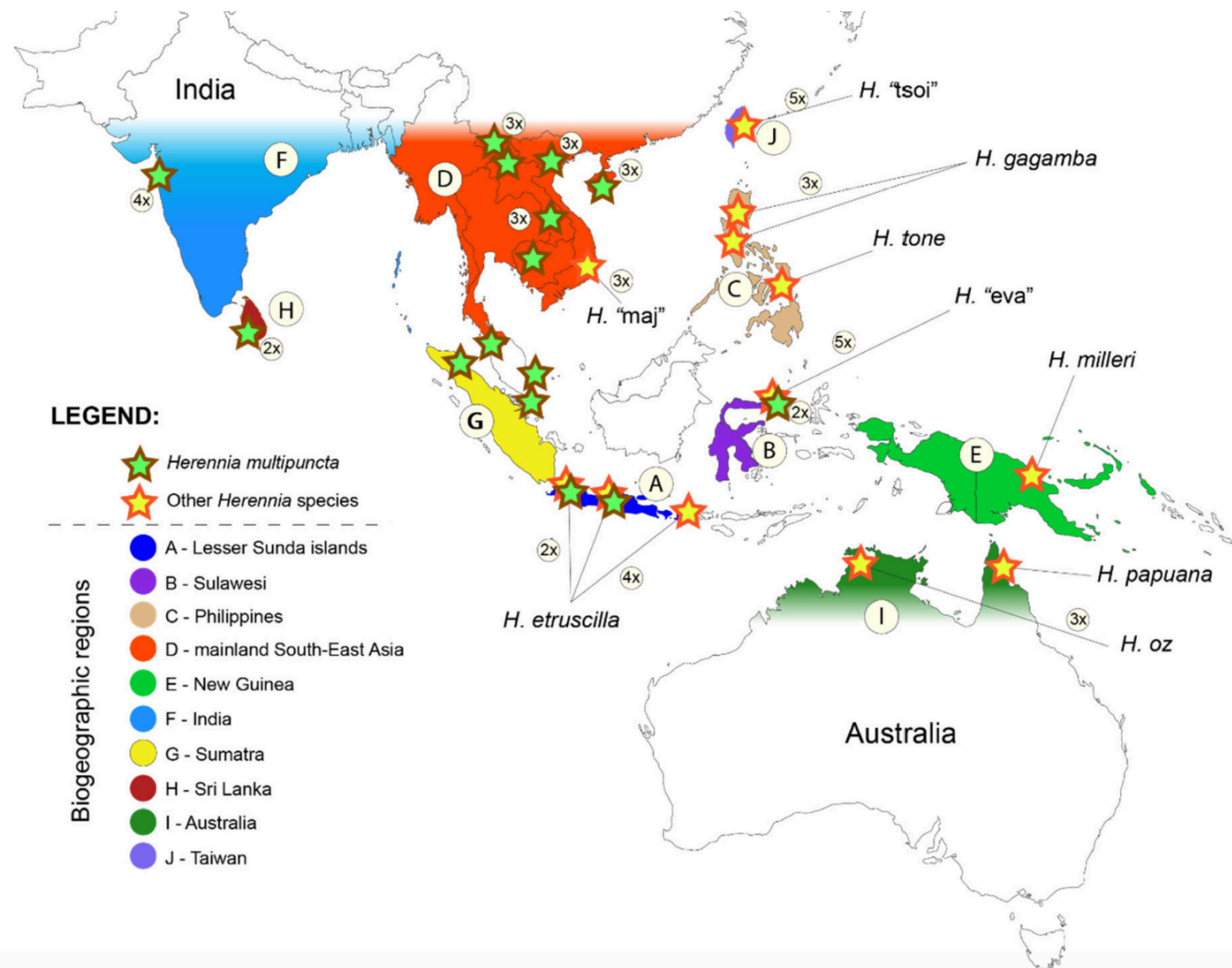
Initial environmental change triggered preadaptation
smaller body size in expanding populations

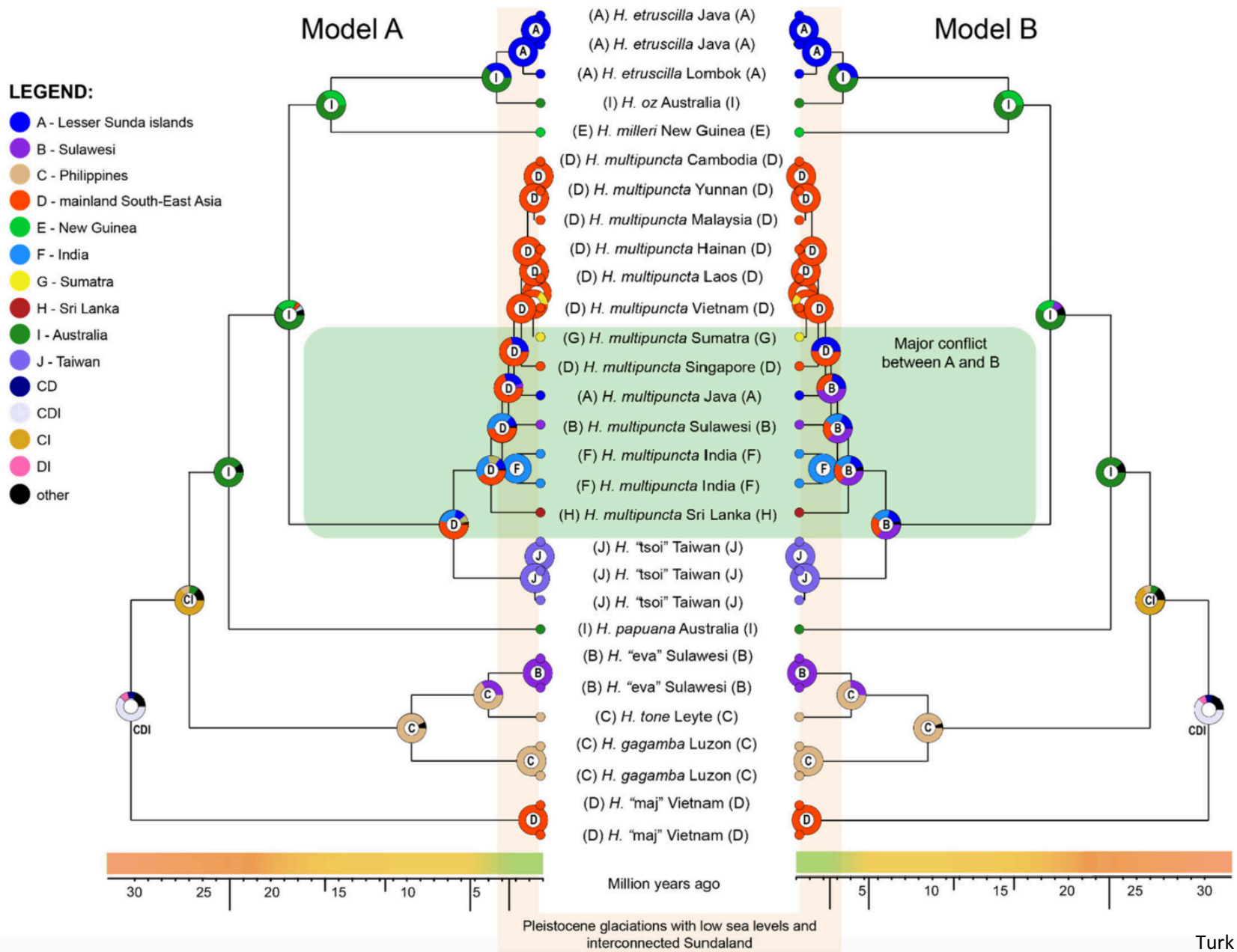


Herennia (Nephilinae) – area expansion or introduction?

Coin spiders presumably do not balloon

Is the Asian distribution range of *H. multipuncta* natural or human-mediated?

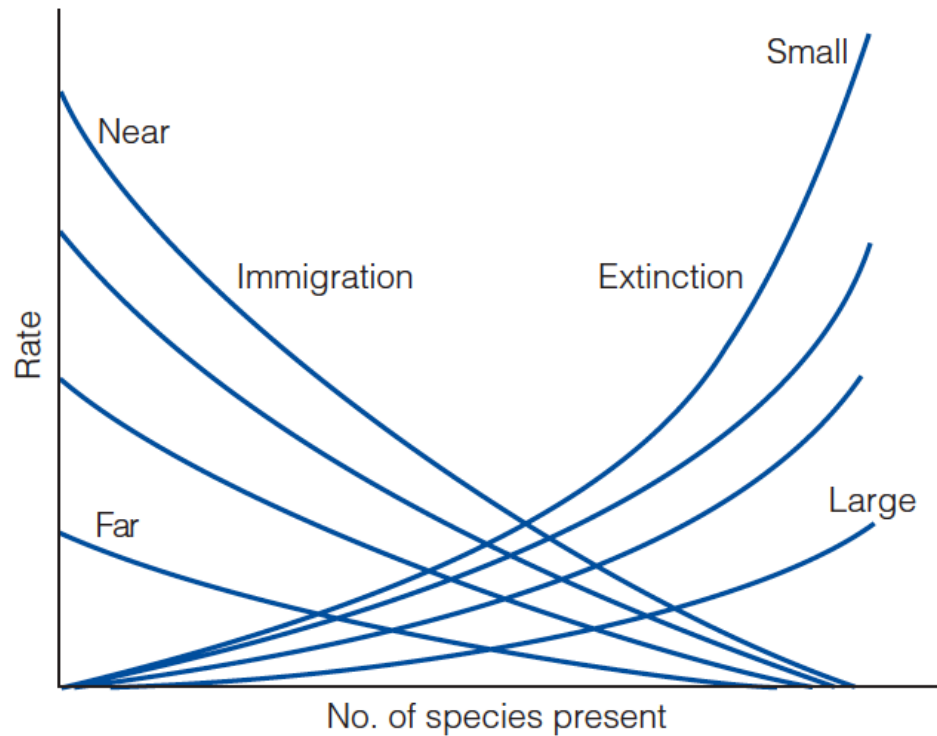




Divergences date back to Pleistocene
 Did *Herennia* regained ballooning capability?

Island biogeography dispersal vs. vicariance

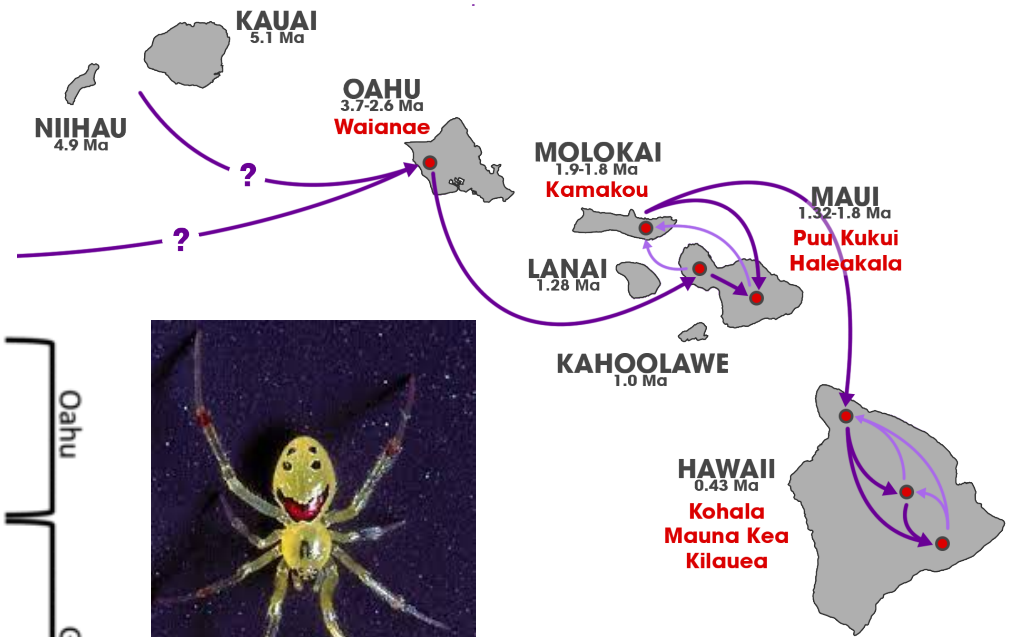
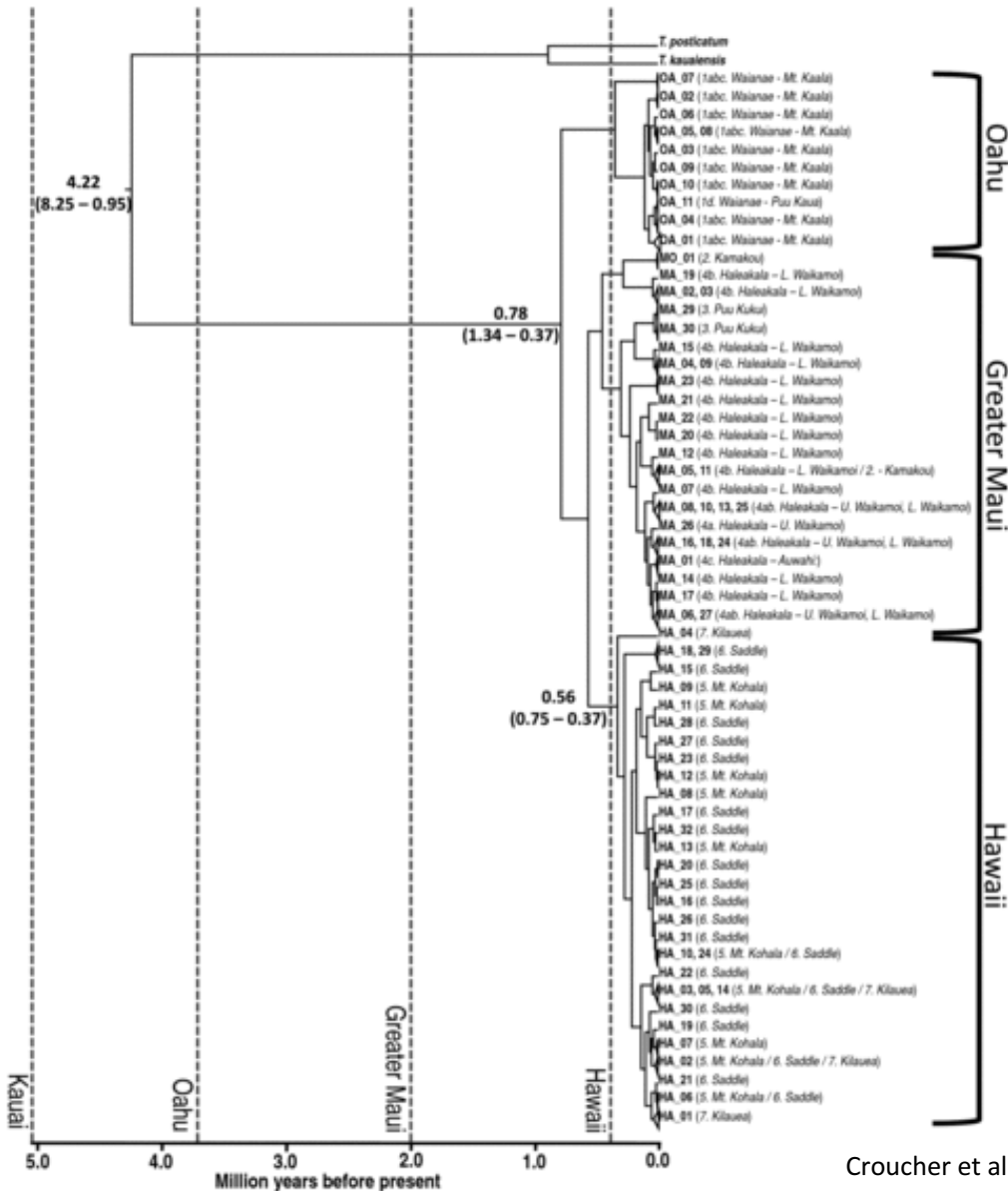
carrying capacity = turnover equilibrium



Cox *et al* 2010

Continental islands – split from a larger landmass; vicariance* + dispersal
Oceanic islands – volcanic *de novo* origin; dispersal

Hawaii *Theridion grallator*

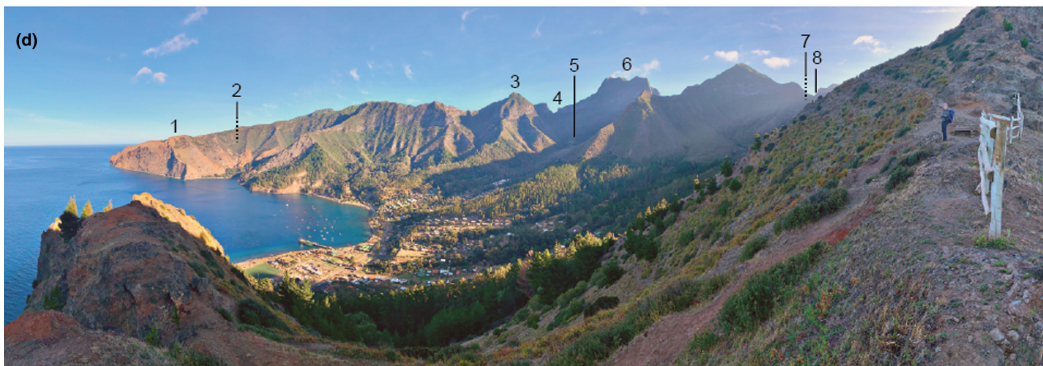
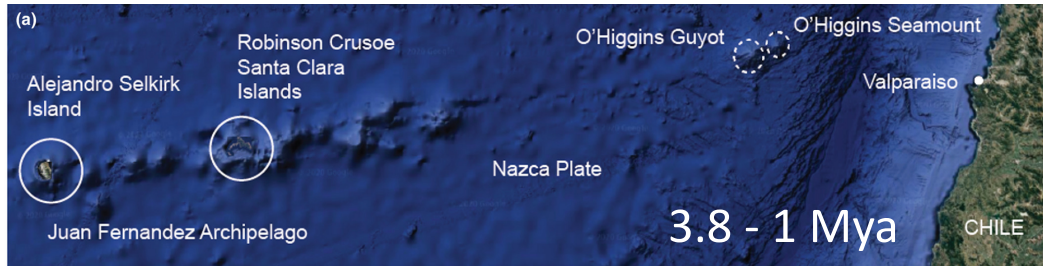


Distinct monophyletic clades
– currently little taxa exchange

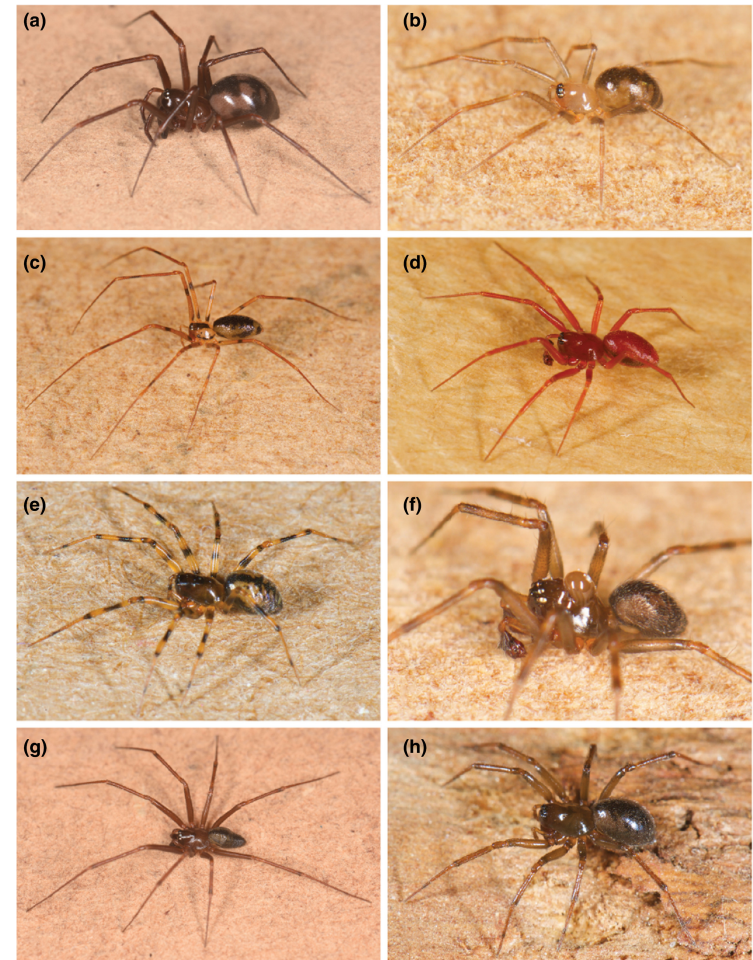
Younger island colonized from older ones
– “progression rule”

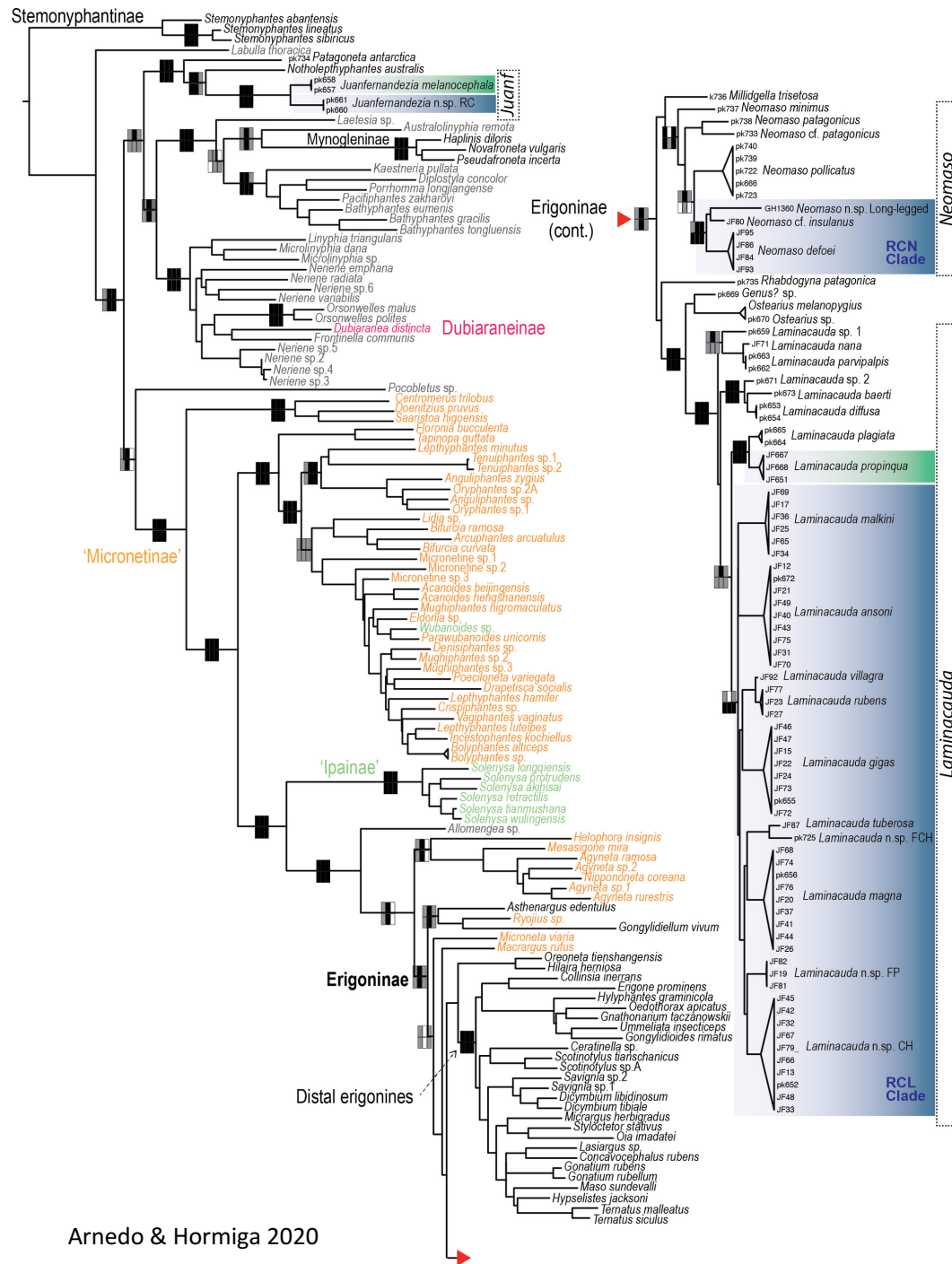
Rapid colonization from Oahu

Juan Fernandez *Linyphiidae*



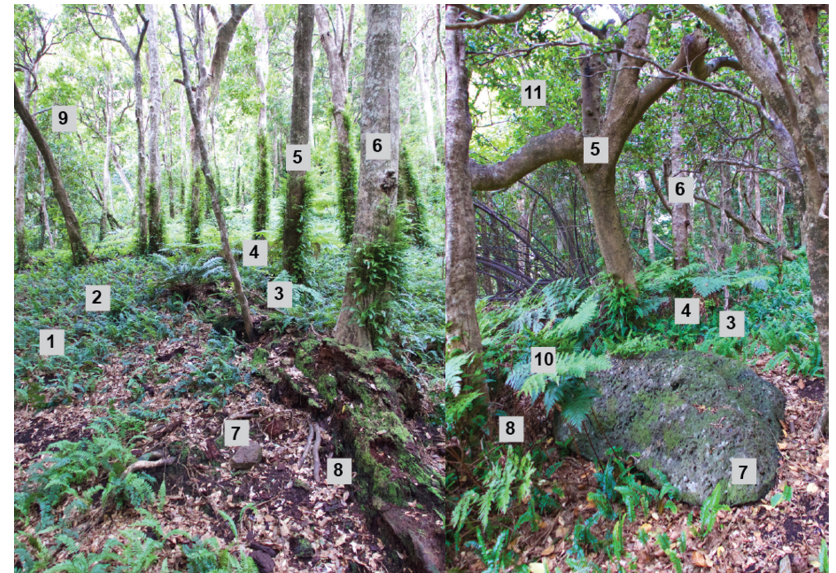
50 native sp. of spiders, ~70 % endemic
40 % Linyphiidae





5 independent colonization events
(*Laminacauda* 2x)

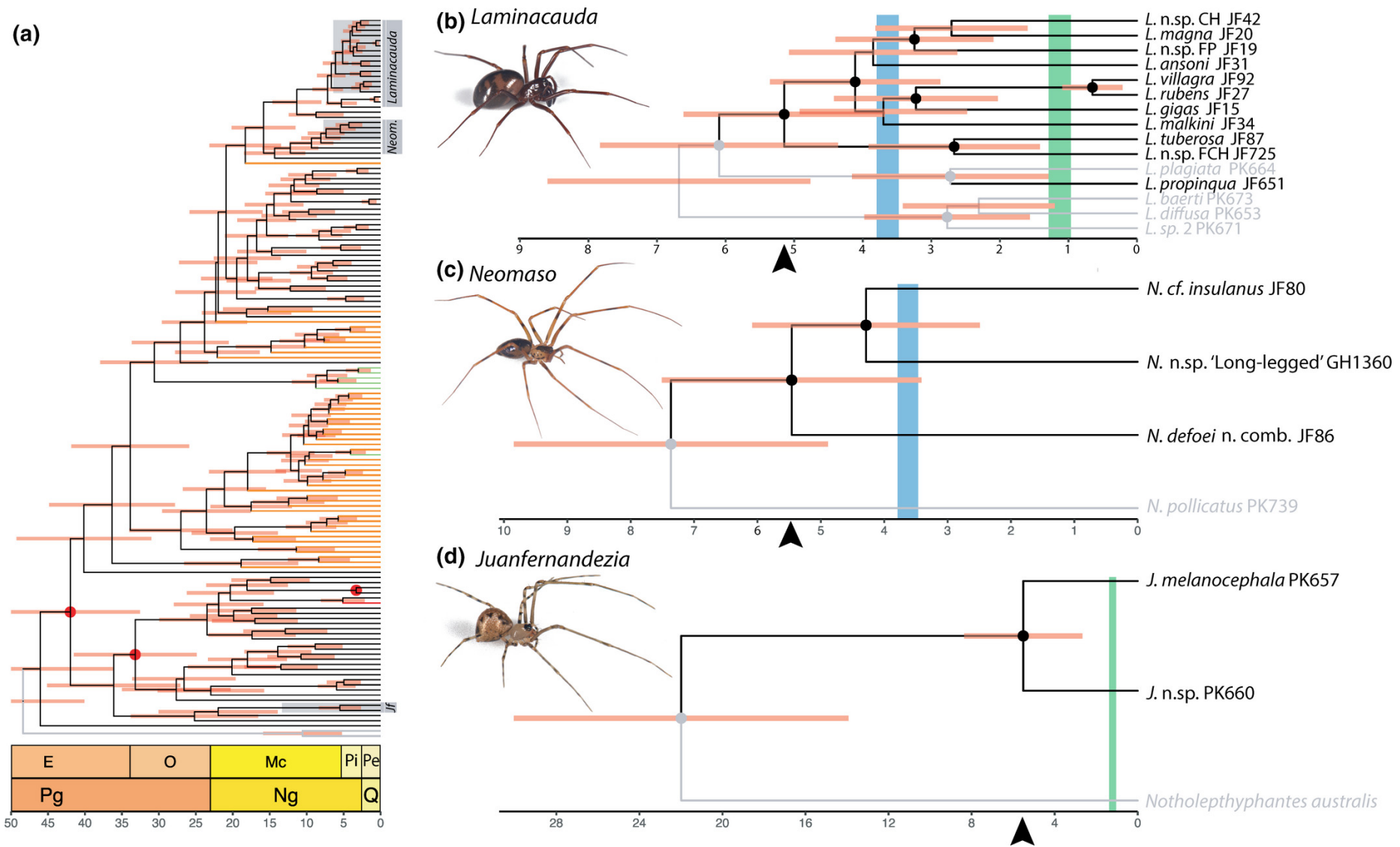
→ adaptive radiation



Laminacauda spp.: 1 = *L. n. sp. CH*; 2 = *L. n. sp. CO*; 3 = *L. n. sp. FC*; 4 = *L. n. sp. FP*; 5 = *L. ansoni*
6 = *L. rubens*; 7 = *L. gigas*; 8 = *L. magna*; 9 = *L. malkini*; 10 = *L. tuberosa*; 11 = *L. villagra*.

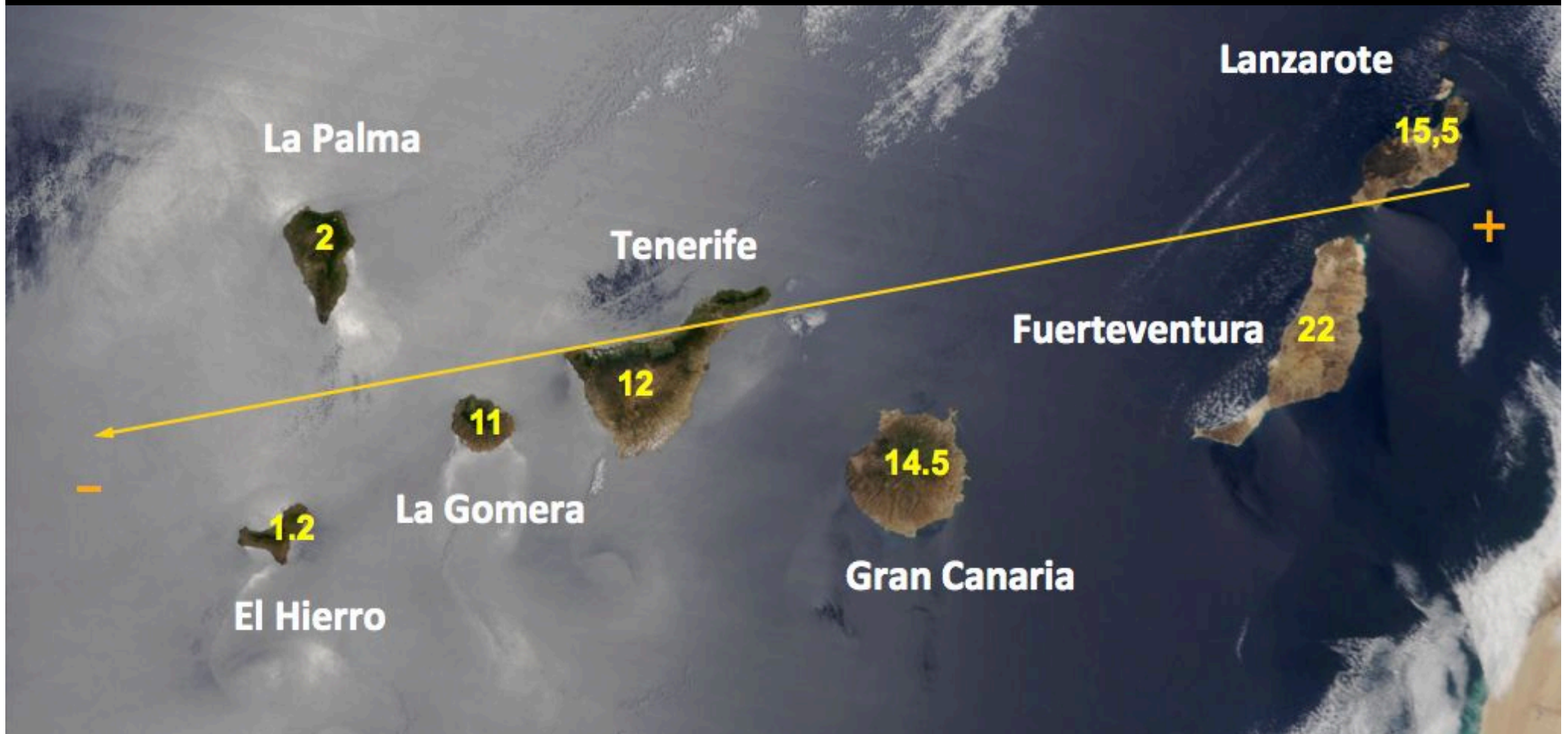
- spatial distribution
- foraging strategy

“Old taxa on young islands dilemma”



Fossil and mt rate calibration in agreement

Canary Islands



high levels of endemic organisms

Canary Islands *Dysdera*

oniscophagous

sedentary

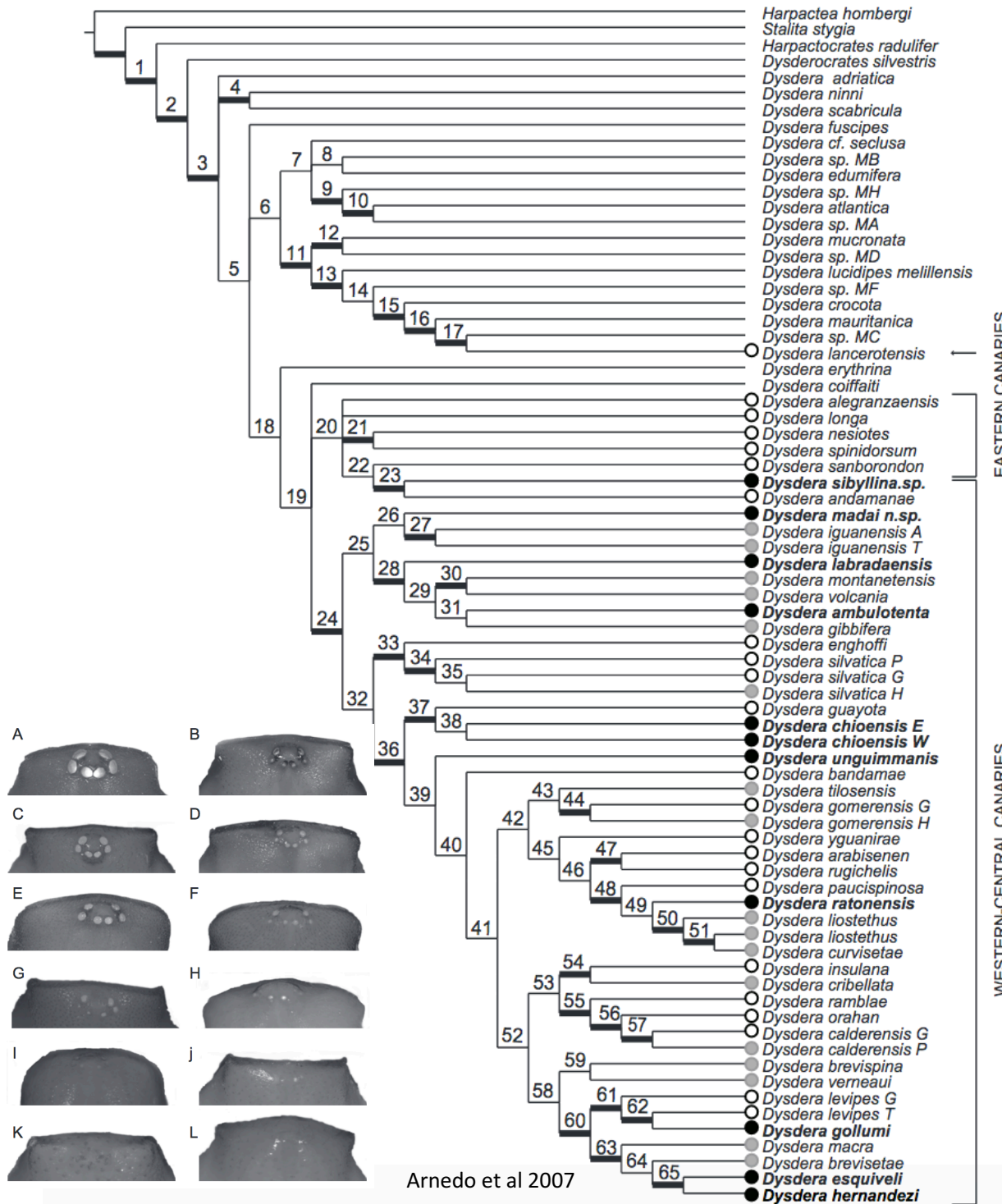
dispersal by rafting



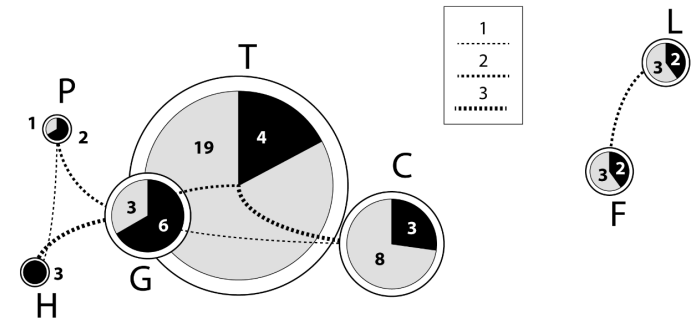
Canary Islands *Dysdera*

48 endemic species

2 x colonization, 1 x radiation

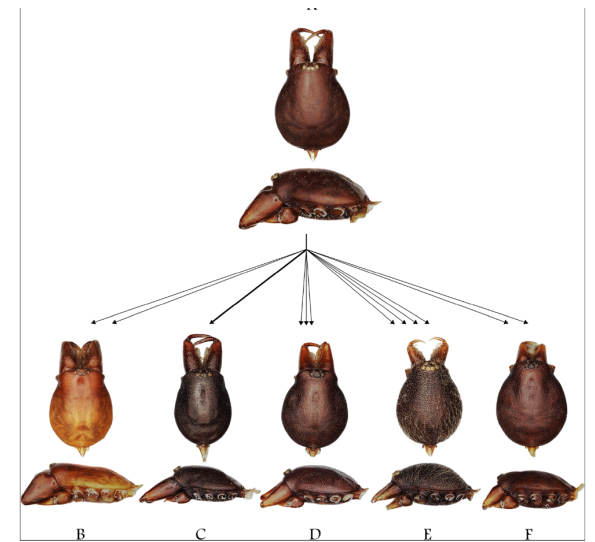
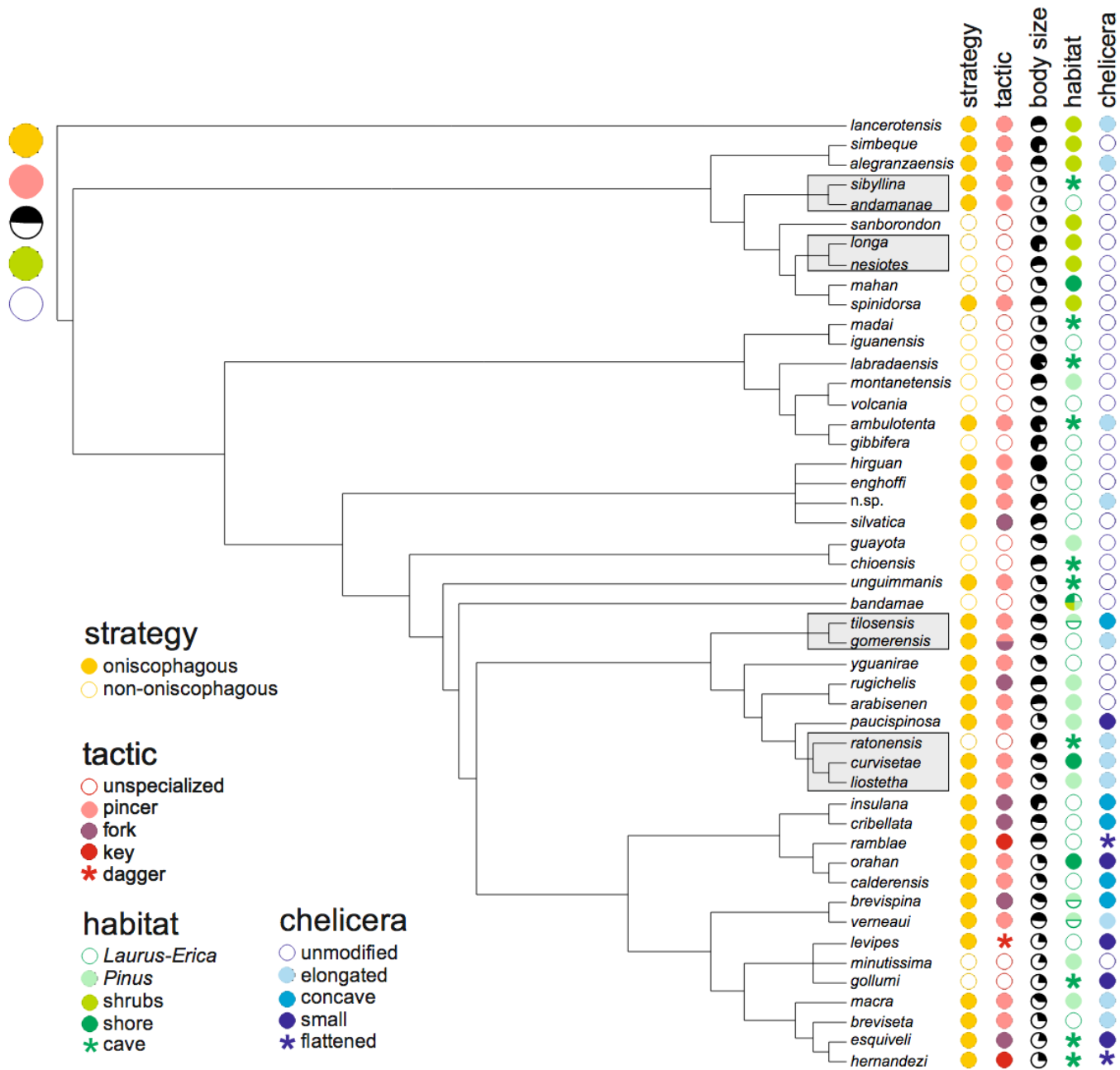


Arnedo et al 2007



black – shared
 grey - endemic

Macías-Hernández et al 2016



Co-occurring species

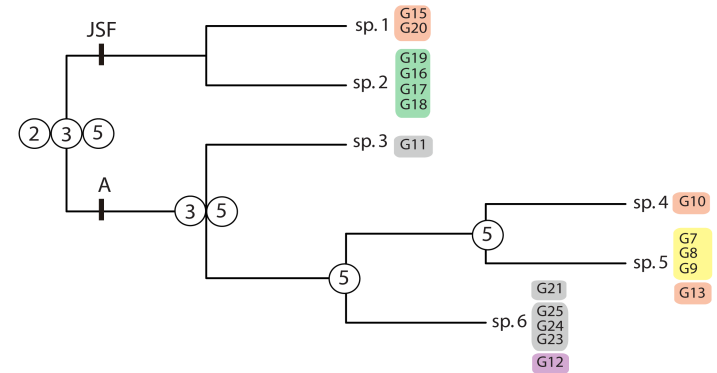
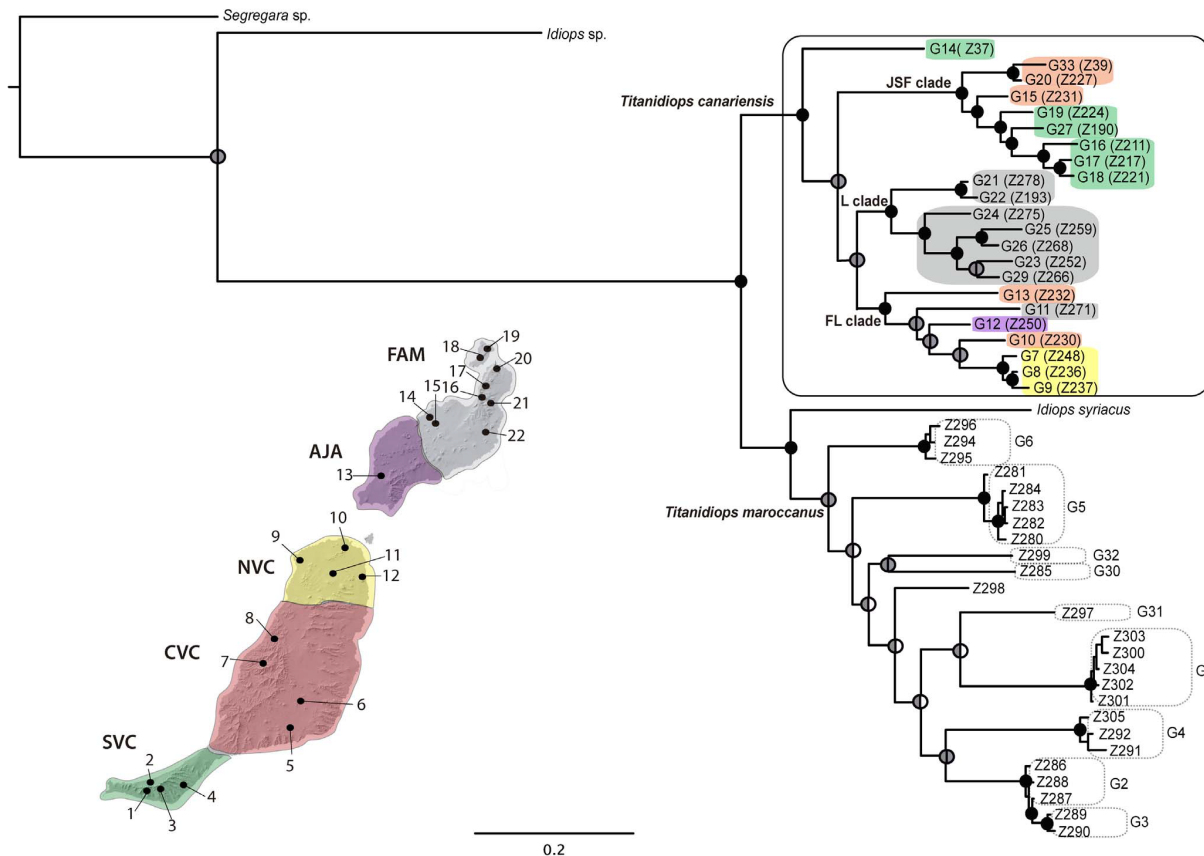
- differences related to prey capture
- different microhabitats

Canary Islands *Titanidiops canariensis*

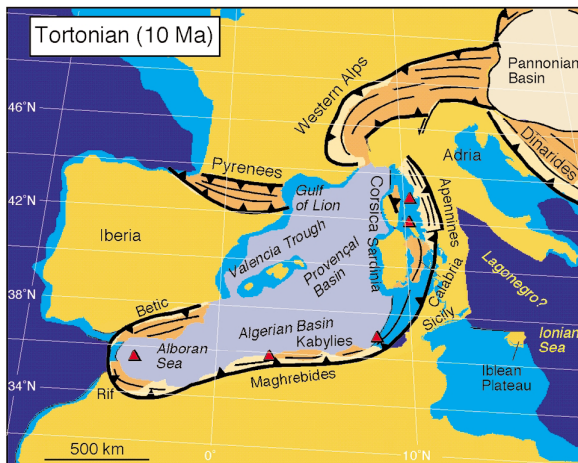
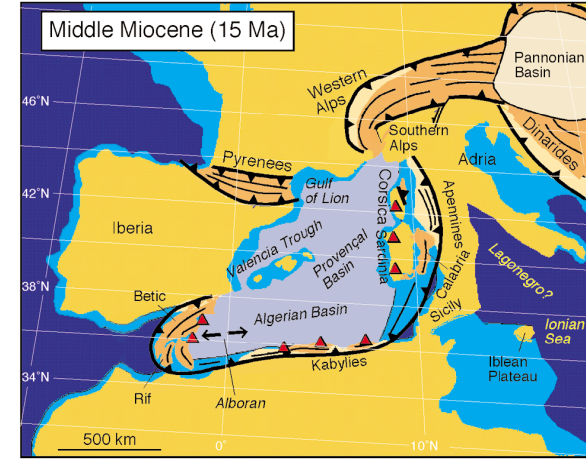
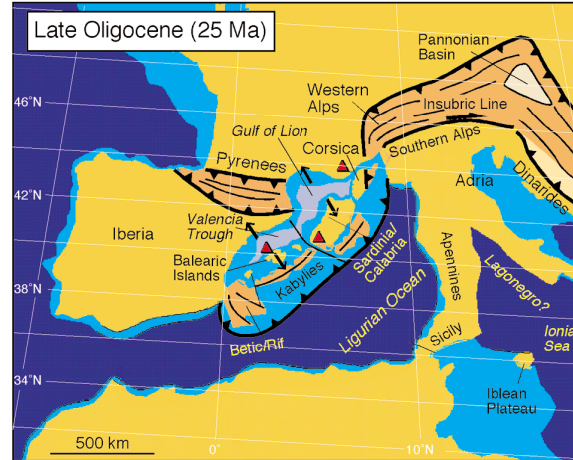
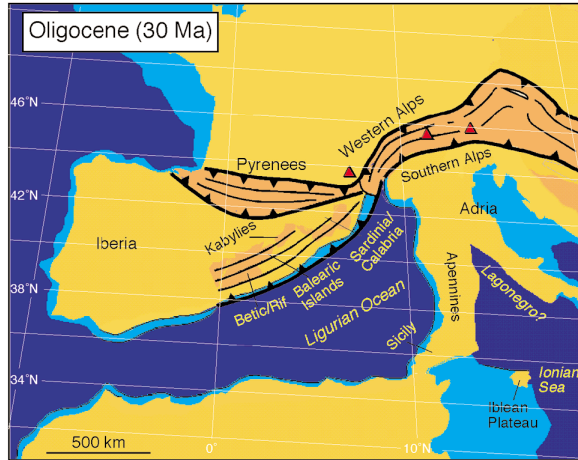
trapdoor spider

sedentary – colonization via rafting, only 1x 7.5 Ma

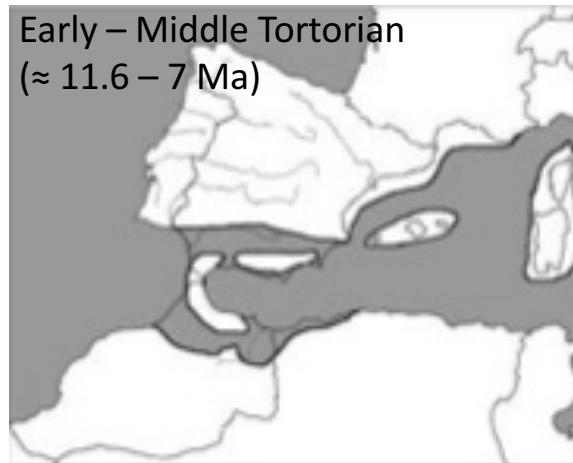
at least 2 cryptic species



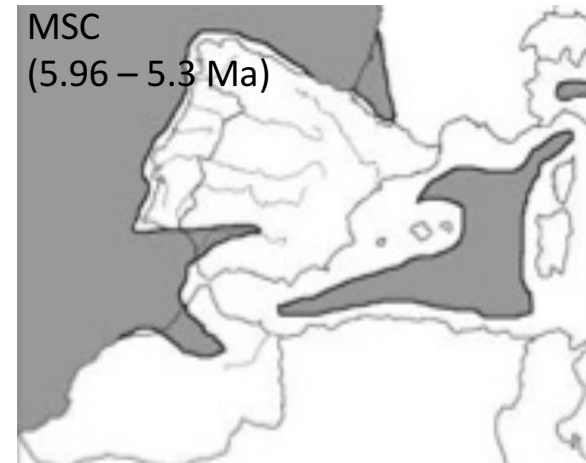
Continental Islands – geological history of the Western Mediterranean



Rosenbaum et al. 2002



Paulo et al. 2008



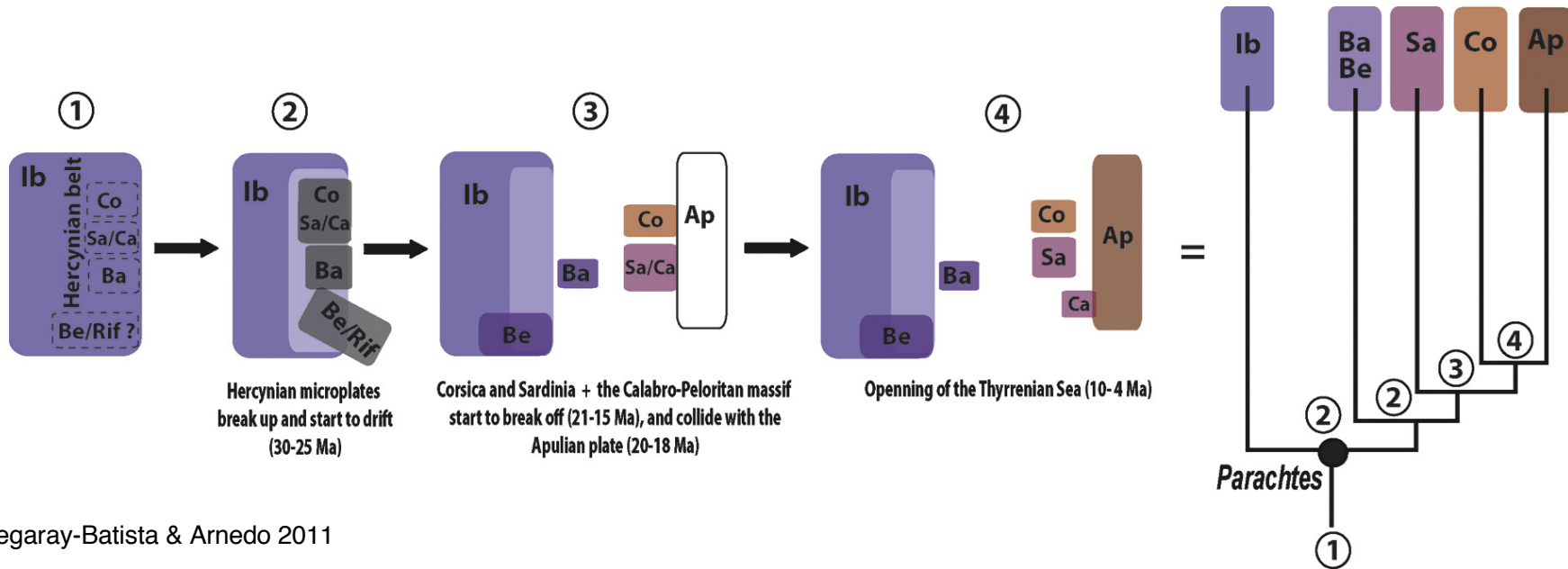
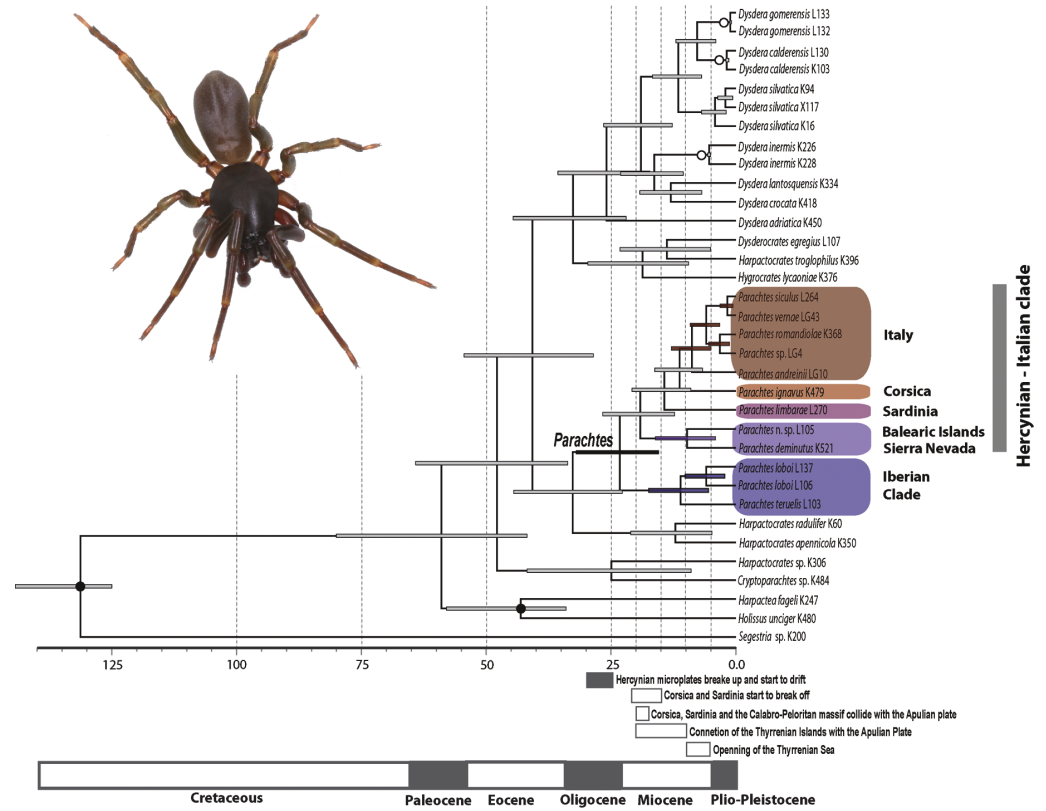
Continental Islands – *Parachtes*

Dysderidae (generalist)

Sedentary

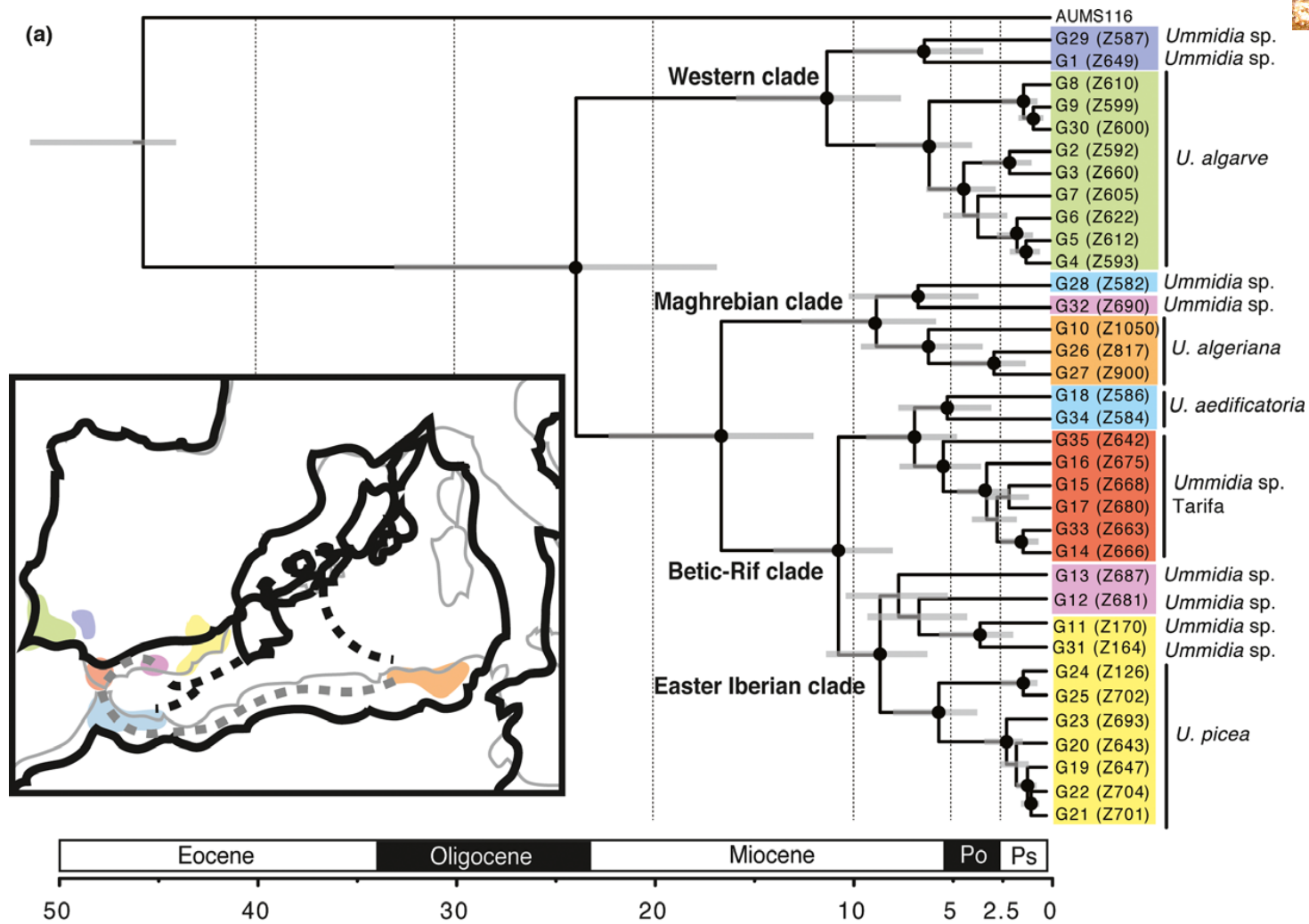


Hercynian belt breakup



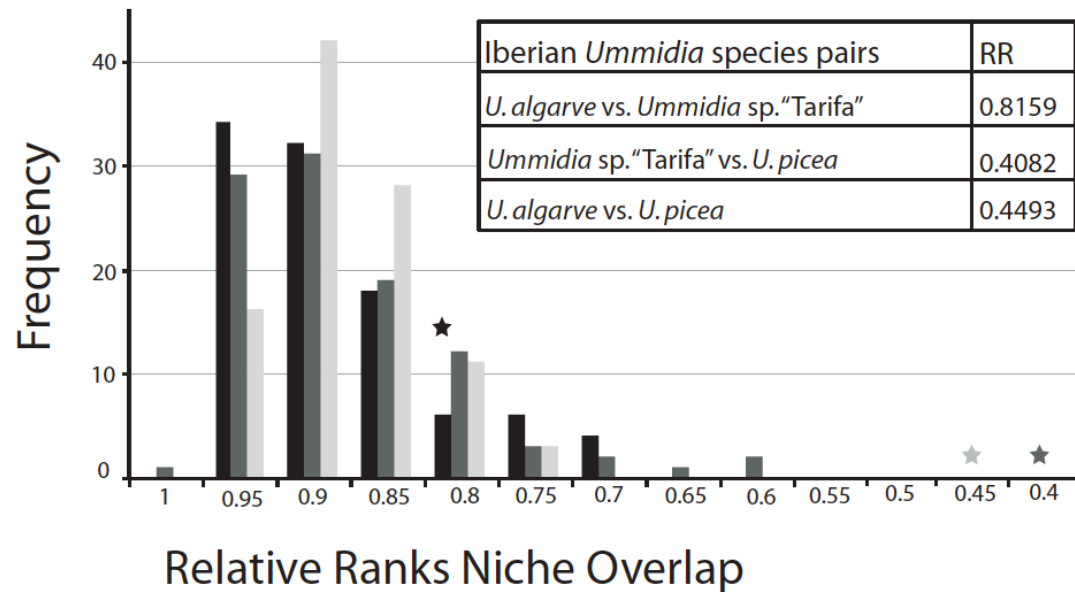
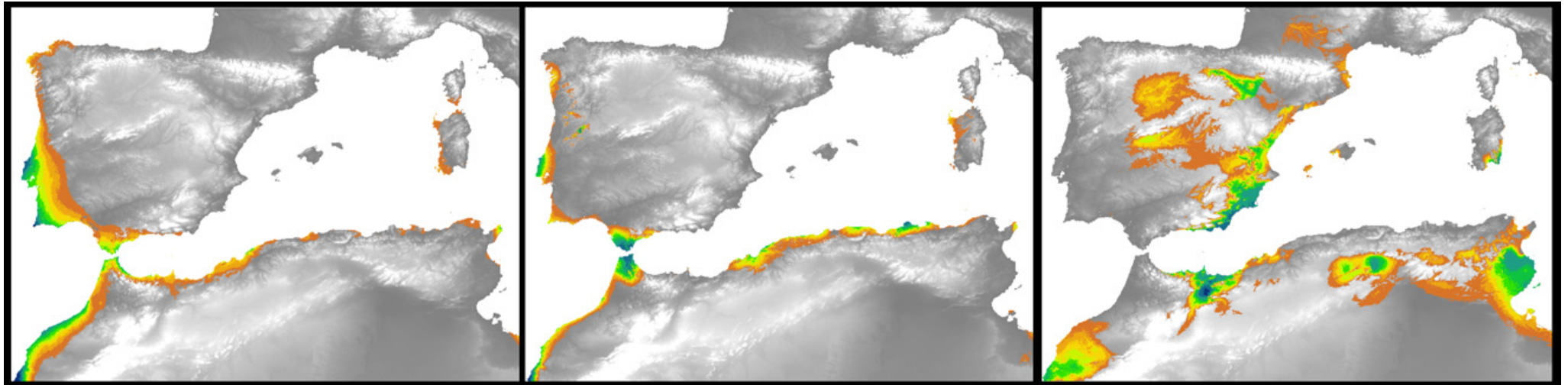
Continental Islands – *Ummidia*

Trapdoor spider with ballooning capability; mostly vicariance



Distribution modelling – assessing ballooning capabilities

Niche overlap, but no gene flow



Pangea formation

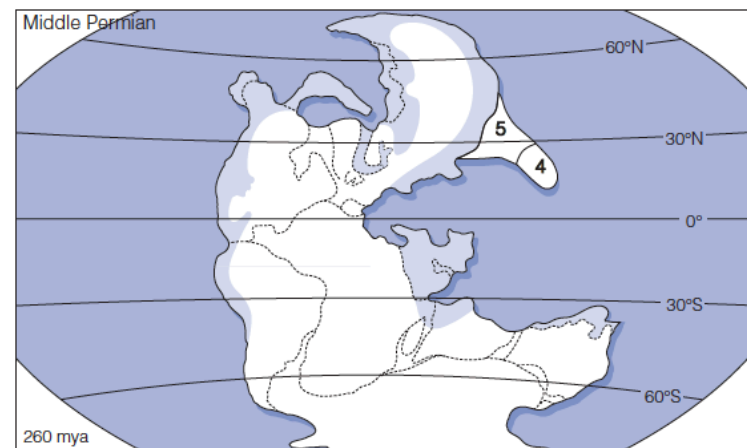
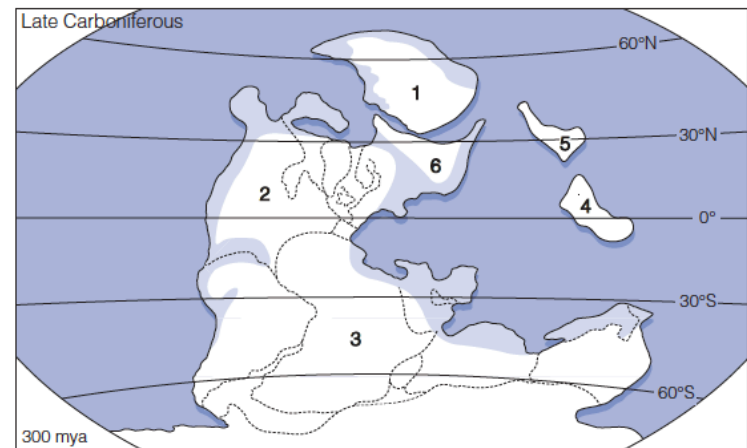
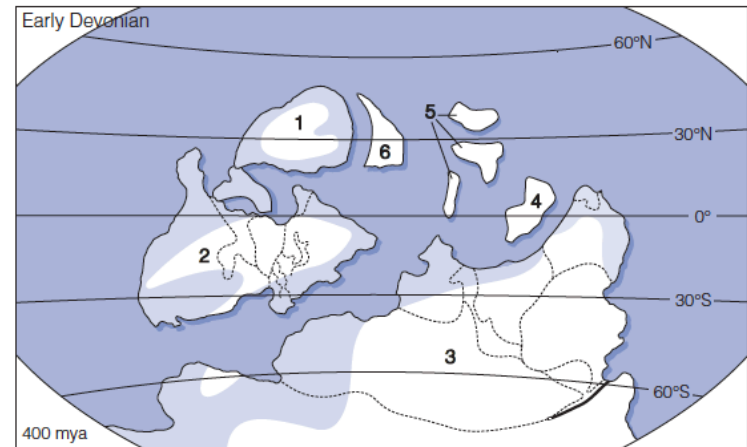
Amalgamation of Gondwana 520 – 510 Ma
- southern hemisphere

Laurentia, Baltica, Siberia – in the north
- formed Laurasia in Paleozoic, ~ 300 Ma

Late Paleozoic – Pangea formation
- lasted ~ 100 Myr

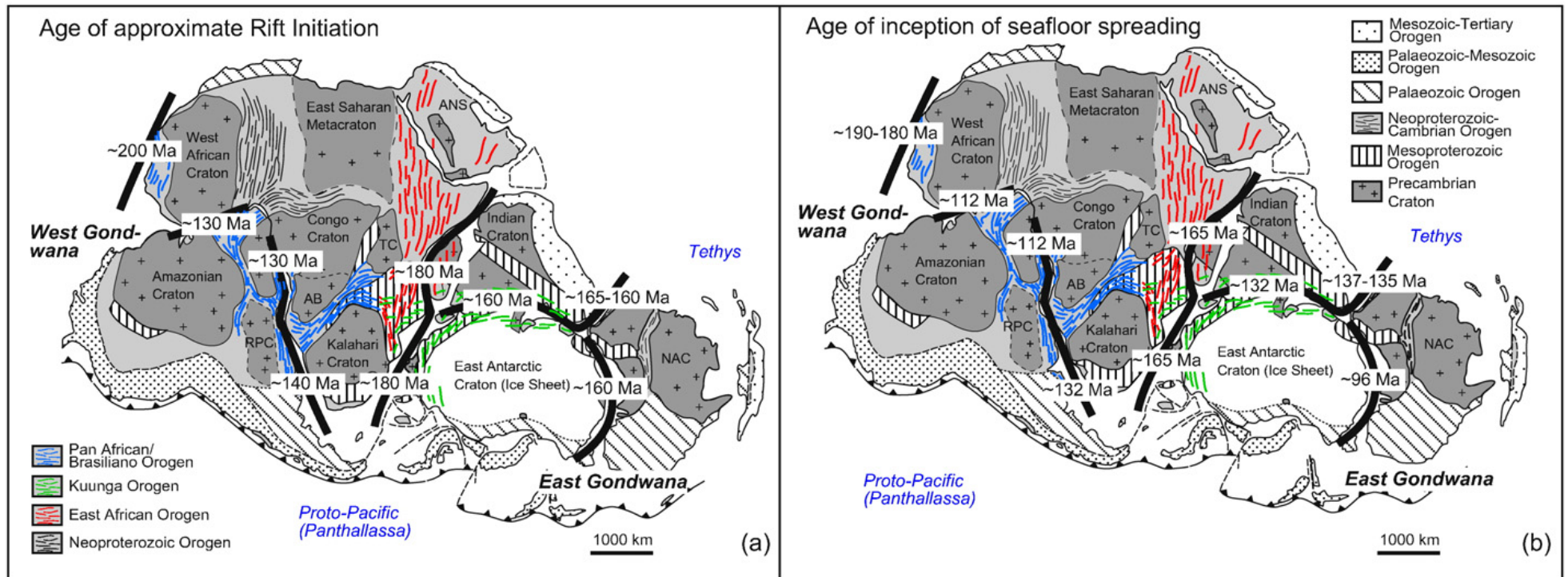
Pangea breakup
- Central Atlantic ridge ~ 200 Ma

Will & Frimmel 2018



Cox et al 2010

Gondwana disintegration



breakup from Laurasia

- Central Atlantic ridge ~ 200 Ma

Will & Frimmel 2018

Lower Jurassic ~ 180 Ma East/ West Gondwana breakup

Upper Jurassic ~ 160 Ma India-Madagascar/Antarctica

~ 160 Ma Antarctica/Australia

Lower Cretaceous ~ 140 – 130 Ma S America/Africa

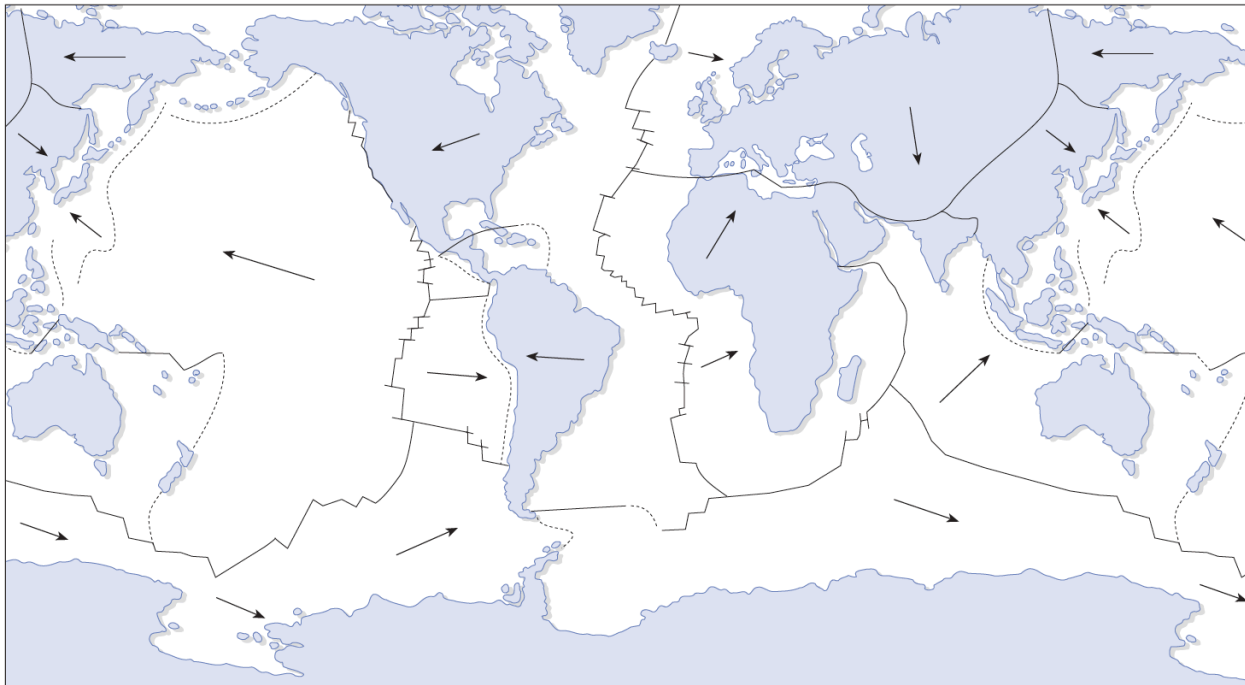
Upper Cretaceous ~ 80 - 90 Ma Madagascar/ India

Laurasia breakup ~ 55 Ma

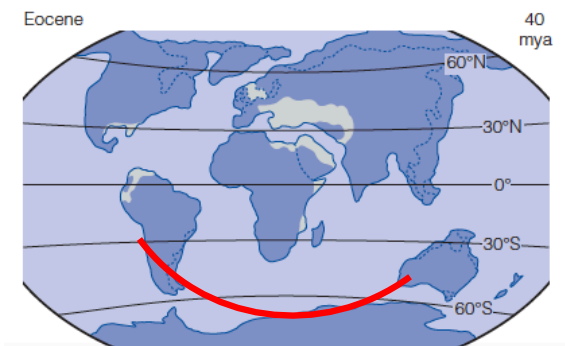
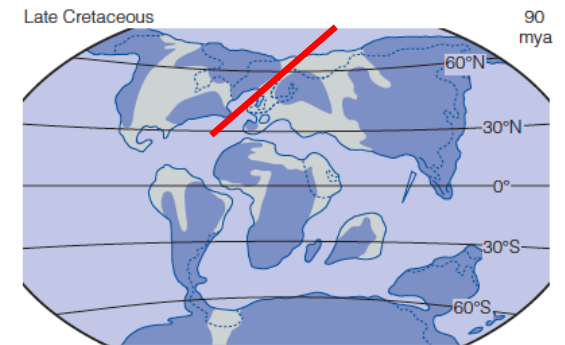
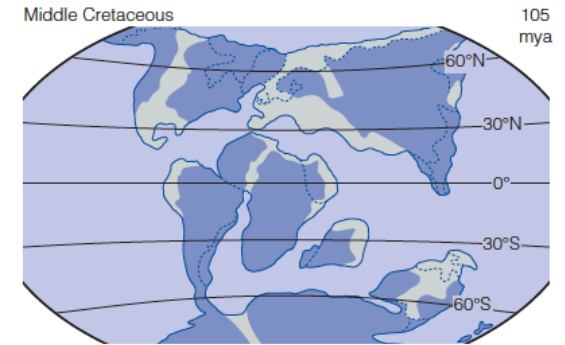
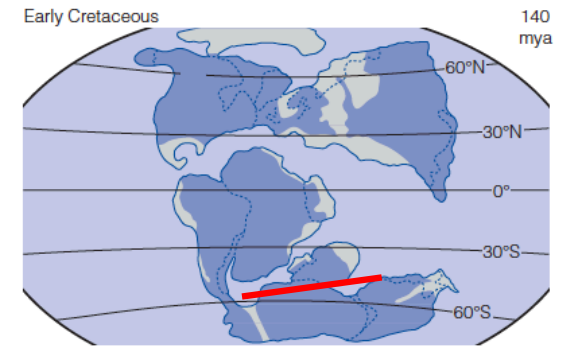
S America – Antarctica – Australia land bridge
up to ~ 30 Ma

timing updated continuously, controversial topics remain

The movement continues ~ 5 – 10 cm/yr



Cox et al 2010



Cox et al 2010

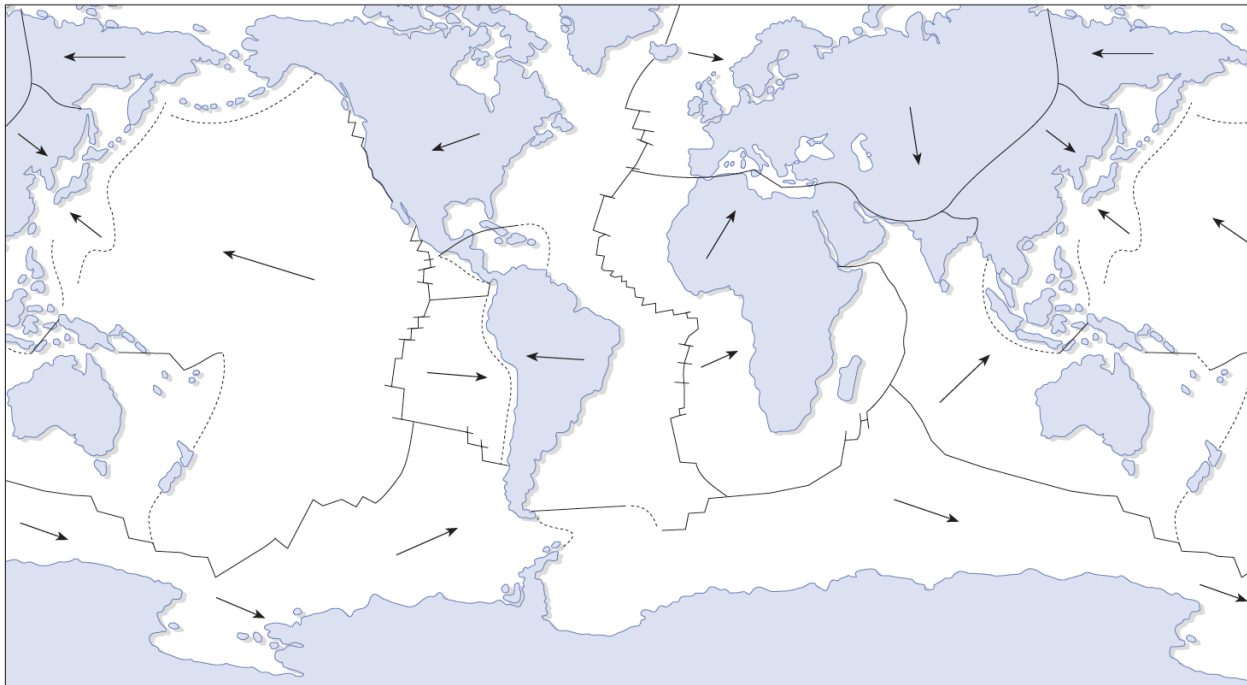
Will & Frimmel 2018

Laurasia breakup ~ 55 Ma, land bridge ~ 25 Ma

S America – Antarctica – Australia land bridge
up to ~ 30 Ma

timing updated continuously, controversial topics remain

The movement continues ~ 5 – 10 cm/yr



Cox *et al* 2010

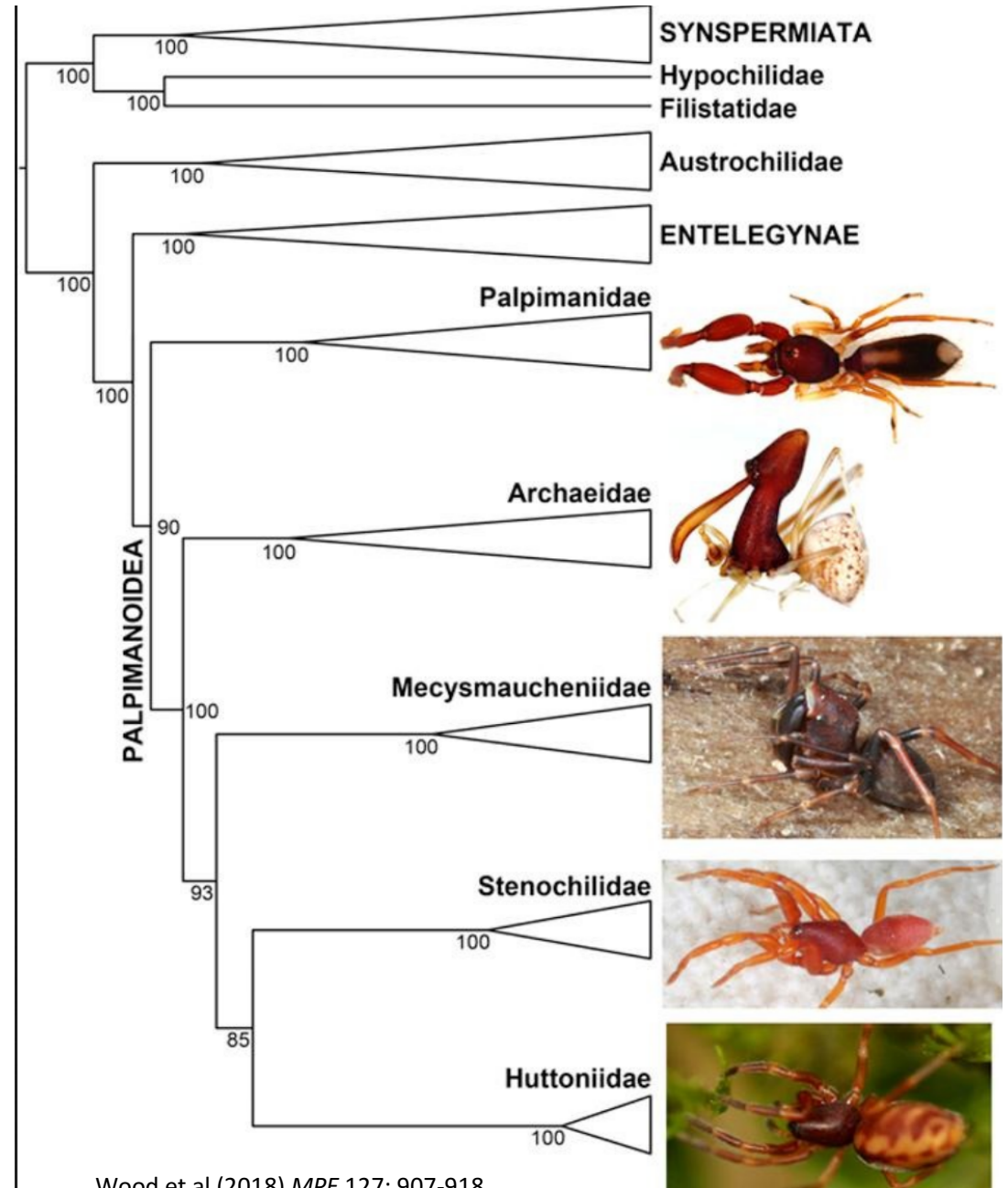
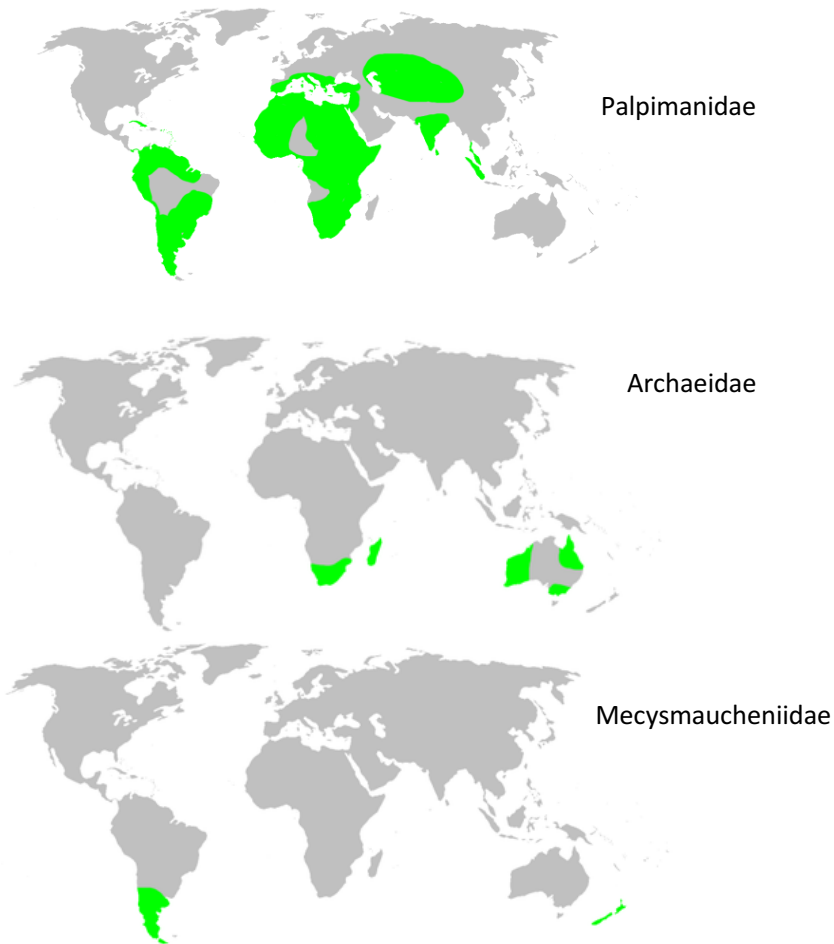


Ms. A. Williams
730 cm/30 yr

Palpimanoidea continental vicariance?

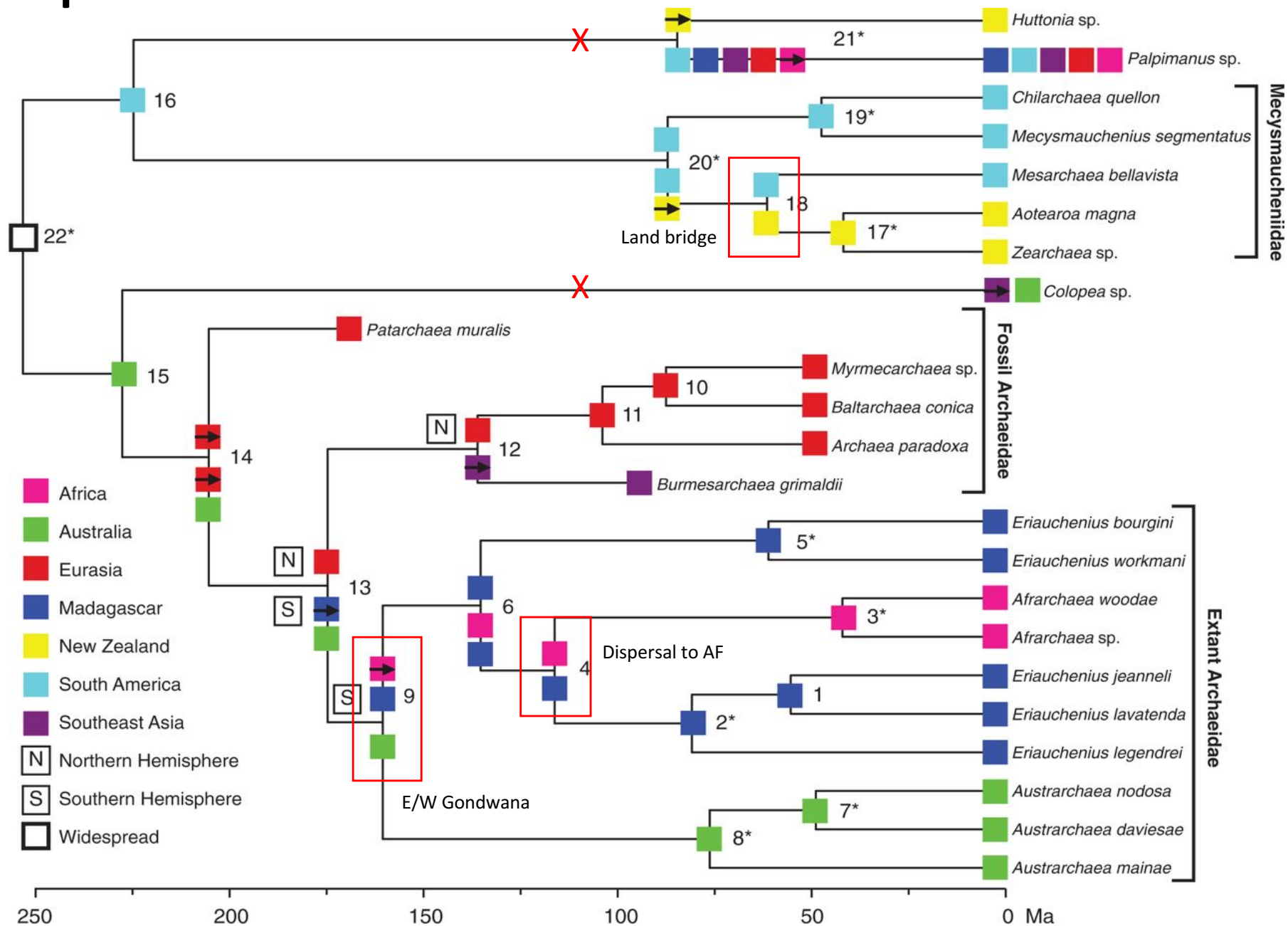
~ Gondwanan distribution

araneophagous: modifications

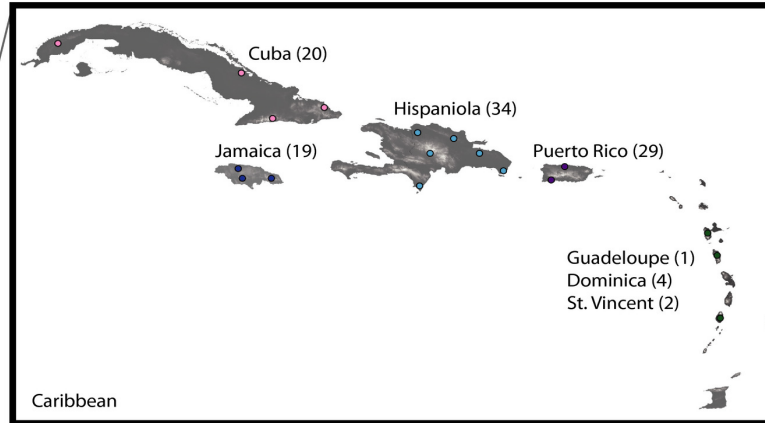
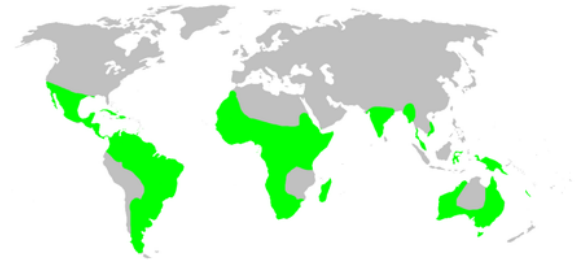


Wood et al (2018) *MPE* 127: 907-918.

Palpimanoidea continental vicariance?



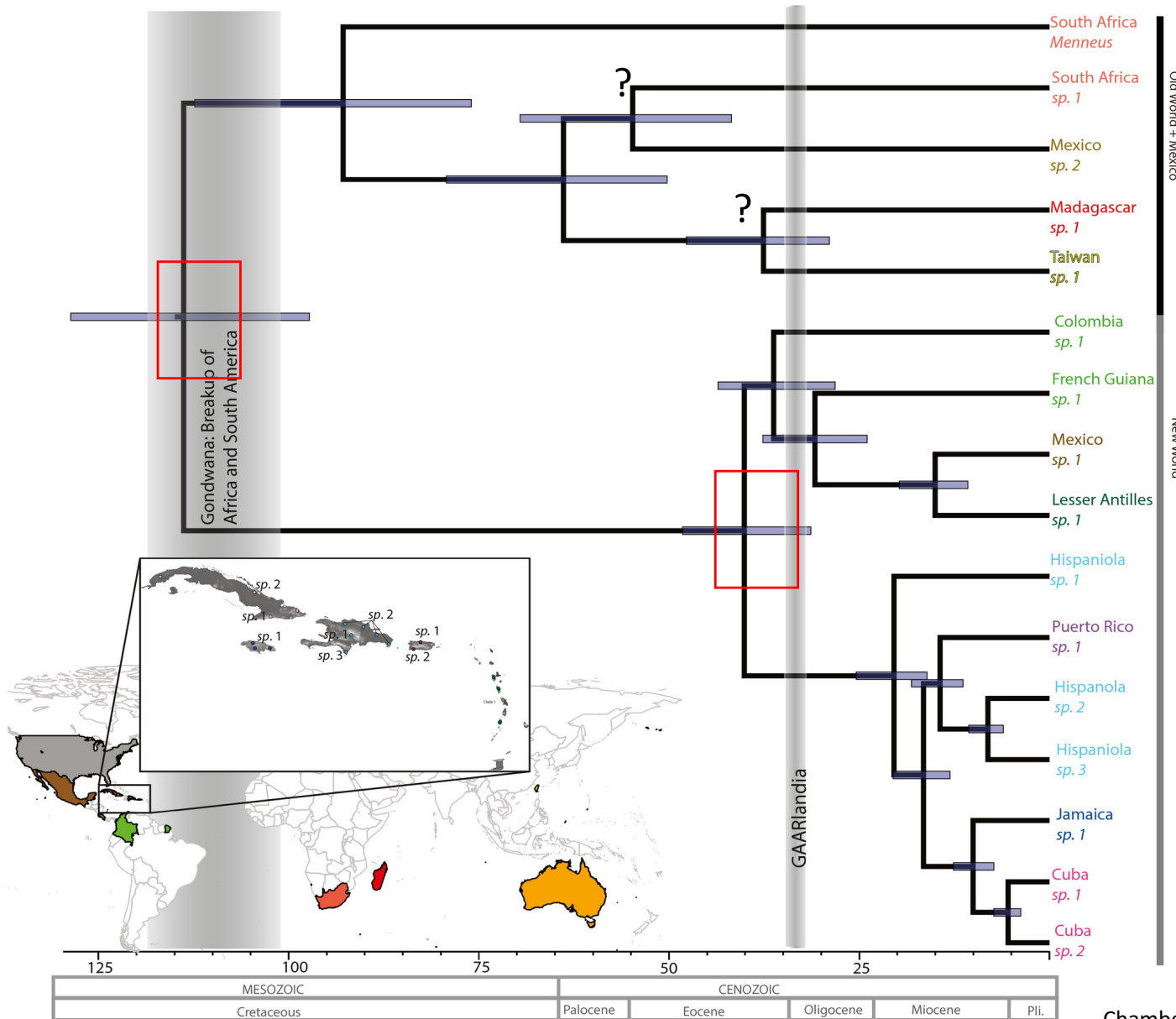
Deinopsis continental vicariance? GAARlandia land bridge



Greater Antilles and Aves Ridge

- land bridge connecting S America with the Greater Antilles
- Eocene – Oligocene ~ 35 – 33 Ma

Deinopsis continental vicariance? GAARlandia land bridge



Gondwanan origin

? long-distance dispersal

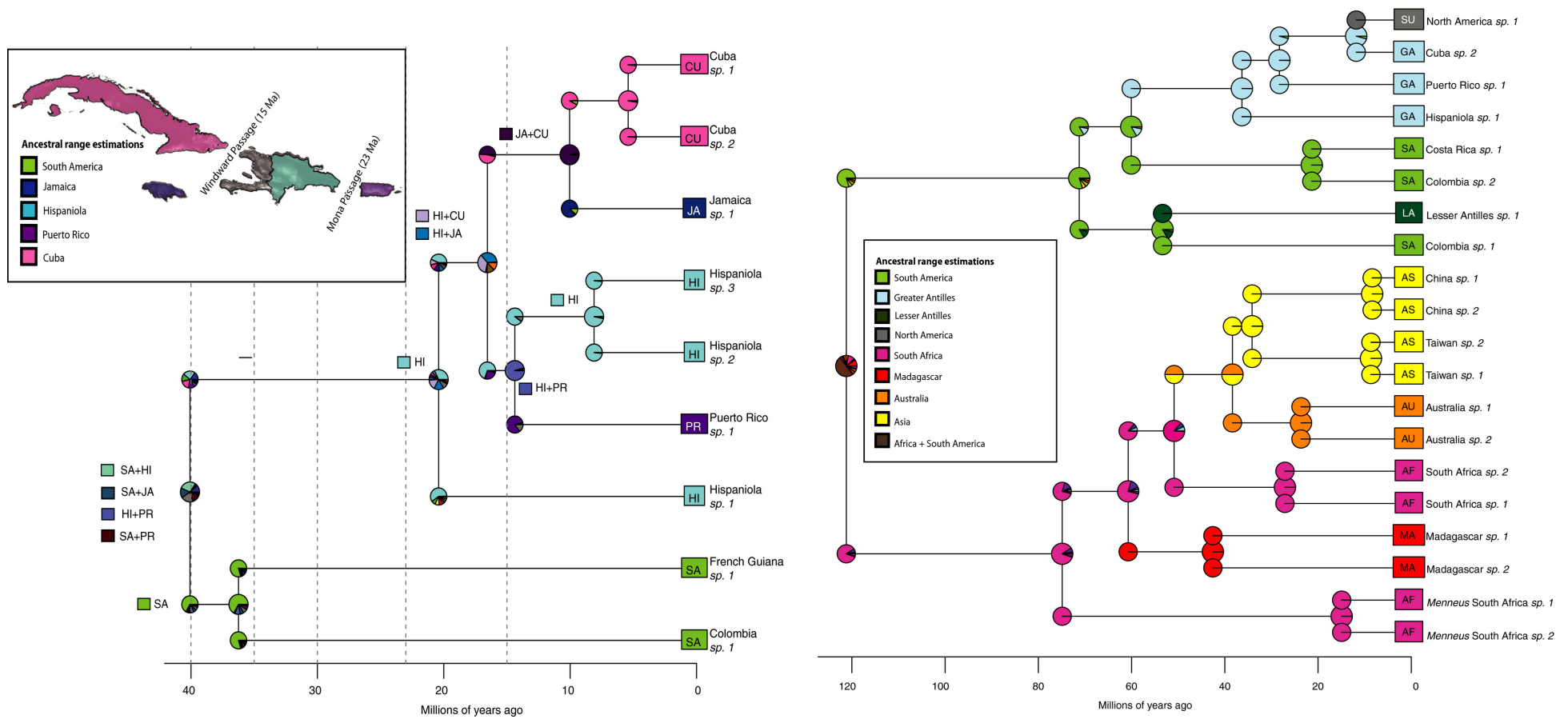
Supports GAARlandia

Also in: *Loxosceles*, *Sicarius*
Heteroctenus scorpions

Not in: *Tetragnatha*,
Selenops

Binford et al (2008)
Crews & Esposito (2020)
Čandek et al (2021)

Deinopis continental vicariance? GAARlandia land bridge

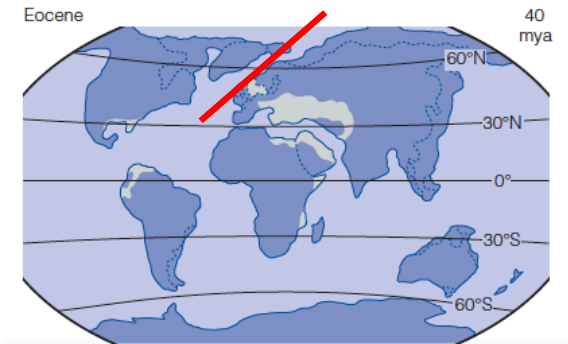
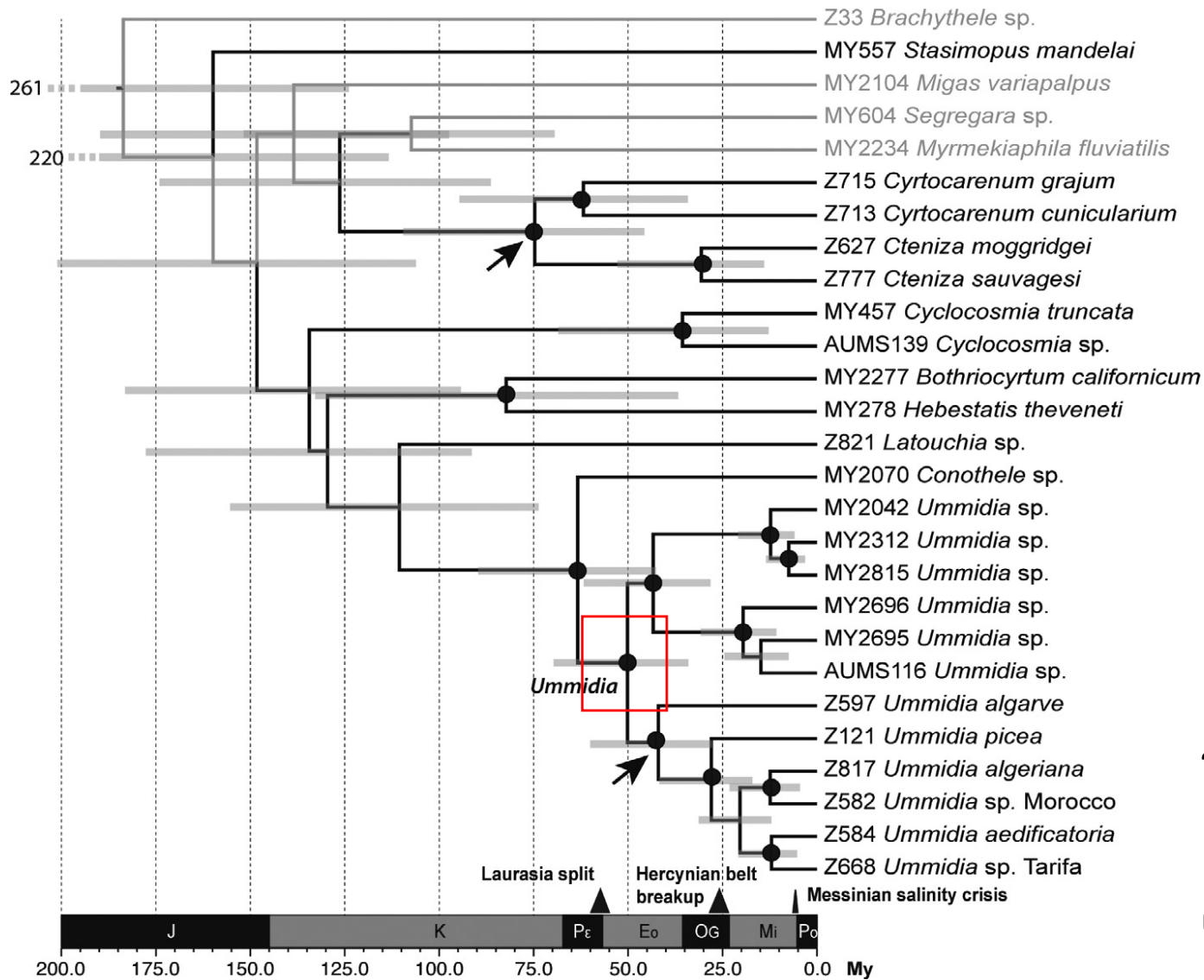


Chamberland et al (2018)

Greater Antilles colonized 1x from S America
 → back colonization
 African origin of the Old World taxa

Ummidia continental vicariance?

Halonoproctidae – Laurasia breakup



Laurasia breakup
~ 55 Ma

North American land bridge
up to ~ 25 ma

Macrothele continental vicariance?

M. calpeiana

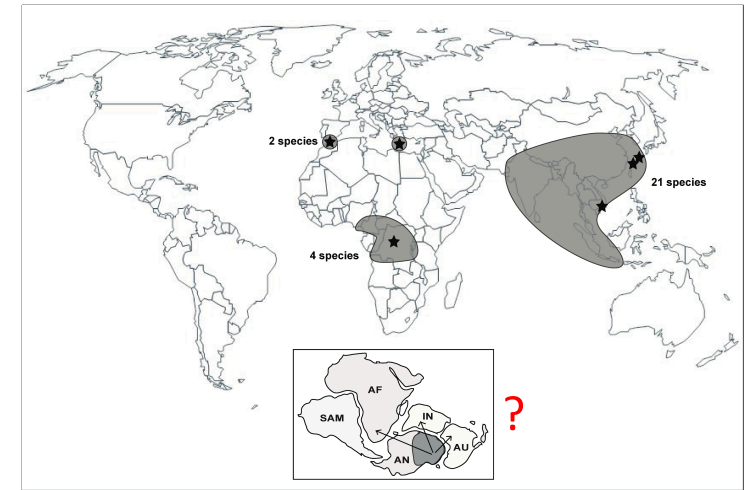
Bern Convention (1987)

EU Habitat Directive (1992)

- Species of community interest, need of strict protection

M. cretica IUCN red list, data deficient category

both considered sedentary



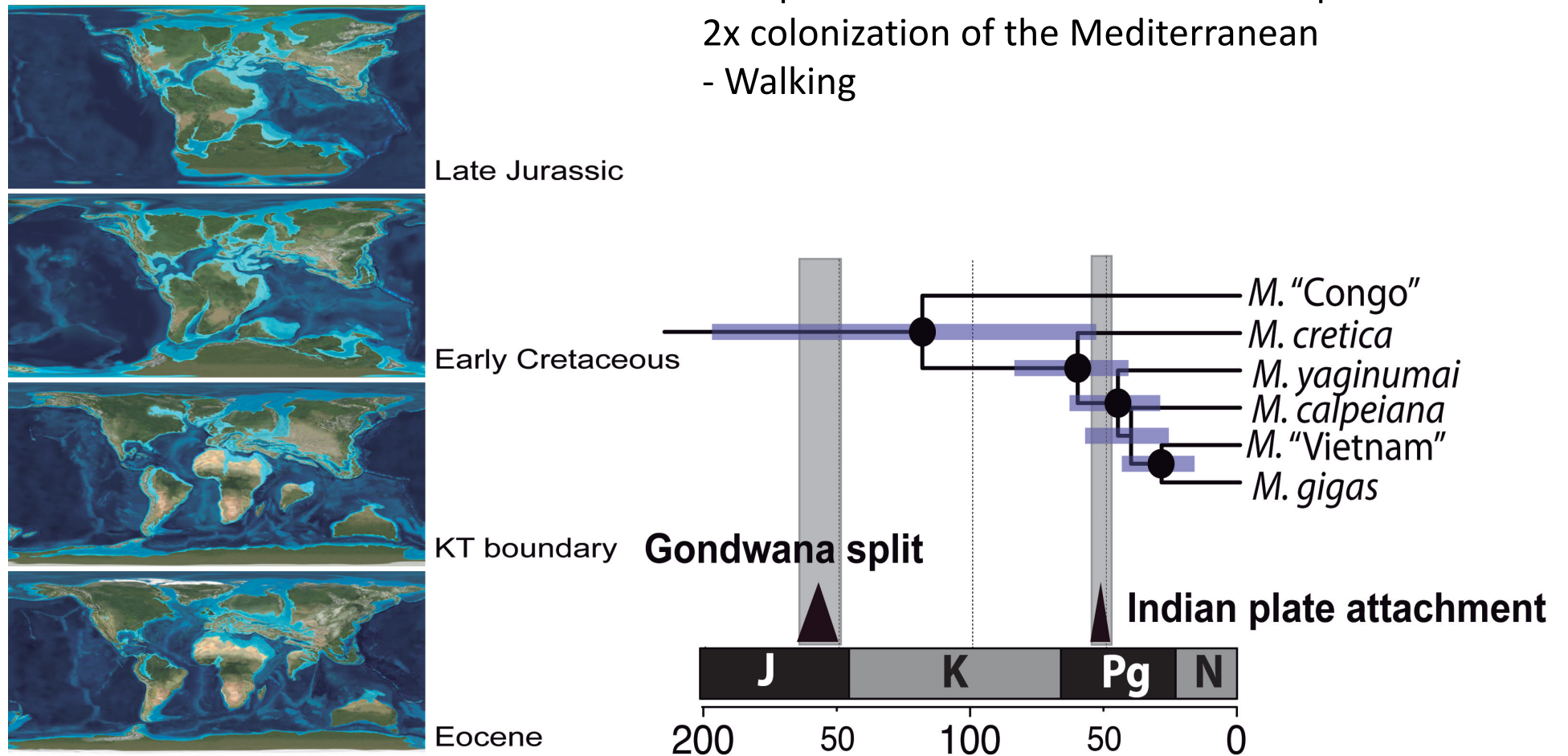
Macrothelidae



Macrothele continental vicariance?

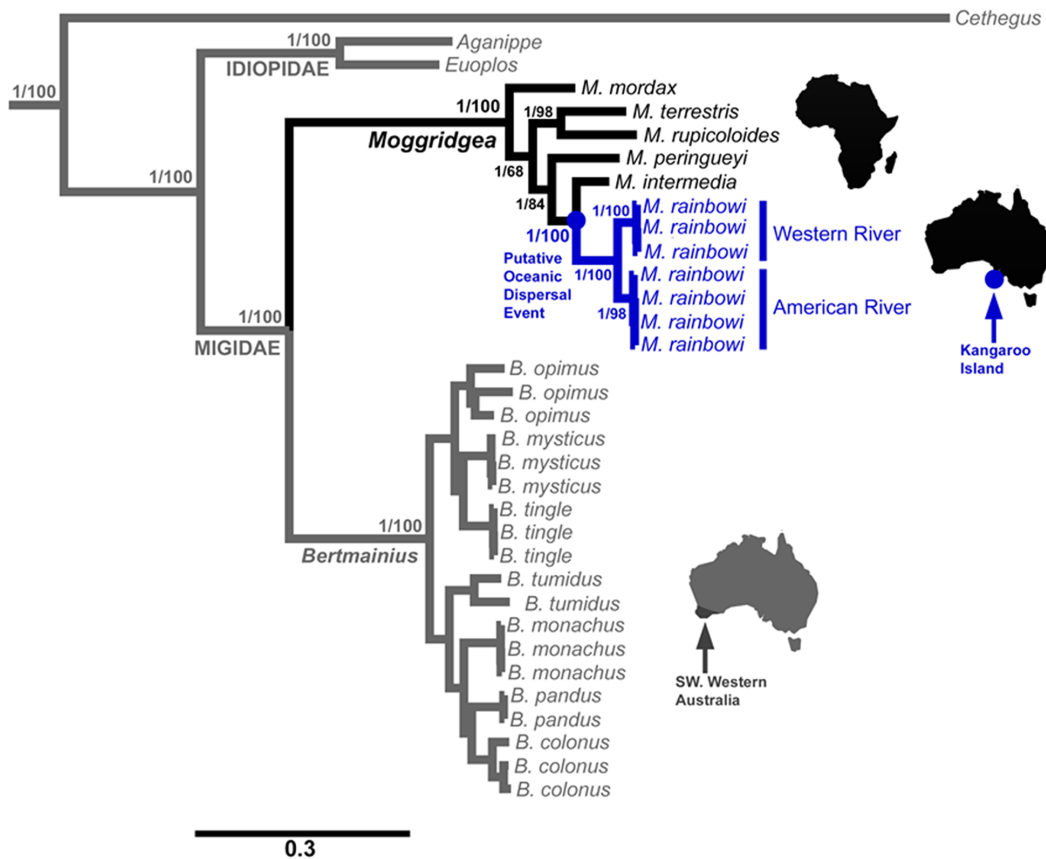
Gondwana breakup

European Macrothele are not sister species
2x colonization of the Mediterranean
- Walking



Moggridgea continental vicariance?

Migidae – Gondwanan distribution



Parameters	TMRCa Node 1 (Moggridgea Dispersal) + 95% Highest Posterior Density	TMRCa Node 2 (KI Population Divergence) + 95% Highest Posterior Density	Posterior Mean	Posterior ESS
Strict Clock, GTR Models, Speciation: Yule Process	16.02 8.87–25.60	6.39 3.48–10.23	-6059.22	20.04 X
Strict Clock, HKY Models, Speciation: Yule Process	2.27	1.10	-6478.87	1449.97
Strict Clock, GTR Models, Speciation: Birth-Death Process	15.98 8.63–25.96	6.35 3.55–10.37	-5813.92	8.93 X
Strict Clock, HKY Models, Speciation: Birth-Death Process	2.27	1.10	-6259.82	1672.67
Exponential Clock, GTR Models, Speciation: Yule Process	10.59 4.01–19.94	4.06 1.59–7.66	-6021.34	5.93 X
Exponential Clock, HKY Models, Speciation: Yule Process	3.54 2.35–4.96	1.75 1.17–2.45	-6155.69	2310.32
Exponential Clock, GTR Models, Speciation: Birth-Death Process	10.49 3.97–19.71	3.69 1.60–7.45	-5789.97	7.62 X
Exponential Clock, HKY Models, Speciation: Birth-Death Process	3.54 2.36–4.92	1.73 1.16–2.40	-5936.87	1607.35
Lognormal Clock, GTR Models, Speciation: Yule Process	15.44 5.36–27.15	5.96 1.62–11.13	-6035.79	9.76 X
Lognormal Clock, HKY Models, Speciation: Yule Process	8.48 3.33–13.97	3.56 1.25–6.53	-6172.48	1701.01
Lognormal Clock, GTR Models, Speciation: Birth-Death Process	15.40 5.41–27.32	5.77 1.51–10.63	-5821.38	10.17 X
Lognormal Clock, HKY Models, Speciation: Birth-Death Process	8.48 3.32–14.12	3.47 1.22–6.37	-5953.30	1825.83

<https://doi.org/10.1371/journal.pone.0180139.t002>

2.27 – 8.48 Ma

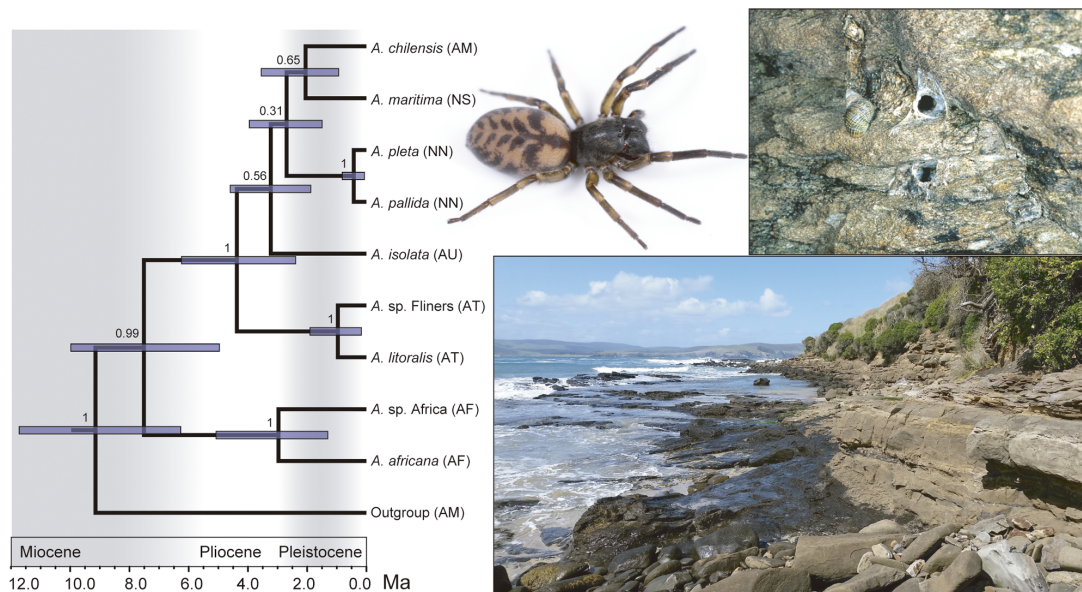
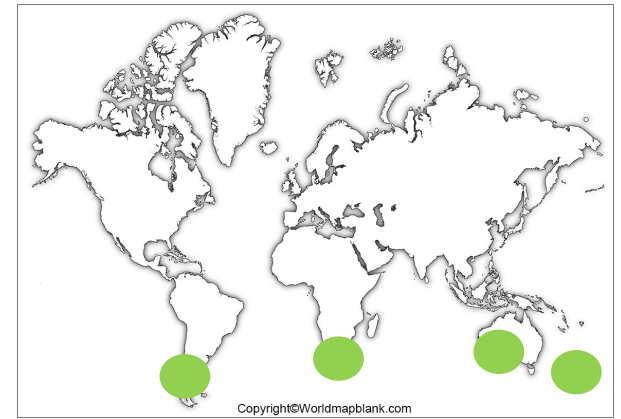
Long distance dispersal – rafting

- Also in *Poecilomigas abrahami*

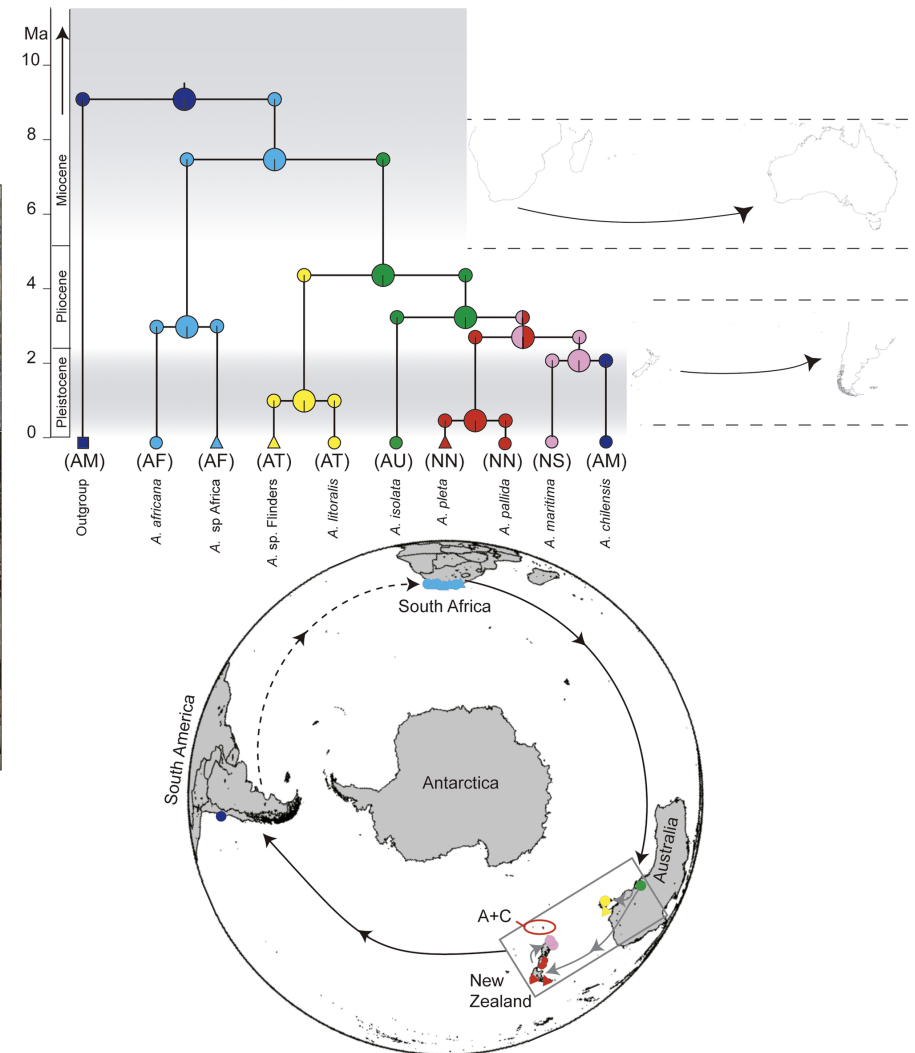
Amaurobioides continental vicariance?

Anyphaenidae

Around the world in 8 million years - rafting



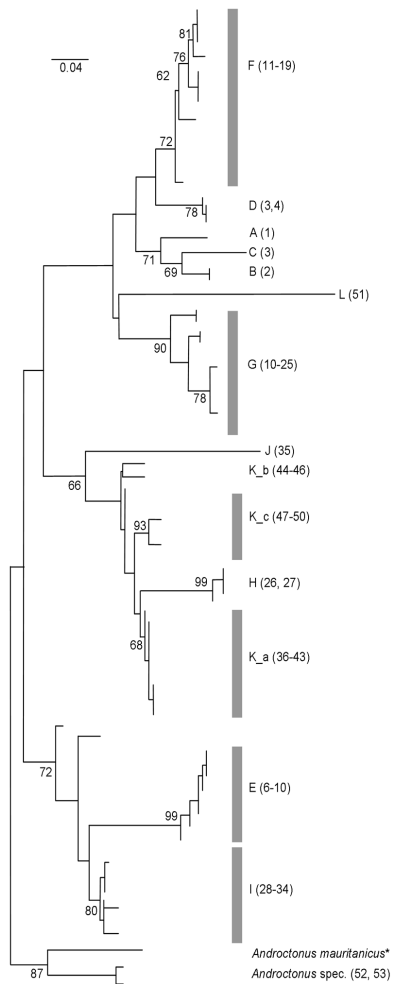
Ceccarelli et al 2016



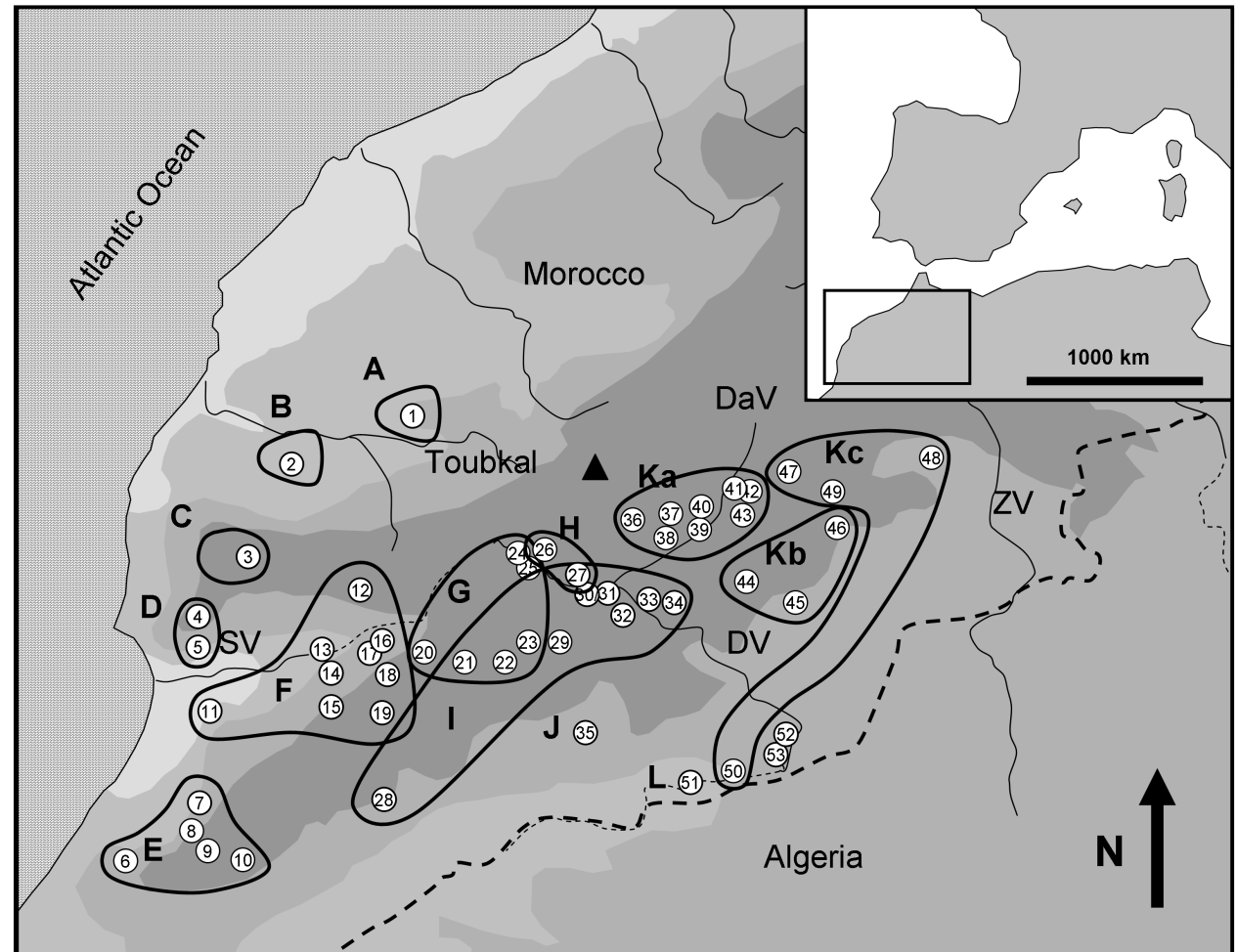
Buthus scorpions in Atlas Mountains

Mountains – *in situ* radiation, microallopatry

Main clades overlap, subclades parapatric



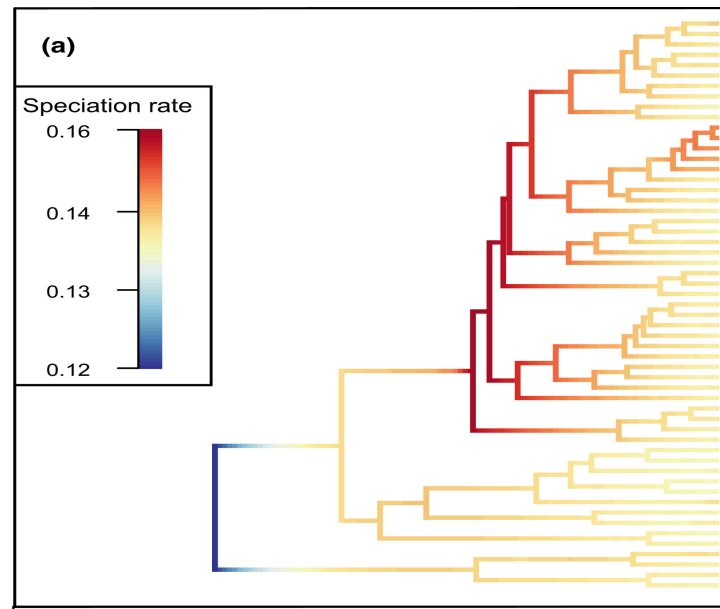
Habel et al 2012



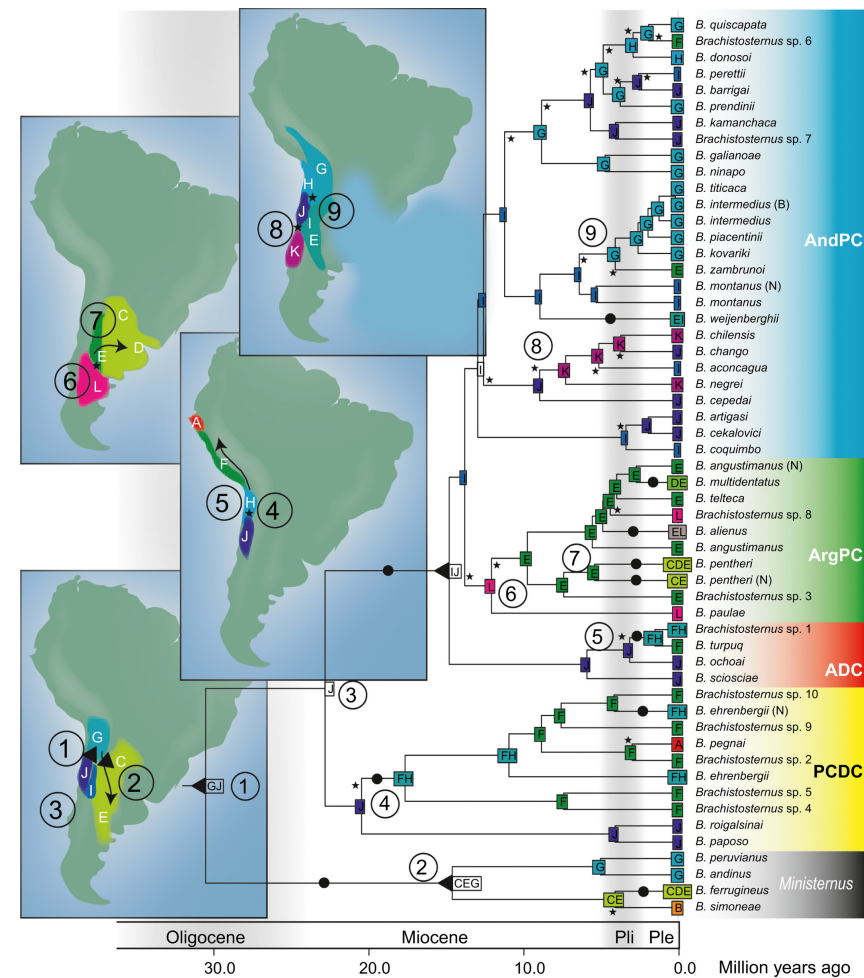
Brachiolesternus scorpions in the Andes

Mountains – *in situ* radiation

Coastal habitats stable – source of colonization of adaptive lineages



Ceccarelli et al 2016

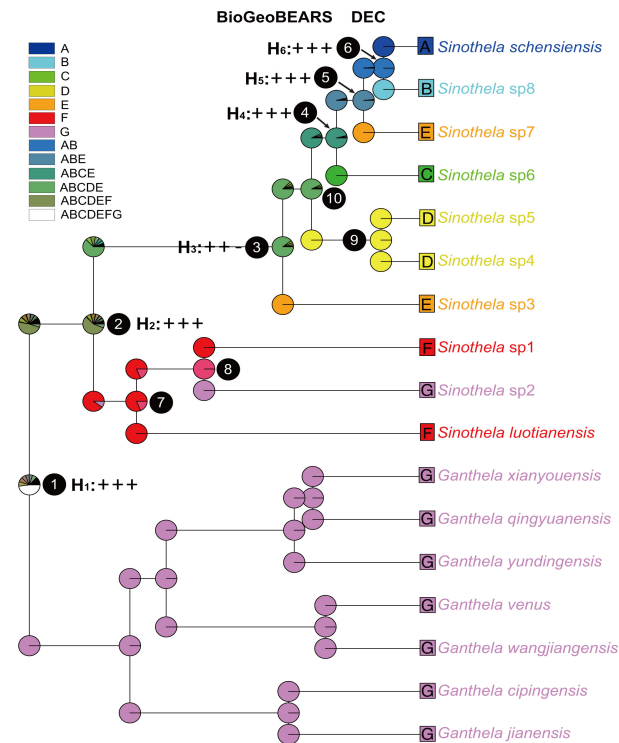
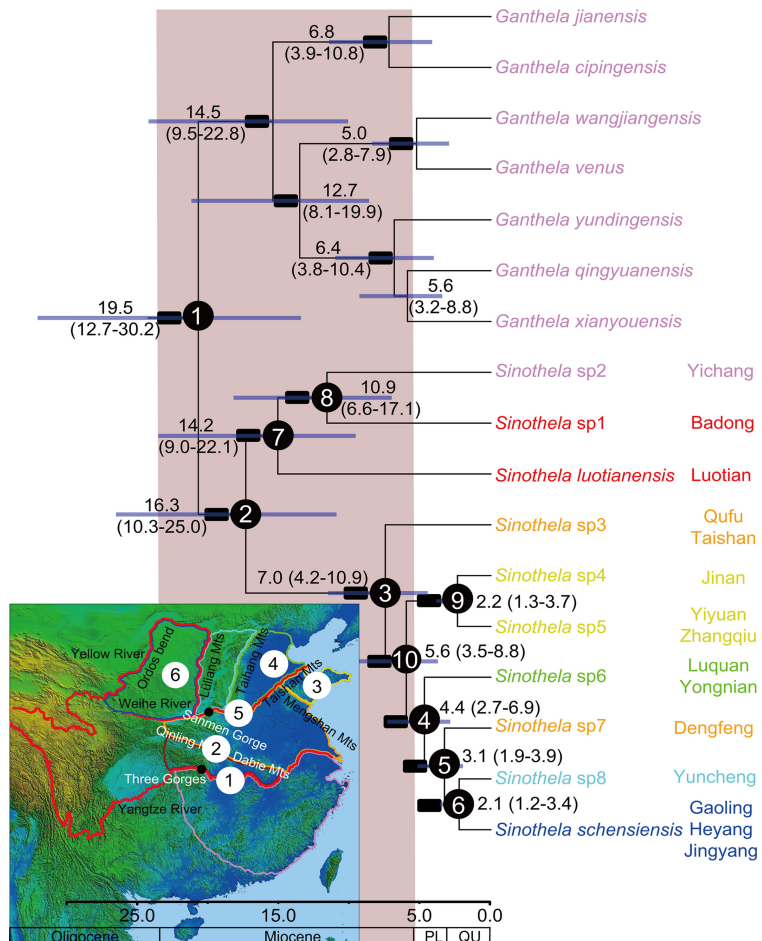


Primitively segmented spiders SE Asia

Ganthela, *Sinothela*



River formation, mountain uplift

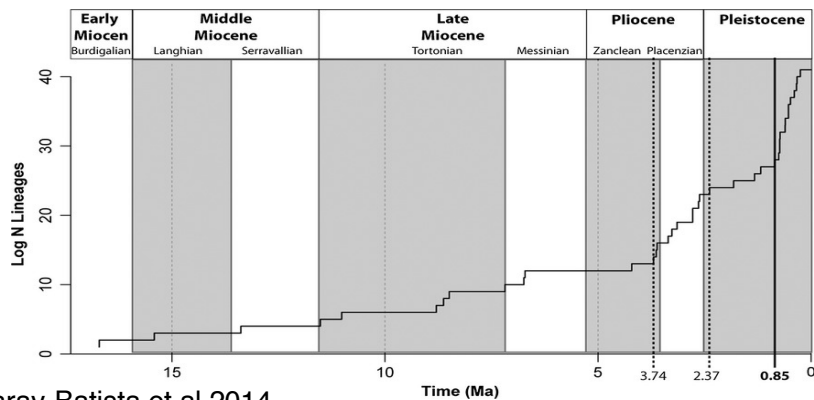
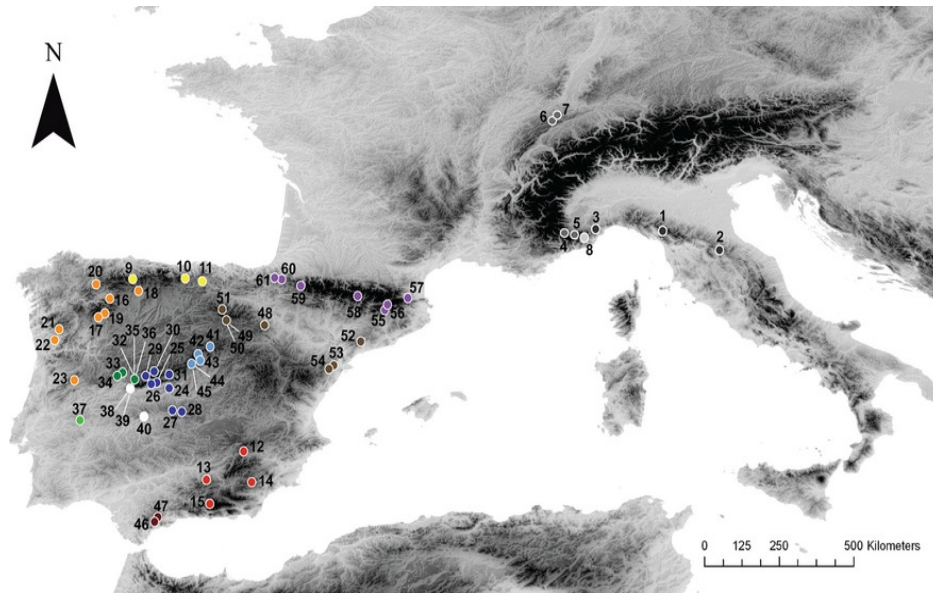


1. Yangtze River formation (23–36.5 Ma)
2. Qinling–Dabie mountains (2.6–23 Ma)
3. Taishan Mts uplift - unsupported
4. Taihang Mts (3.6–5.3 Ma)
5. Yellow River formation (1.8–3.6 Ma)
6. Ordos bend coincide with its origin (1.6 Ma)

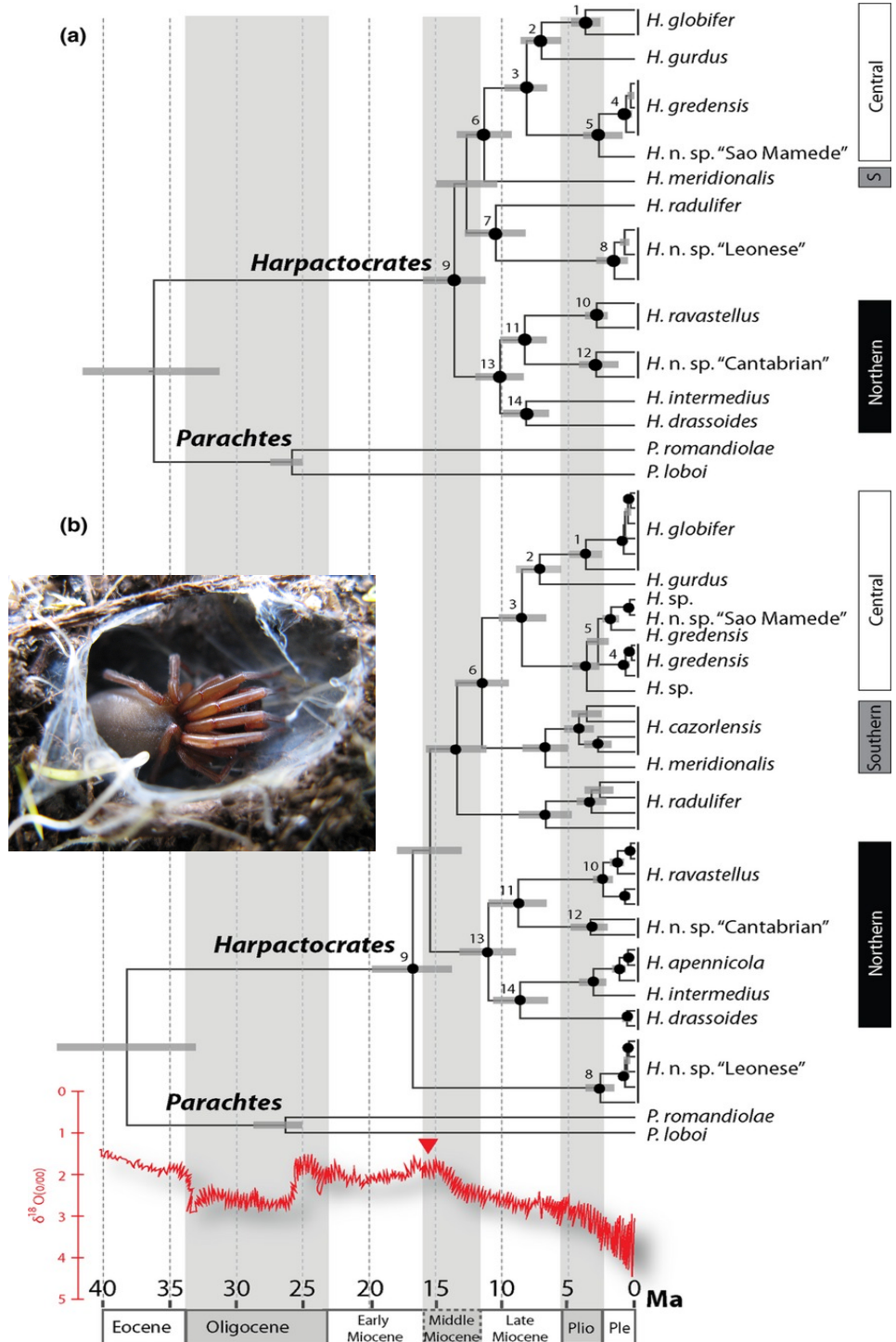
Harpactocrates Climate: Glaciations

Dysderidae

Ground-dwelling, sedentary
Western Mediterranean

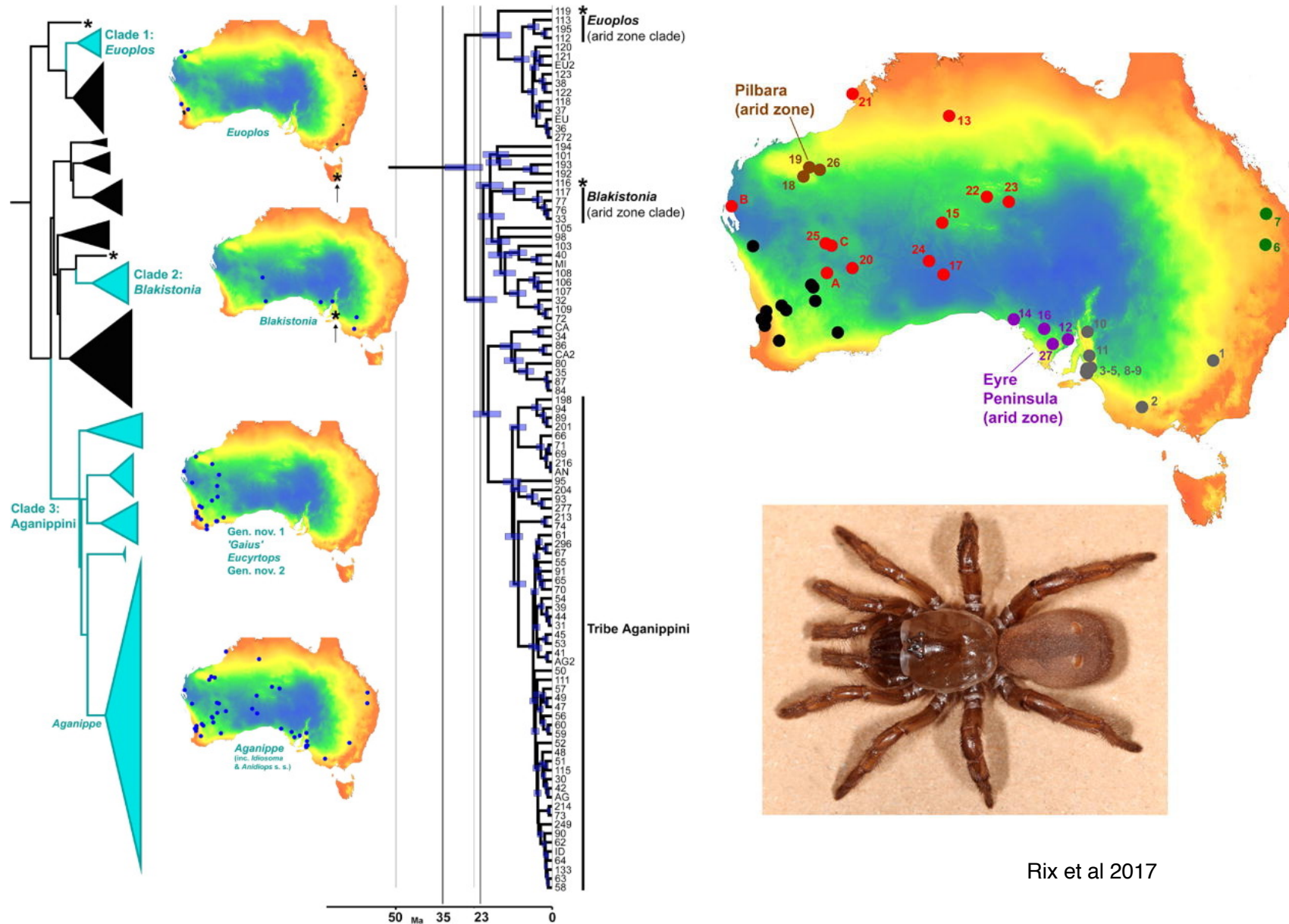


Bidegaray-Batista et al 2014



Idiopidae Climate: Aridification of Australia

Trapdoor spiders



Rix et al 2017

Summary

Amalgamation of Gondwana 520 – 510 Ma
- southern hemisphere

Laurentia, Baltica, Siberia – in the north
- formed Laurasia in Paleozoic, ~ 300 Ma

Late Paleozoic – Pangea formation
- lasted ~ 100 Myr

Pangea breakup
- Central Atlantic ridge ~ 200 Ma

Lower Jurassic ~ 180 Ma E/W Gondwana breakup

Upper Jurassic ~ 160 Ma India-Madagascar/Antarctica
~ 160 Ma Antarctica/Australia

Lower Cretaceous ~ 140 – 130 Ma S America/Africa

Upper Cretaceous ~ 80 - 90 Ma Madagascar/ India
~ 50 Ma India collided with Asia

Laurasia breakup ~55 Ma
N Am land bridge ~ 25 ma

Hercynian belt breakup 30 - 25Ma

Beatic Rif broke of Sardinina + Corsica
~20Ma

→ Sardinia + Corsica collided with Italy,

10 – 5 Ma final separation of Sardinia + It

Messinian Salinity crisis 5.93 – 5.3 Ma

5.3 Ma Opening of Strait of Gibraltar

Last glaciation: 2.58 Ma to 12 000 ya