



WHAT DRIVES BIOLOGICAL DIVERSIFICATION?

DETECTING TRAITS UNDER SPECIES SELECTION

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University of British Columbia
Macquarie University





Gould 1839

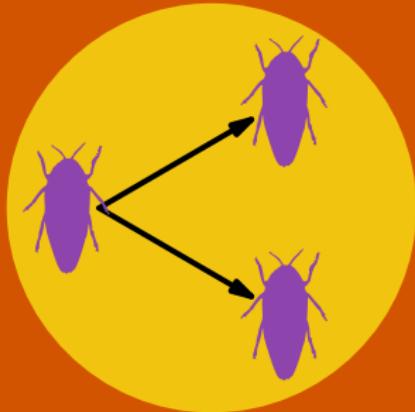


Gould 1839



SPECIES SELECTION

Variation

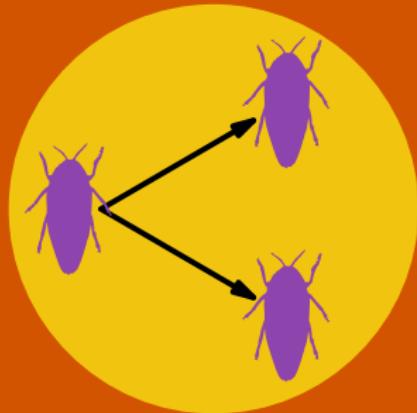


Heritability



Differential fitness

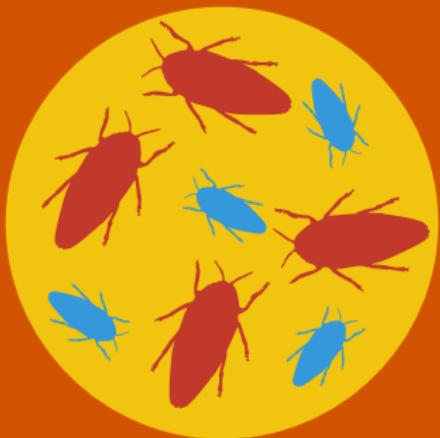
Variation



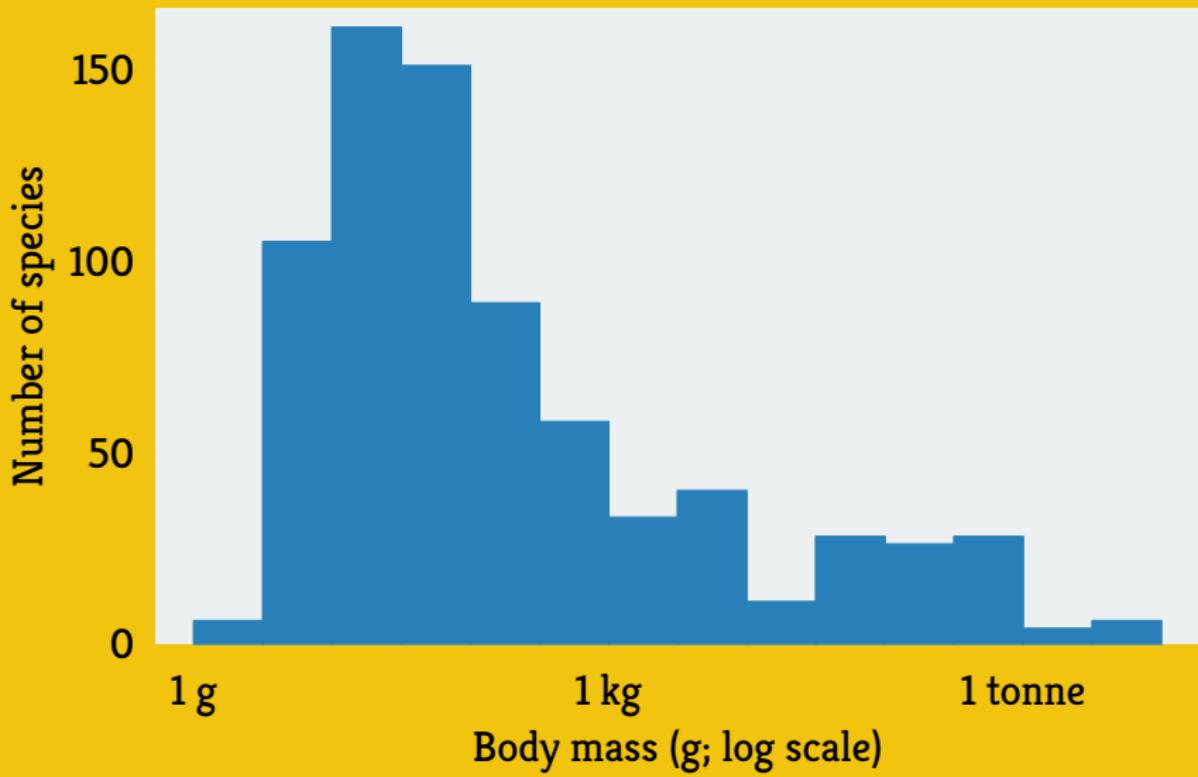
Heritability

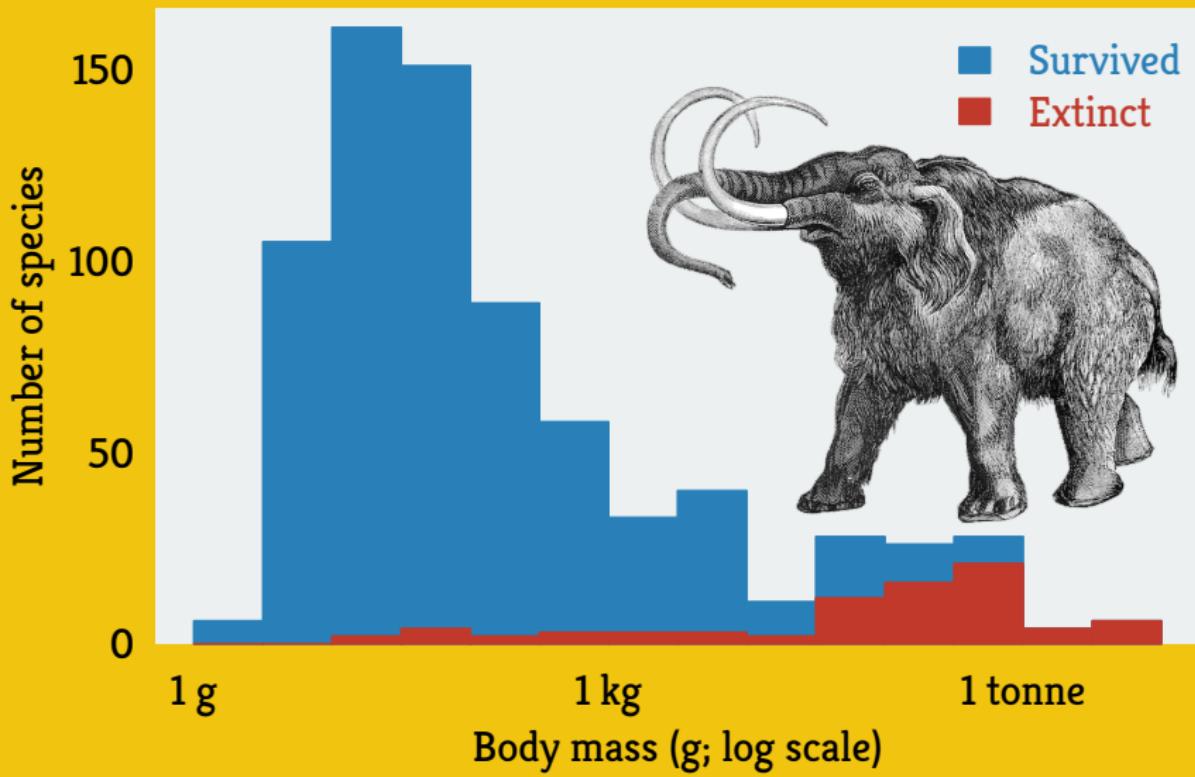


Differential fitness

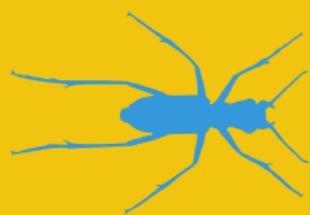
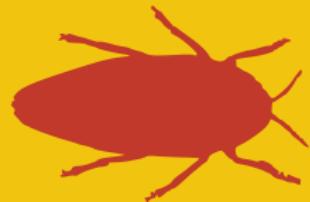


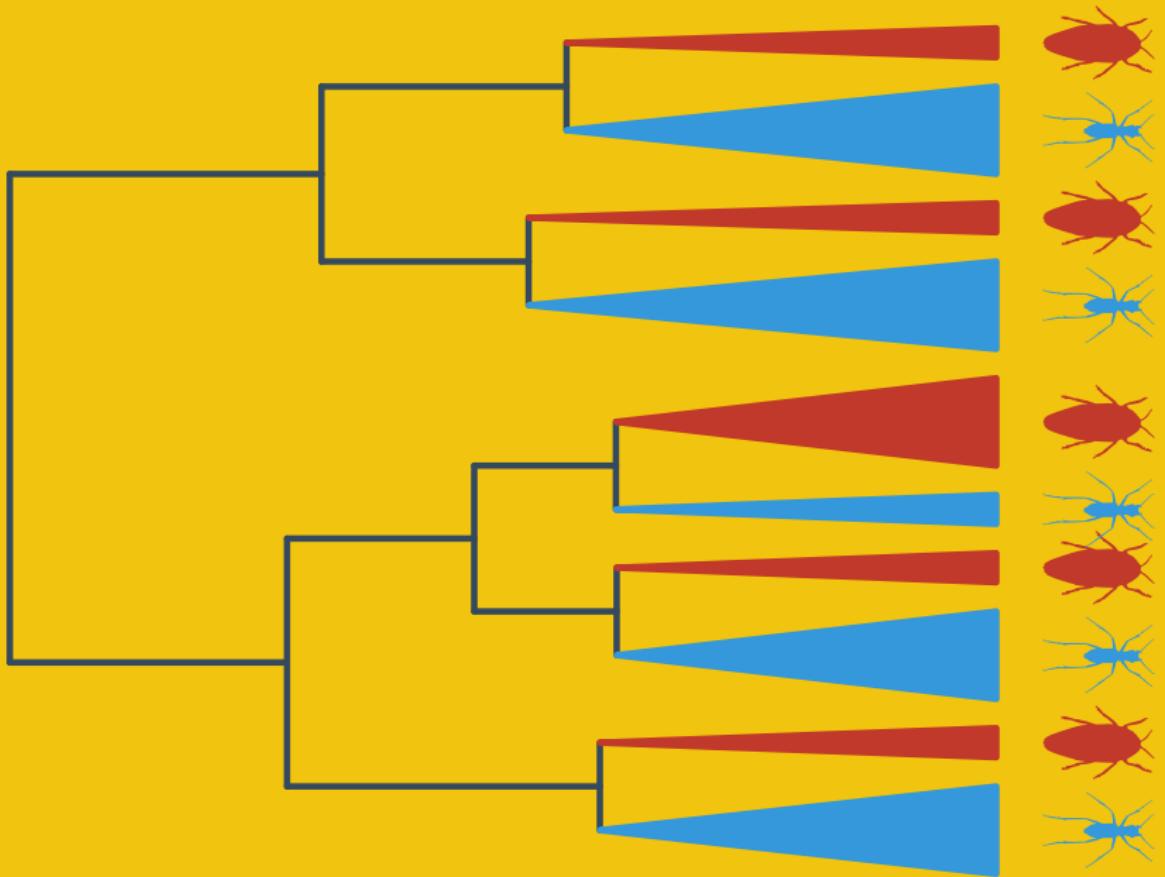
**SPECIES SELECTION
HAS NOT BEEN POPULAR**



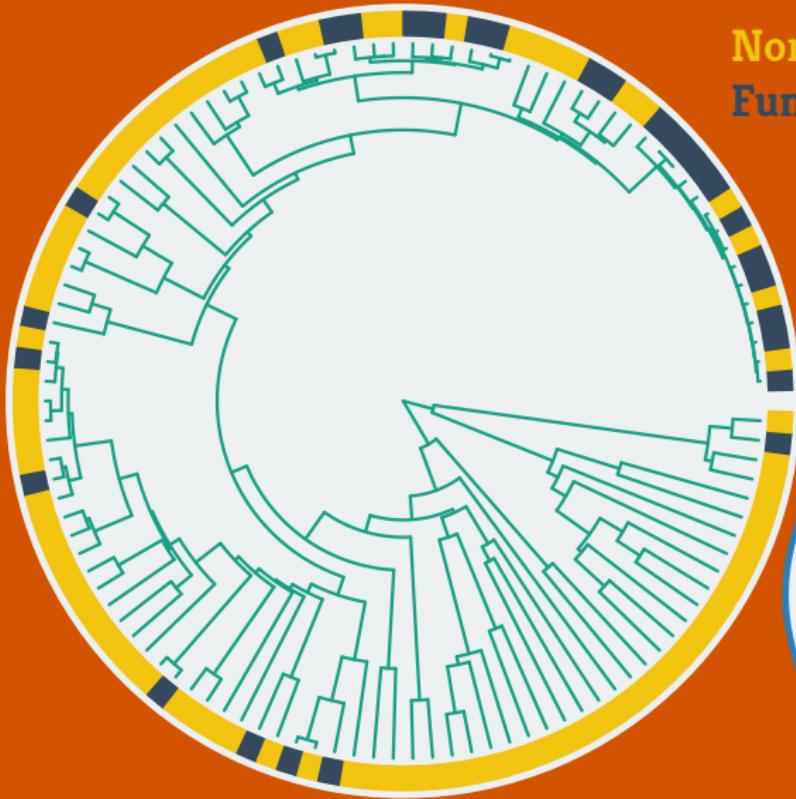


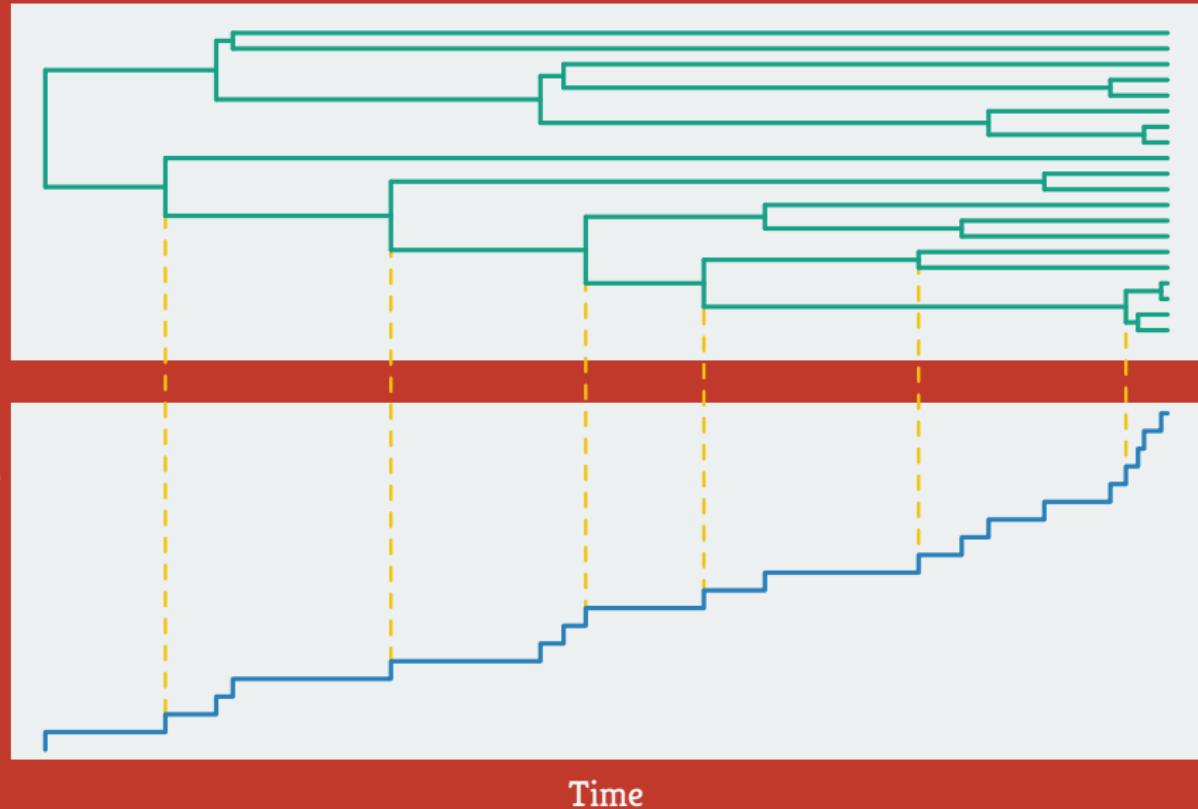
HOW TO DETECT SPECIES SELECTION?

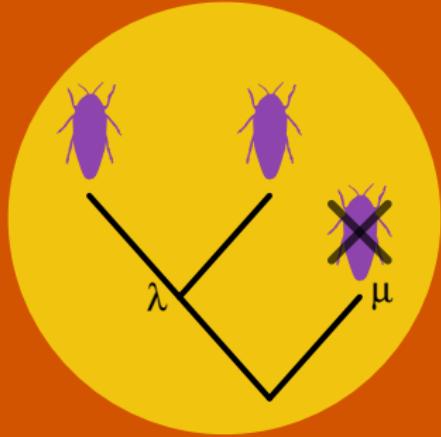




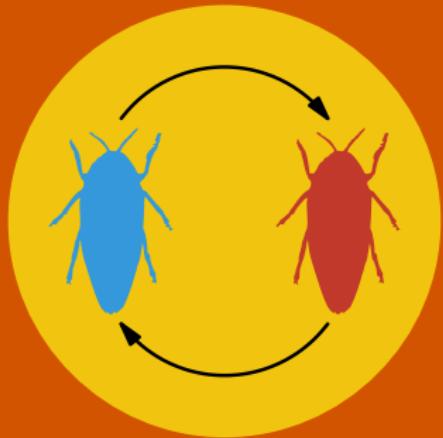
Normal
Functionally asexual



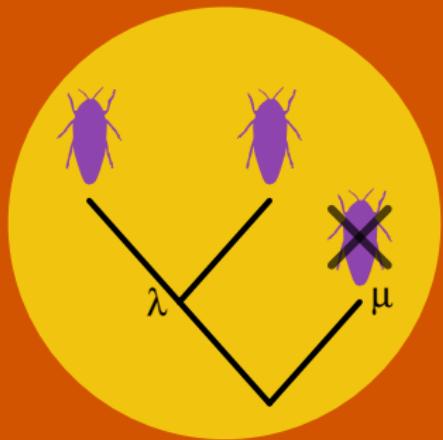




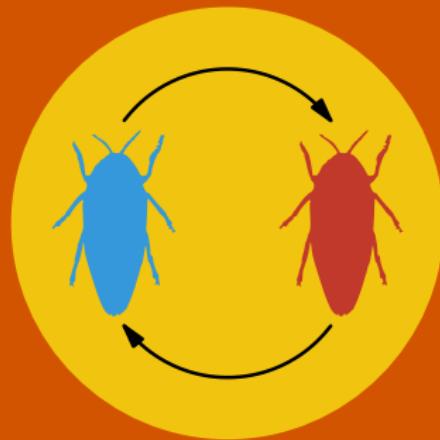
Speciation / extinction: birth death model



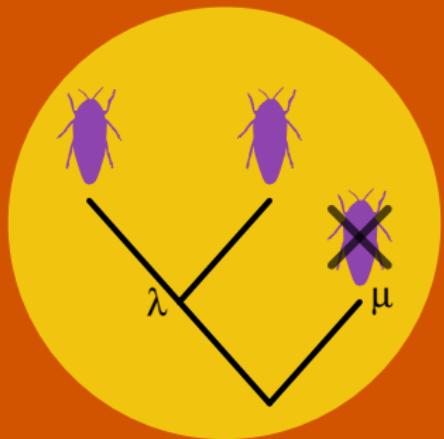
Trait evolution: Markov model



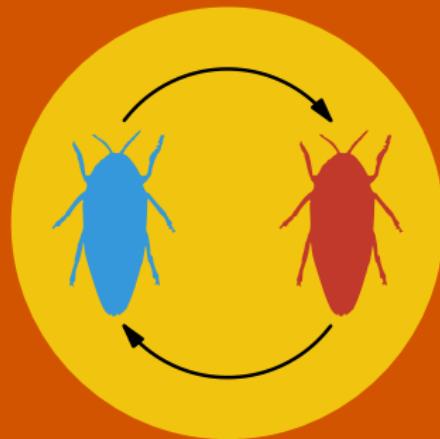
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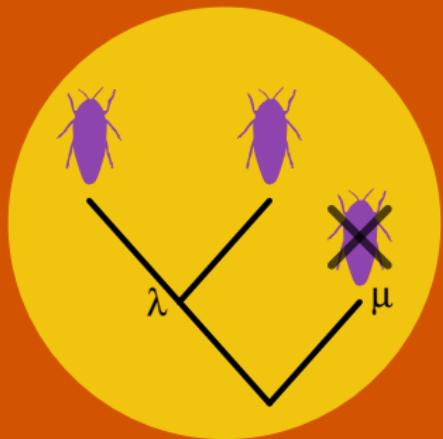
BiSSE: Binary State Speciation & Extinction



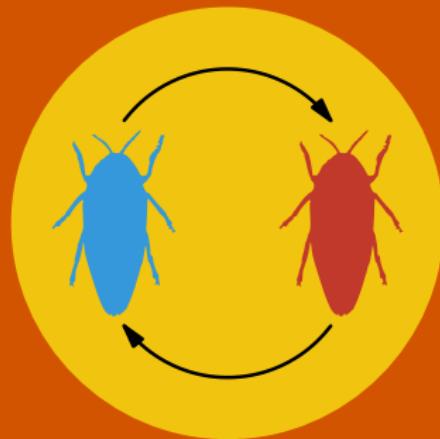
+



$$\Pr \left[\text{[Diagram of insects and phylogenetic tree]} \right]$$



+



λ

μ

p

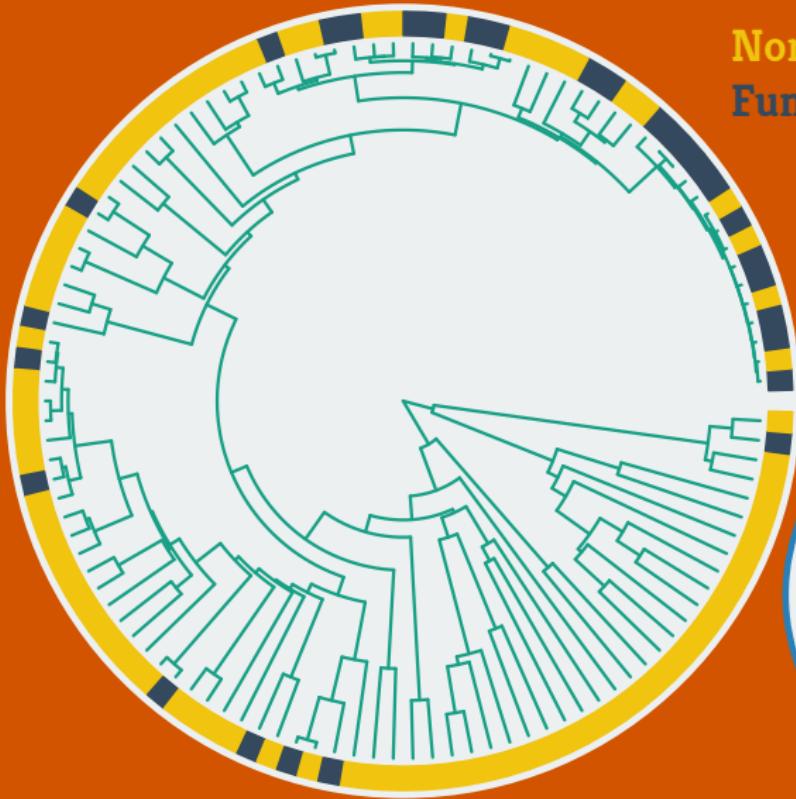
λ

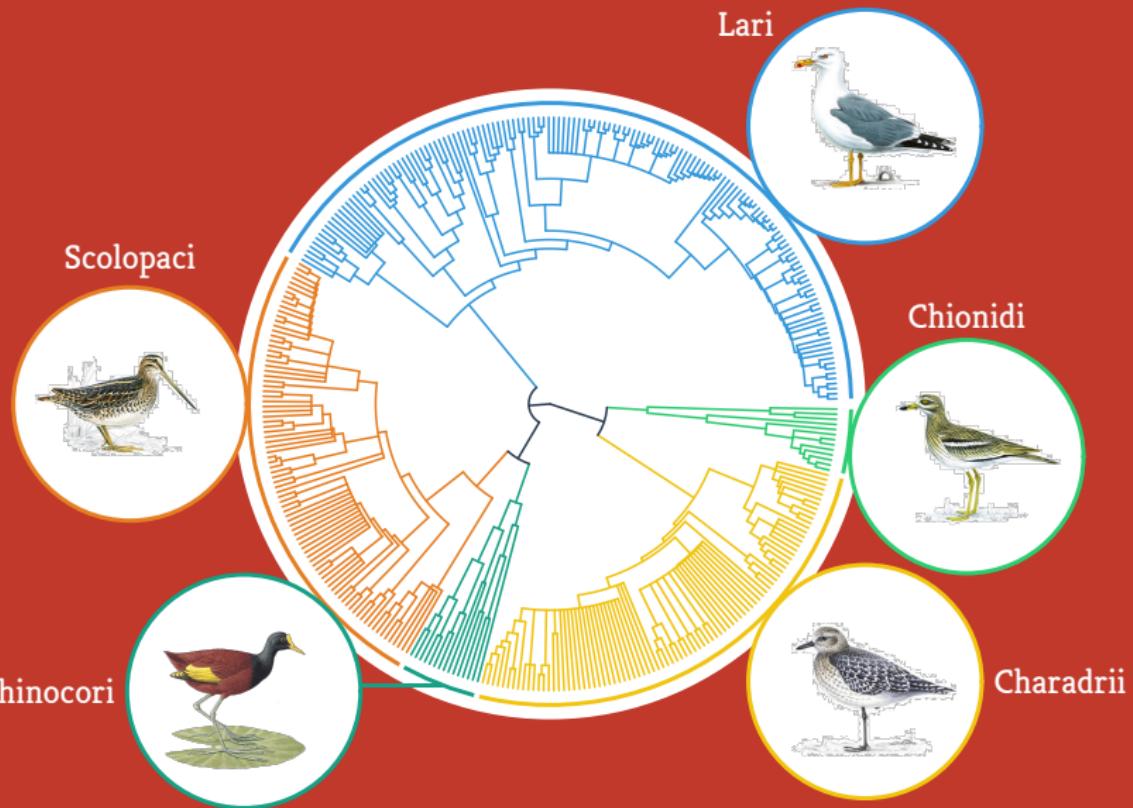
μ

p

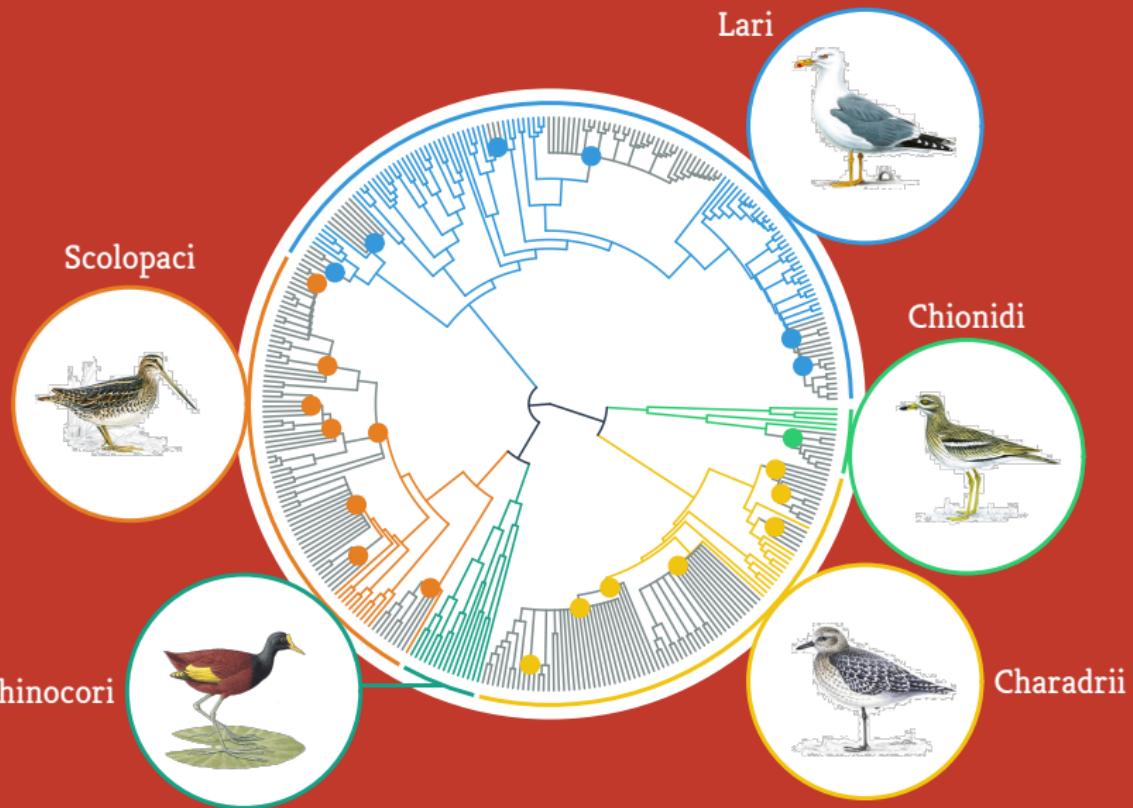
COPING WITH PHYLOGENETIC INCOMPLETENESS

Normal
Functionally asexual

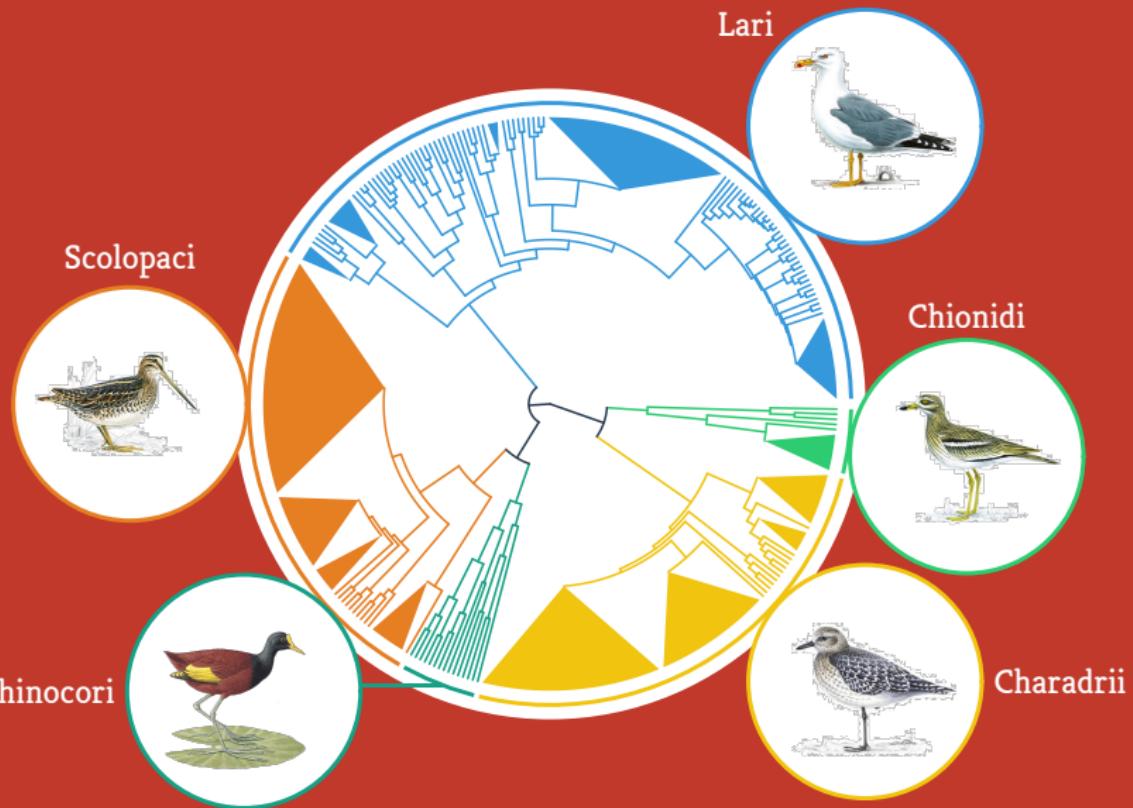




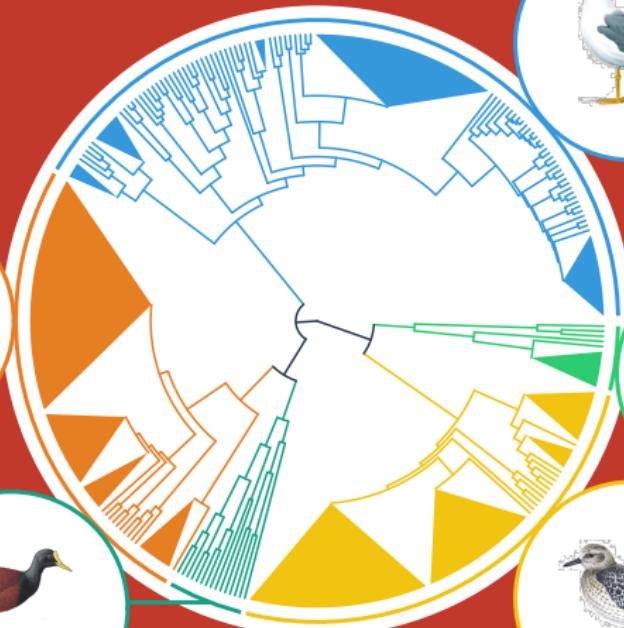
Tree: Thomas, Wills & Székely 2004



Tree: Thomas, Wills & Székely 2004



Tree: Thomas, Wills & Székely 2004



Scolopaci



Lari



Chionidi



Thinocori



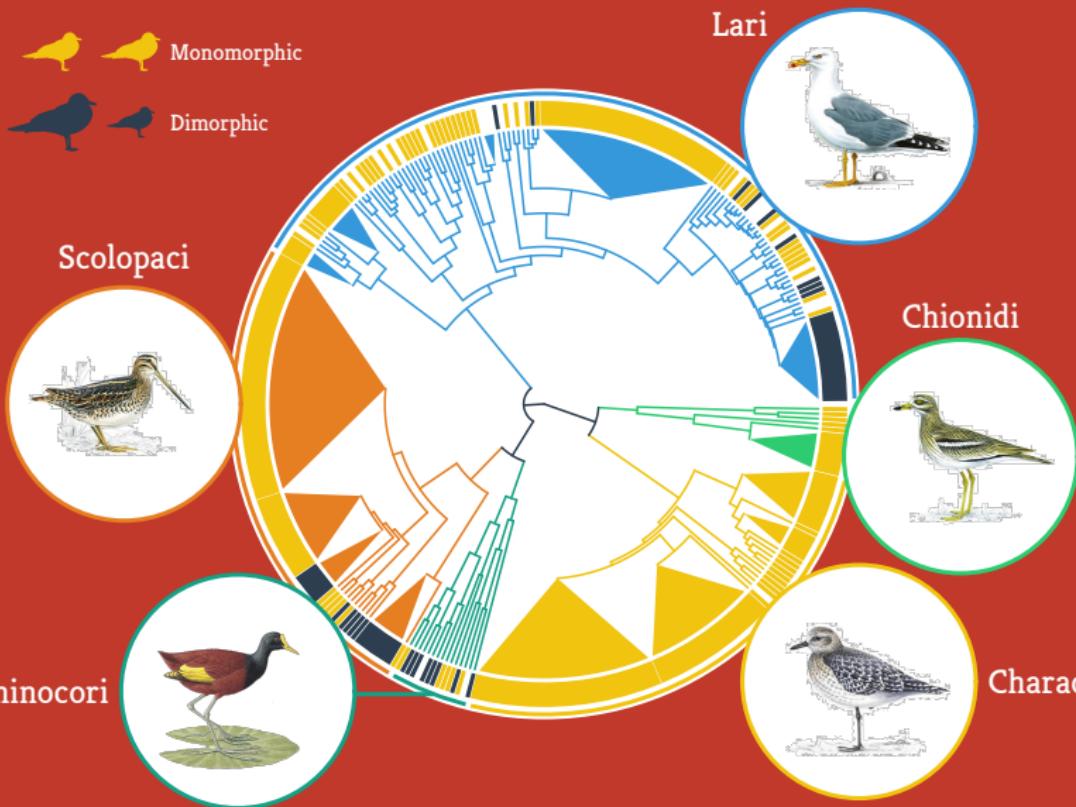
Charadrii





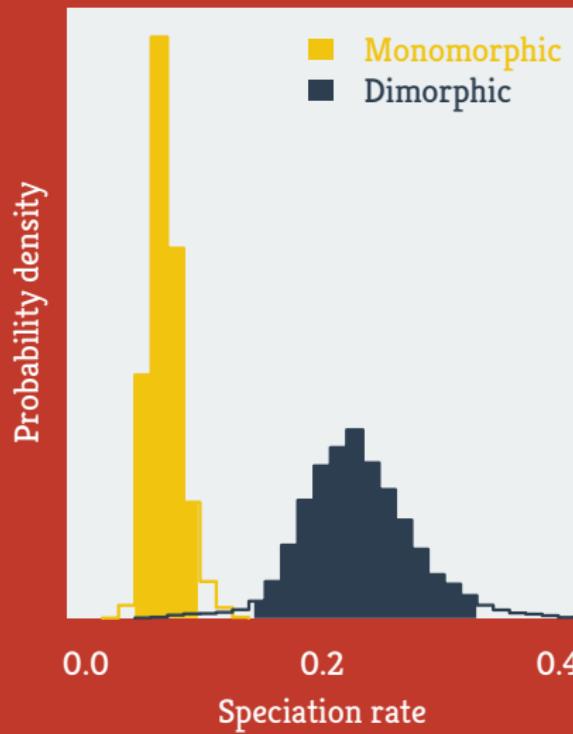
Photo: Thor Veen

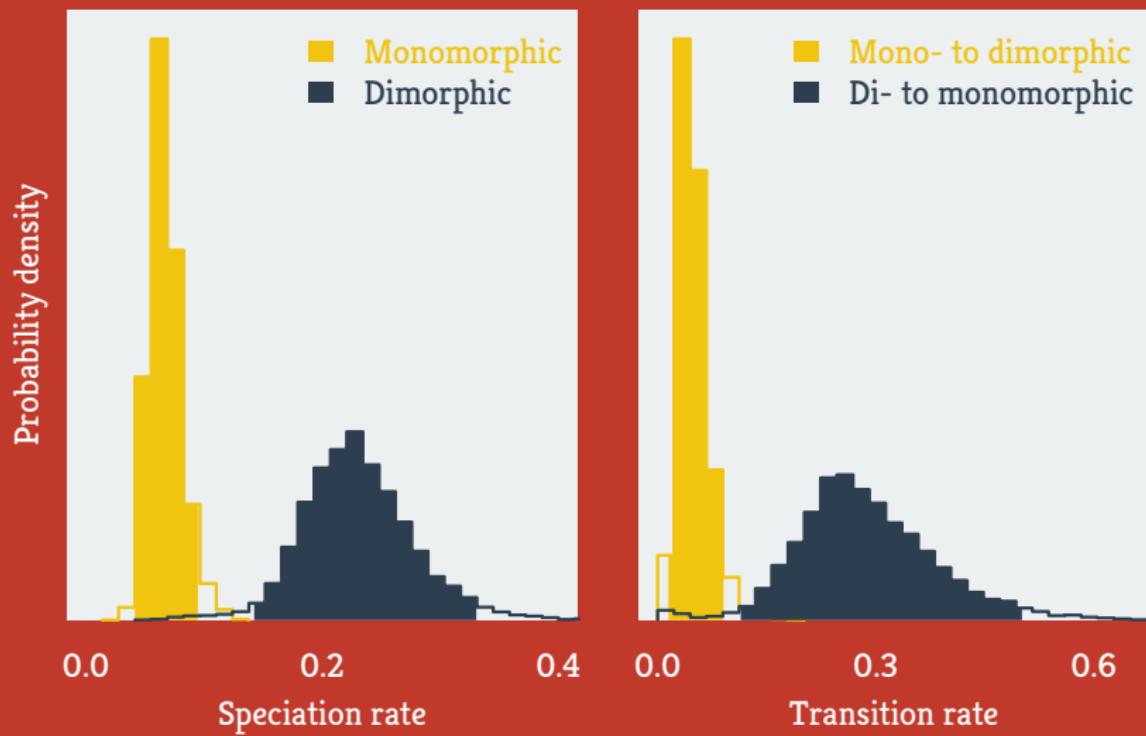
Is
sexual size dimorphism
under species selection?



Tree: Thomas, Wills & Székely 2004

Data: Lislevand, Figuerola & Székely 2007

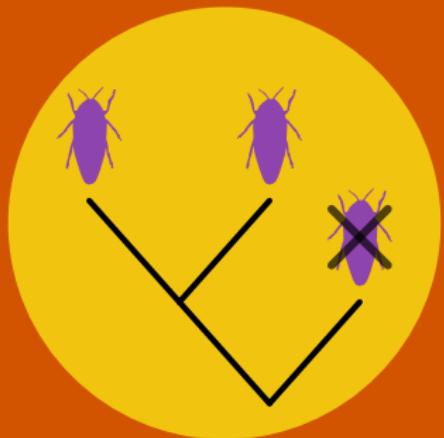




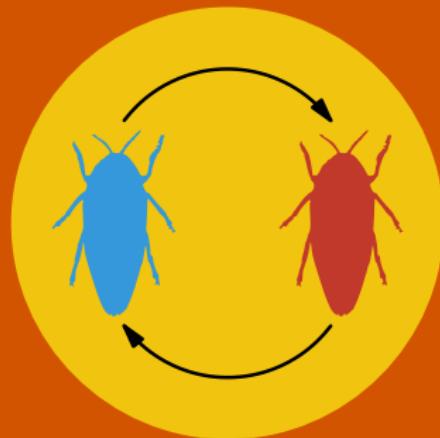
Is
sexual size dimorphism
under species selection?

Perhaps briefly

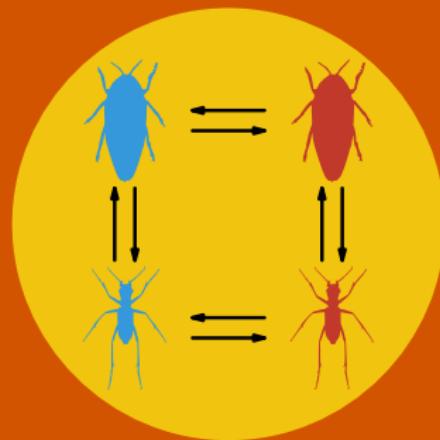
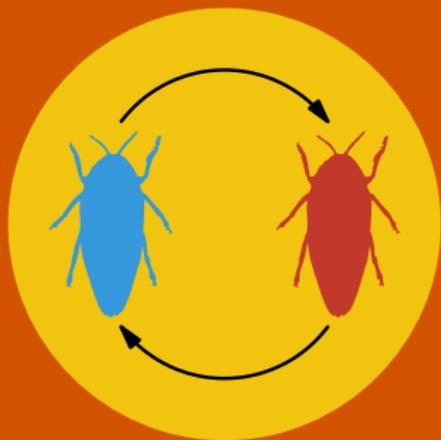
DEALING WITH MULTIPLE TRAITS

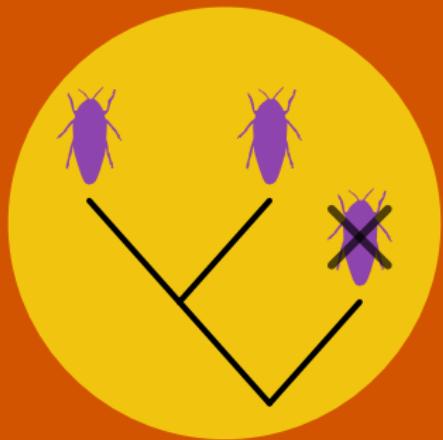


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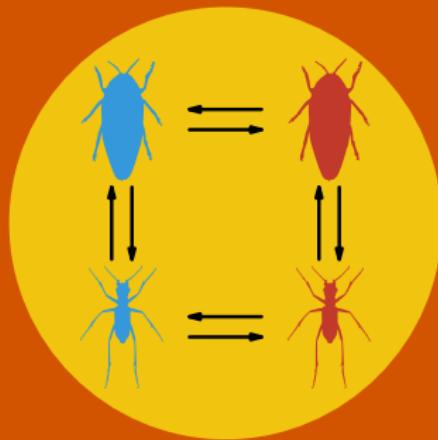


$$\Pr \left[\text{[Diagram of a phylogenetic tree with 10 nodes and 9 edges, where the first node is blue and the remaining 9 are red]} \right]$$

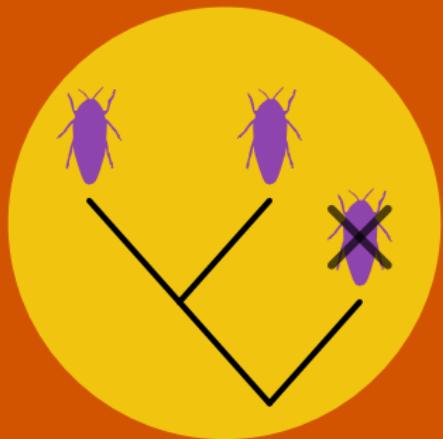




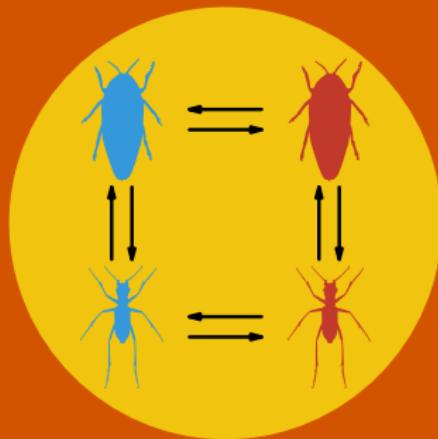
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$$\Pr \left[\begin{array}{c} \text{insects} \\ \text{lineage tree} \end{array} \right]$$

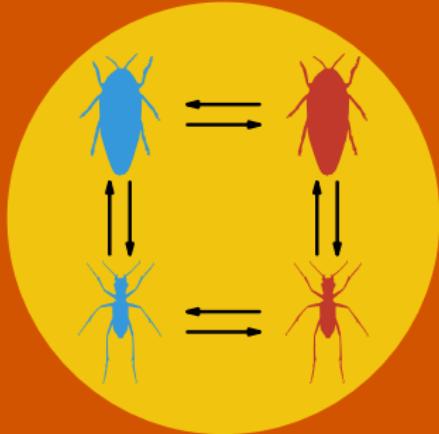
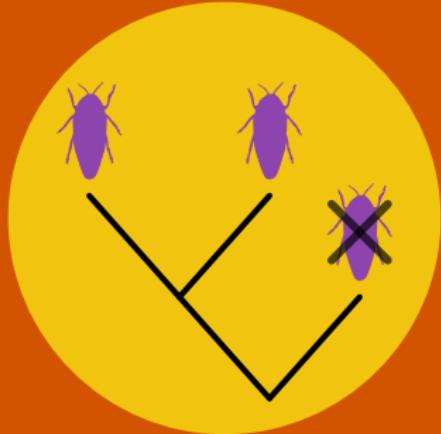


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MuSSE: Multi State Speciation & Extinction





Speciation rate:

$$\lambda = \lambda_0 + \lambda_{\bullet} + \lambda_{\star} + \lambda_{\bullet} \times \lambda_{\star}$$

Monogamous:  Solitary: 

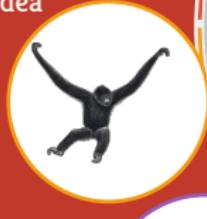


Cercopithecoidea

Lorisiformes



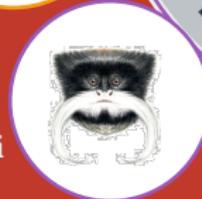
Hominoidea



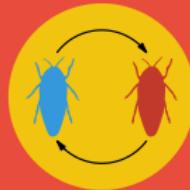
Lemuriformes



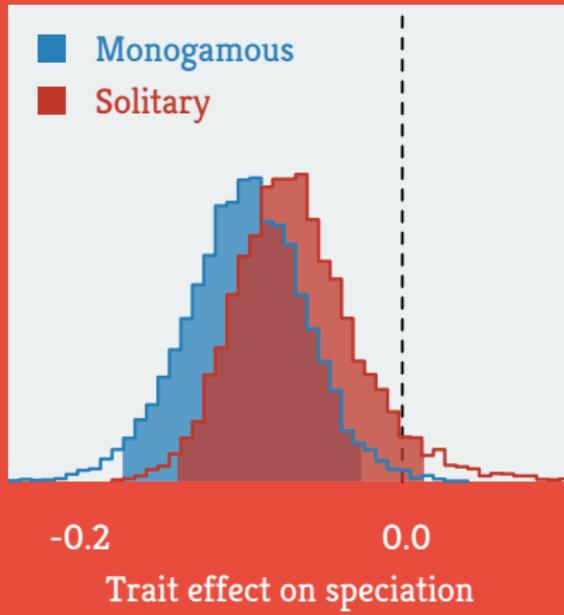
Platyrrhini

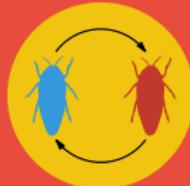


Tarsiformes

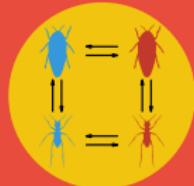
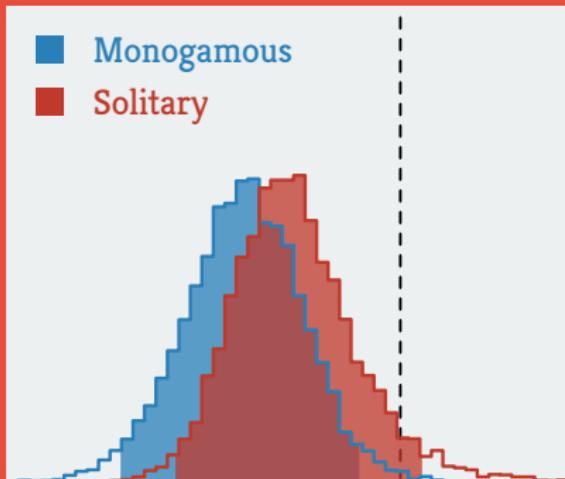


BiSSE (independent)

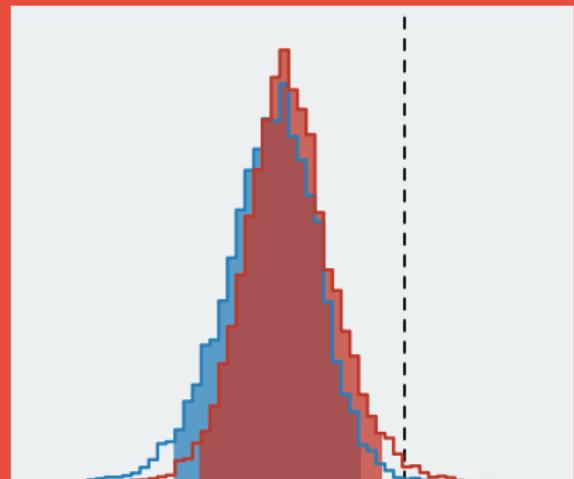




BiSSE (independent)



MuSSE (simultaneous)

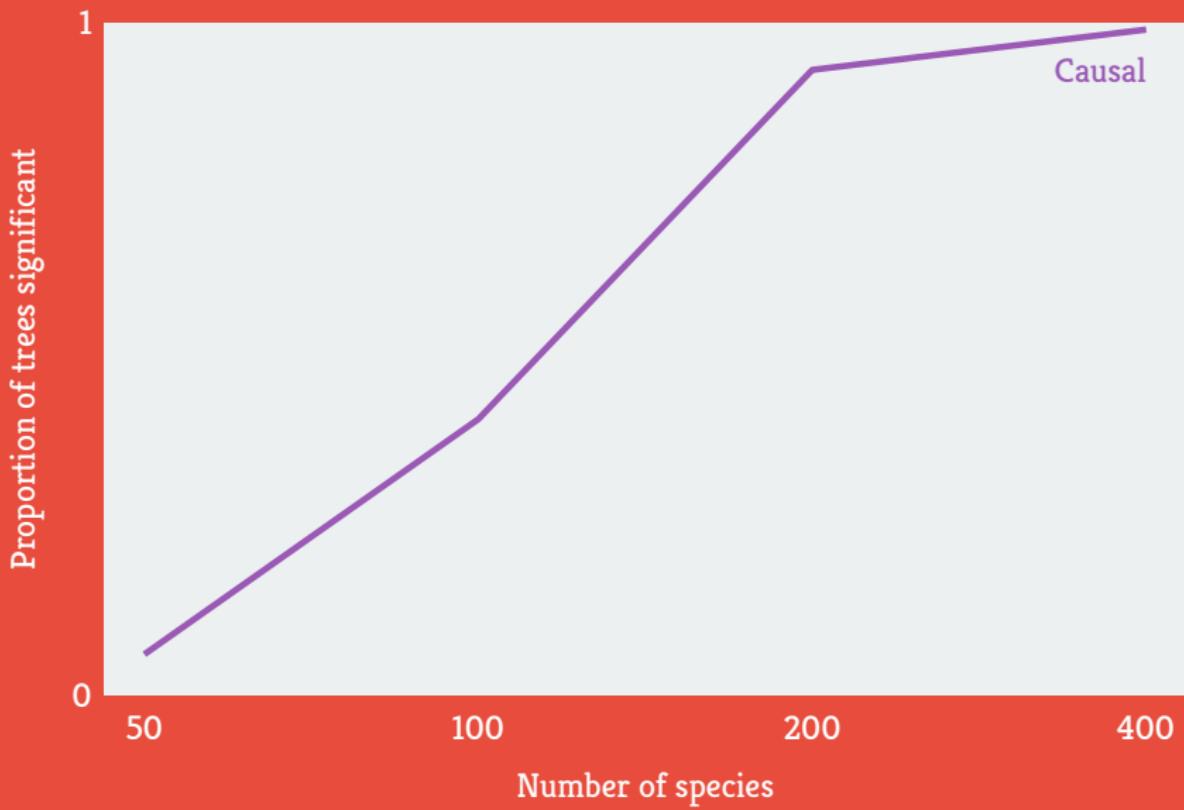


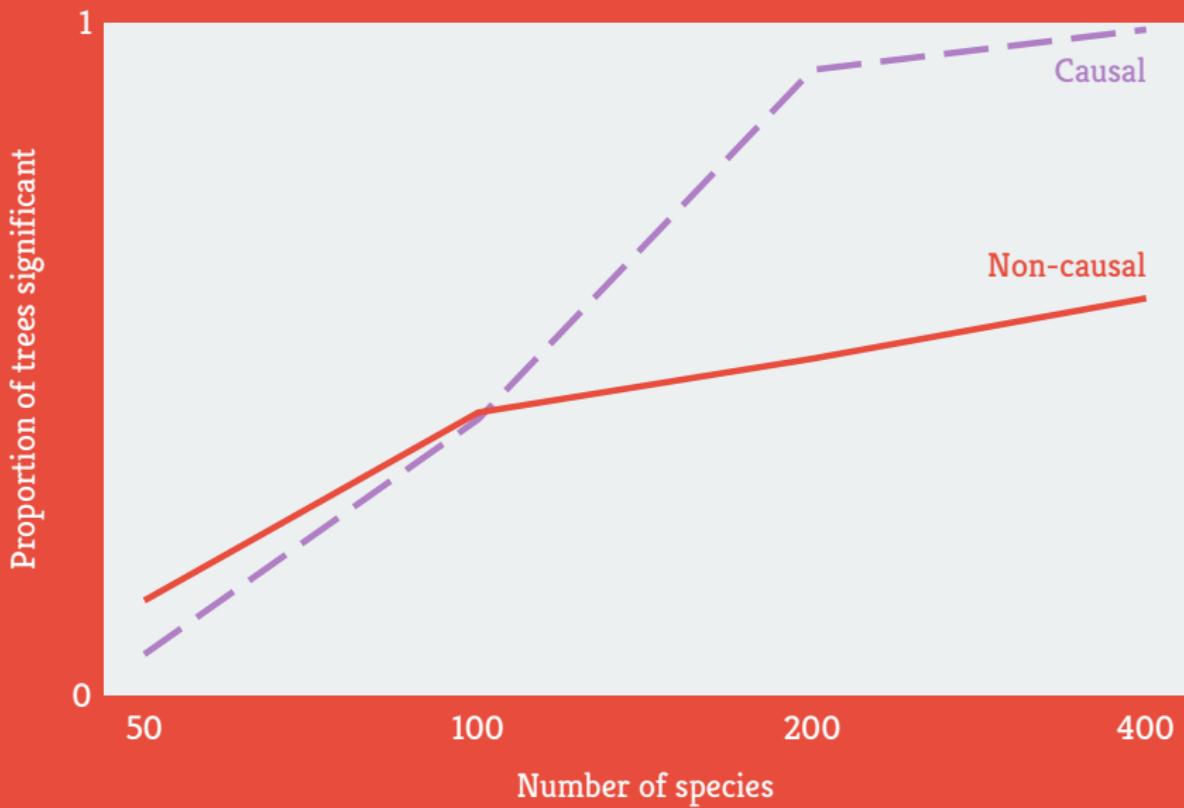
-0.2

0.0

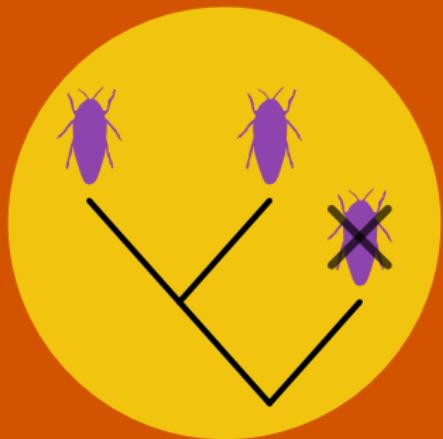
Trait effect on speciation

0.0

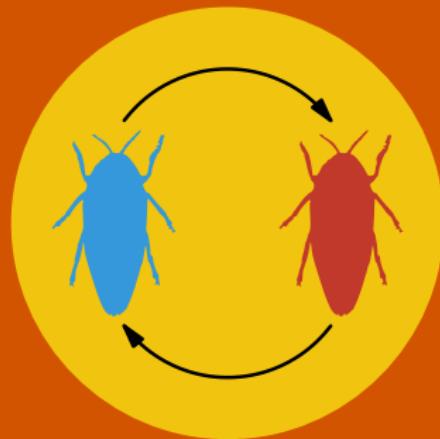




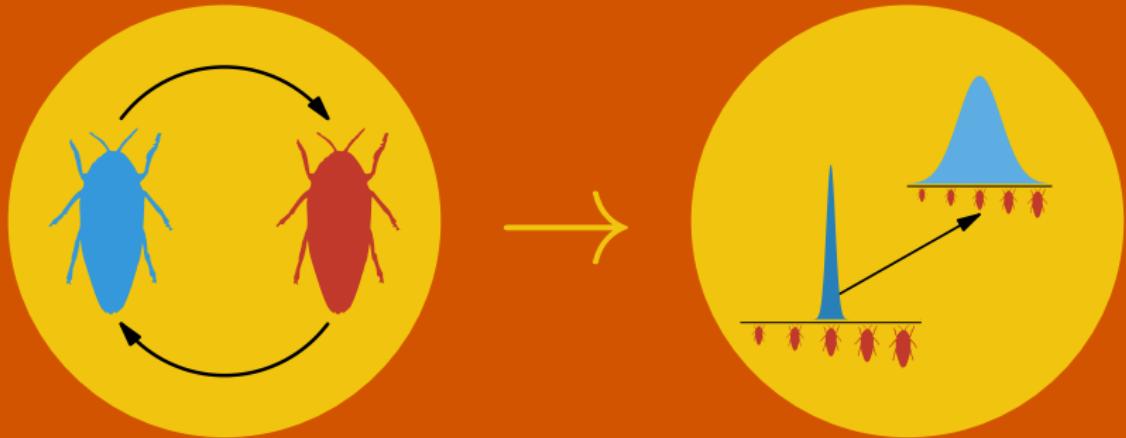
QUANTITATIVE TRAITS



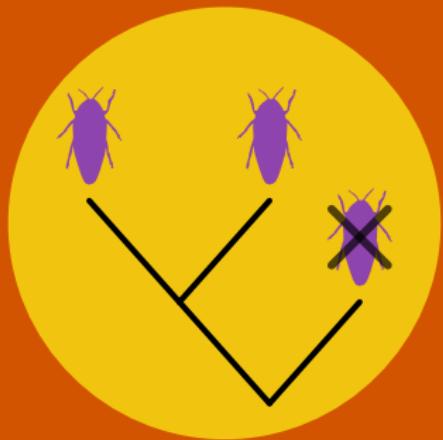
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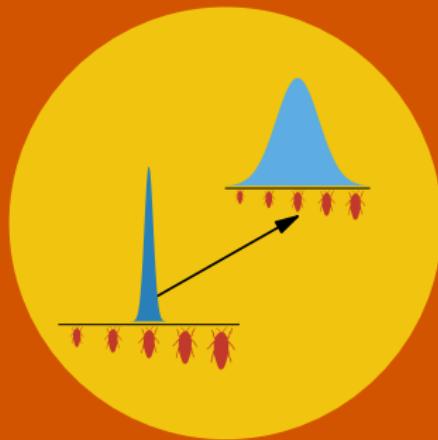
$$\Pr \left[\text{[Diagram of a phylogenetic tree with 10 nodes and 9 edges, where the first node is blue and the last 9 are red]} \right]$$



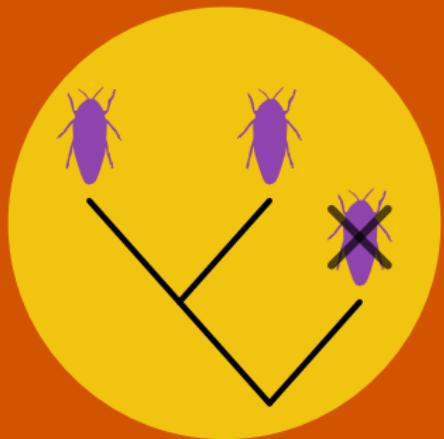
Trait evolution: Brownian Motion



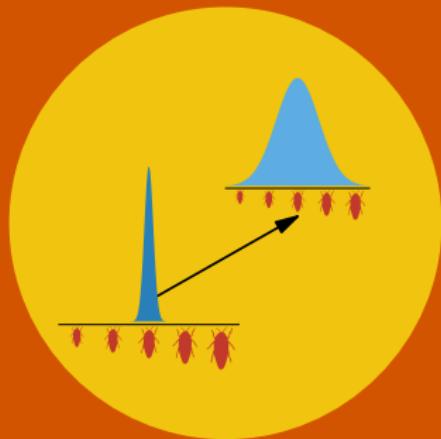
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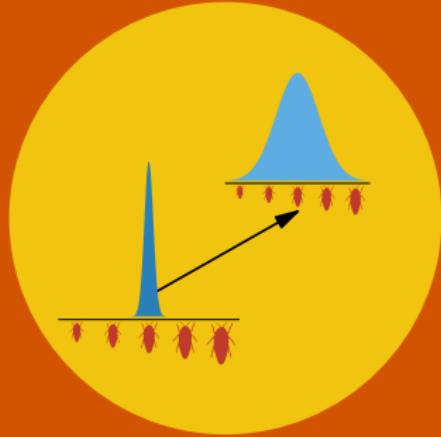
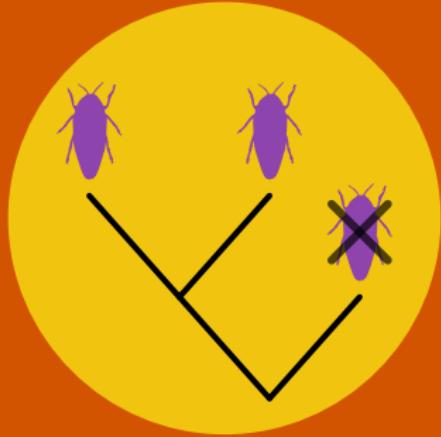
QuaSSE: Quantitative State Speciation & Extinction



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$$\Pr \left[\text{red insects} \mid \text{blue curve} \right]$$



speciation rate = function []

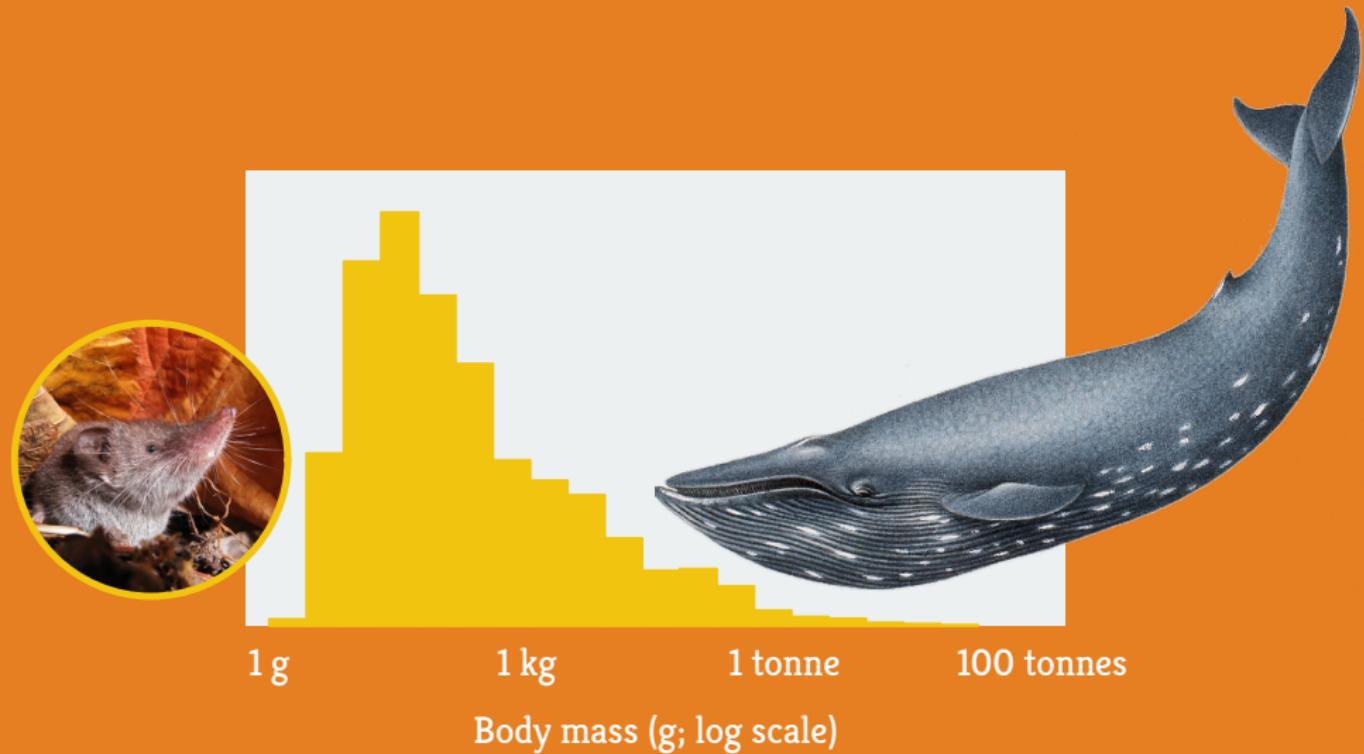
Why are there
so many small
mammal species?

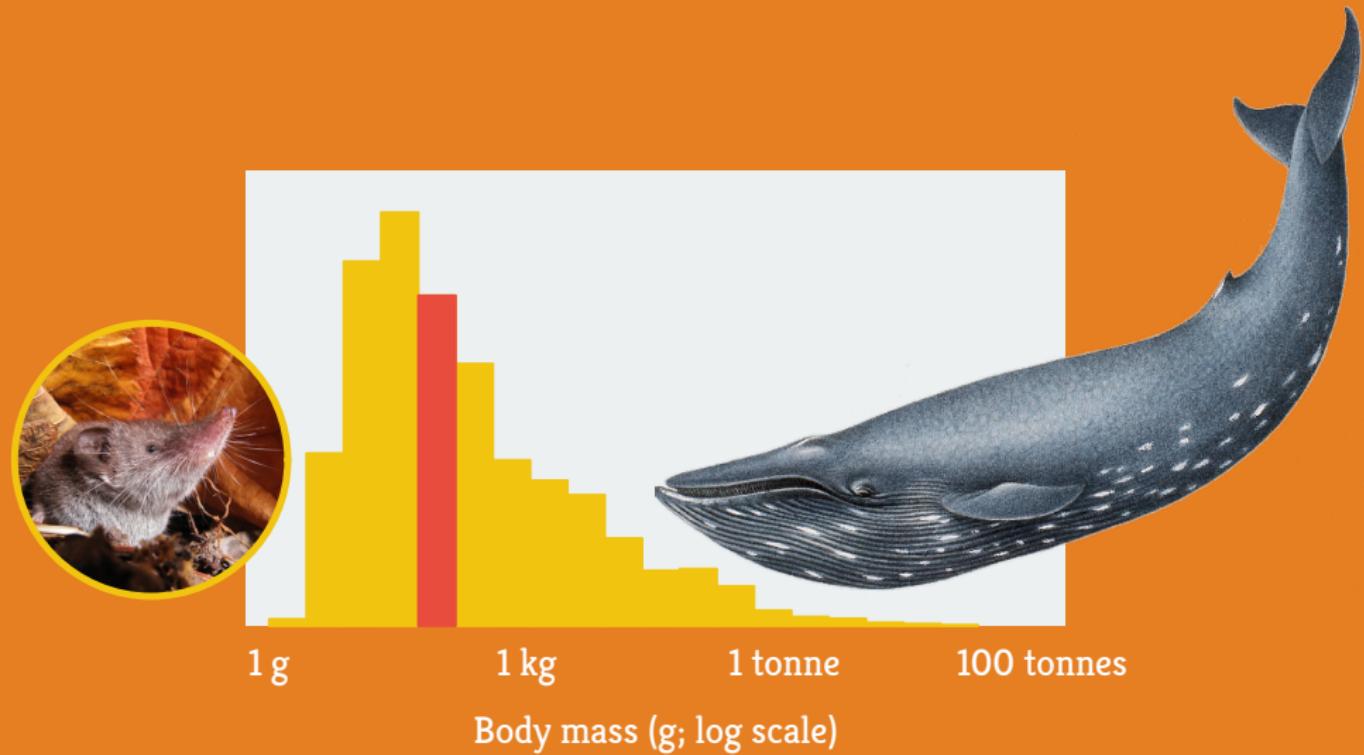


1.8g

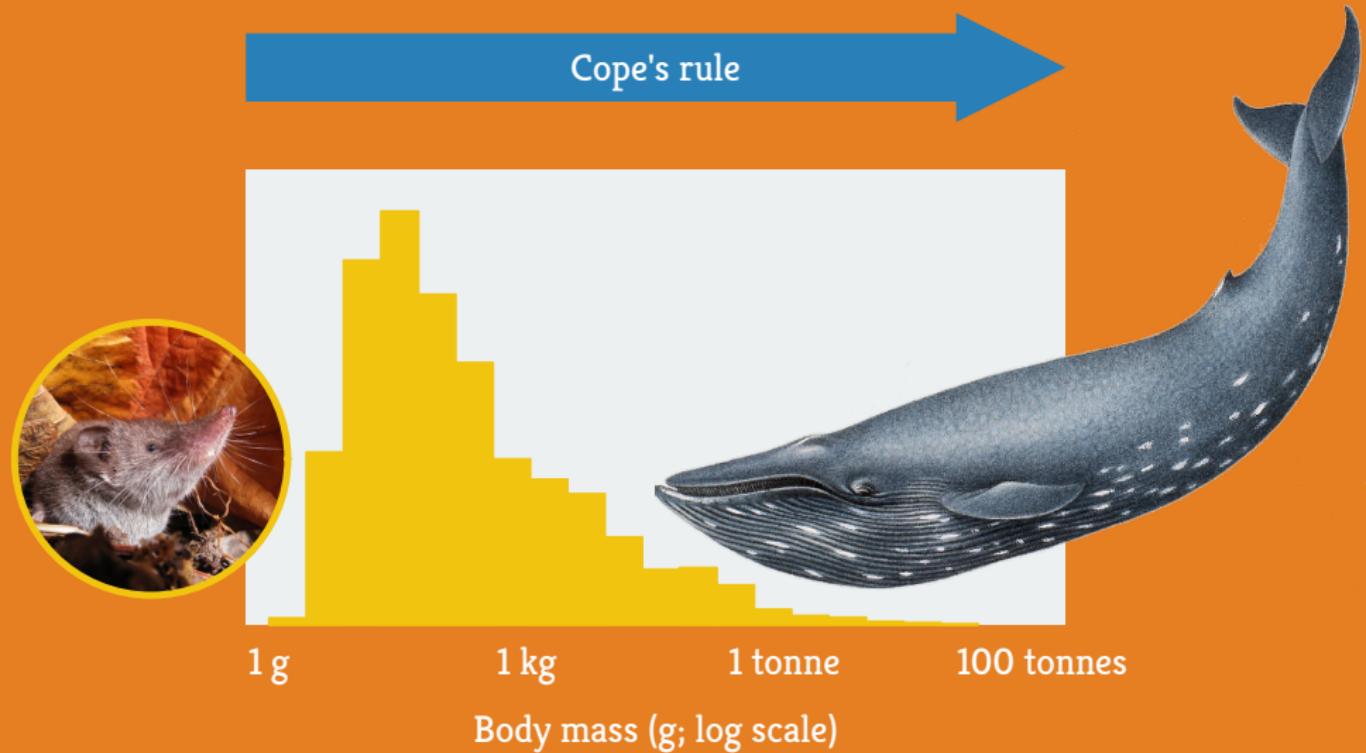


> 170 tonnes





Cope's rule



Cope's rule



Species selection



Tree: Bininda-Emonds et al. 2007

Data: Jones et al. 2009

Speciation rate

0.3

0.0

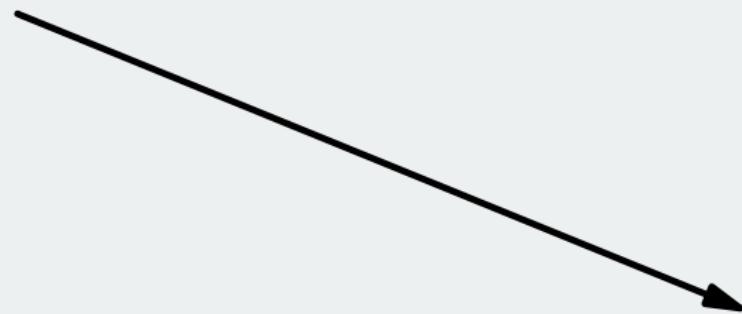
1 g

1 kg

1 tonne

100 tonnes

Body mass (g; log scale)



Speciation rate

0.3

0.0

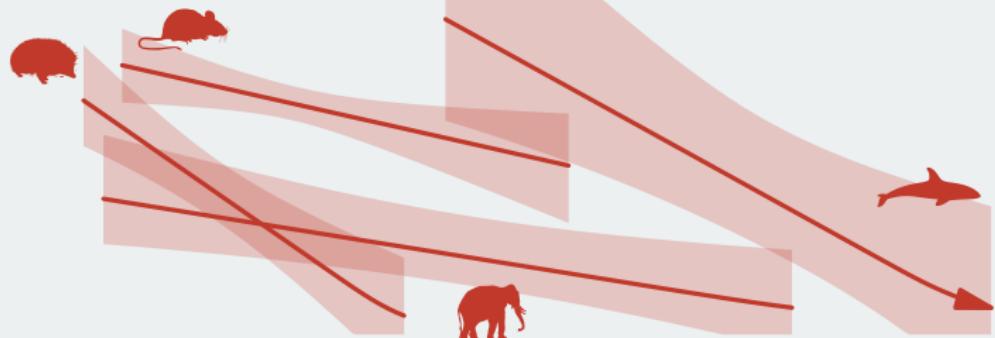
1 g

1 kg

1 tonne

100 tonnes

Body mass (g; log scale)



Speciation rate

0.3

0.0

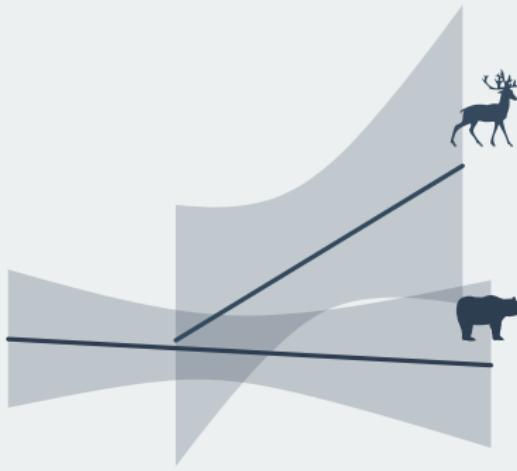
1 g

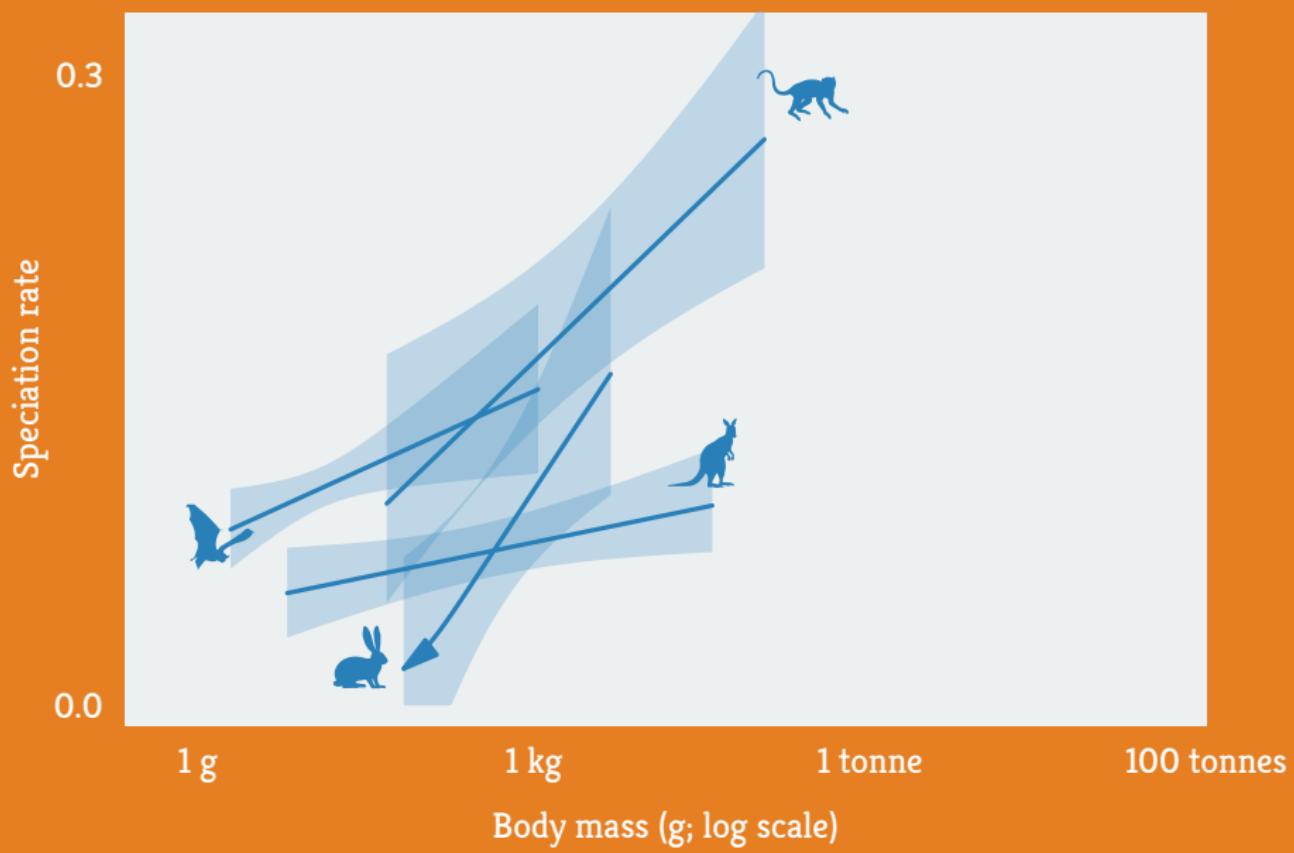
1 kg

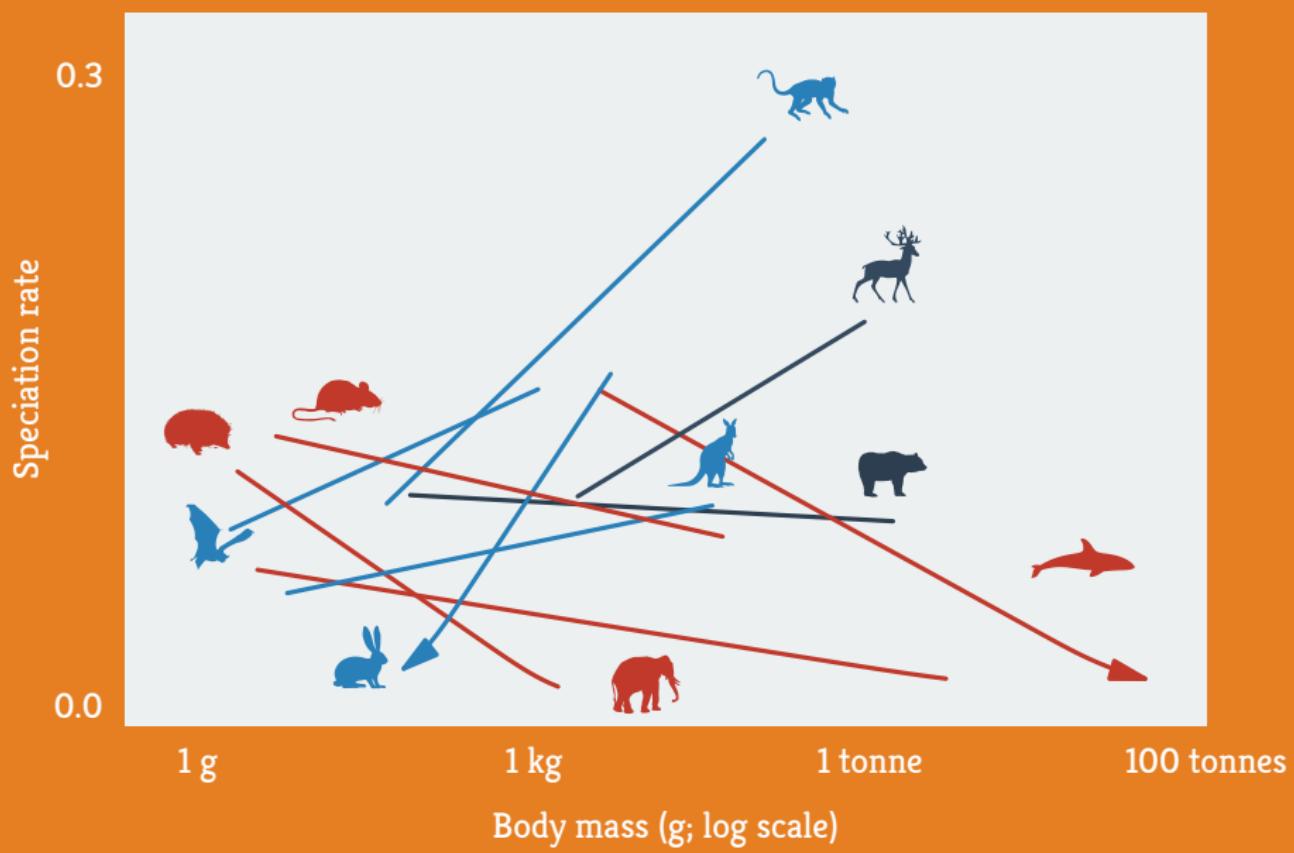
1 tonne

100 tonnes

Body mass (g; log scale)





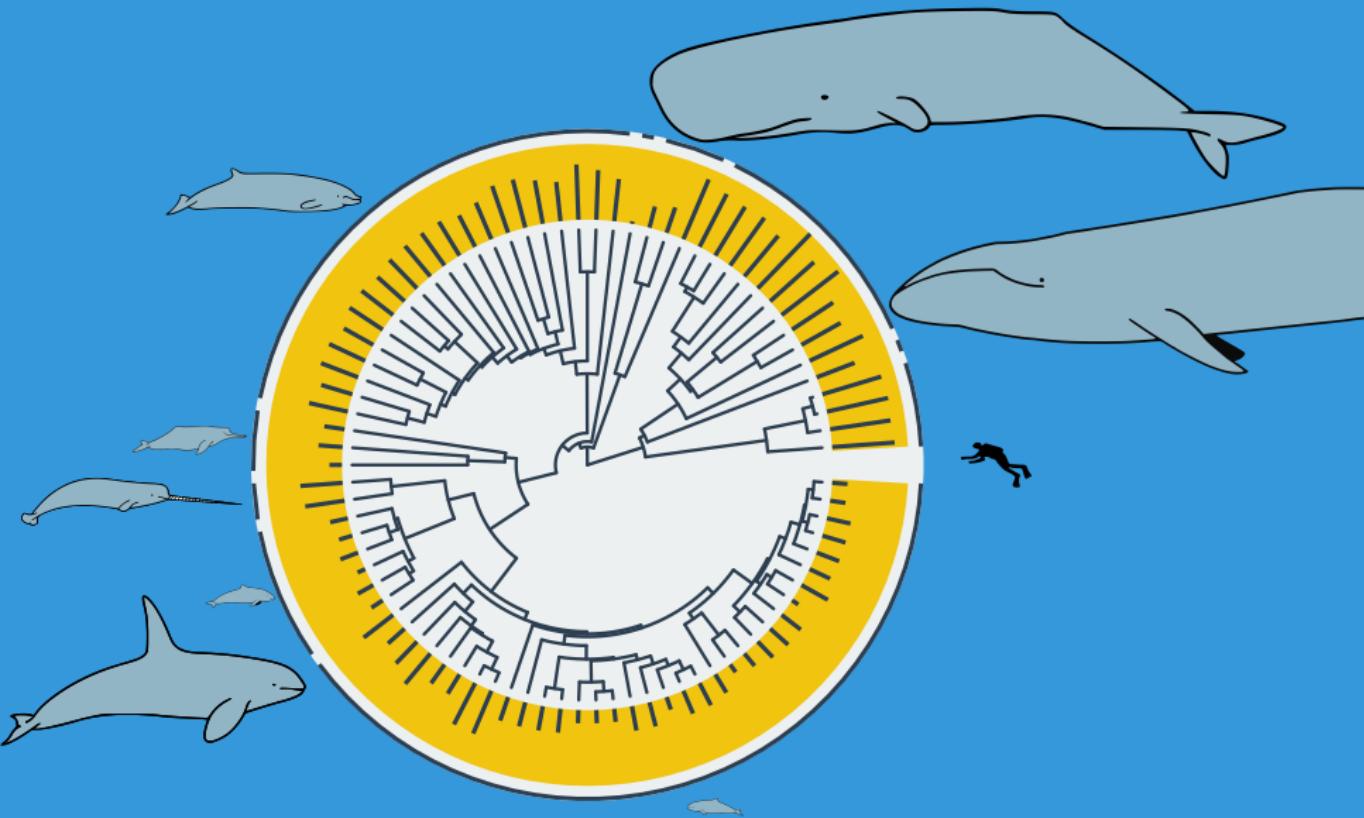


Why are there
so many small
mammal species?

Probably not species selection

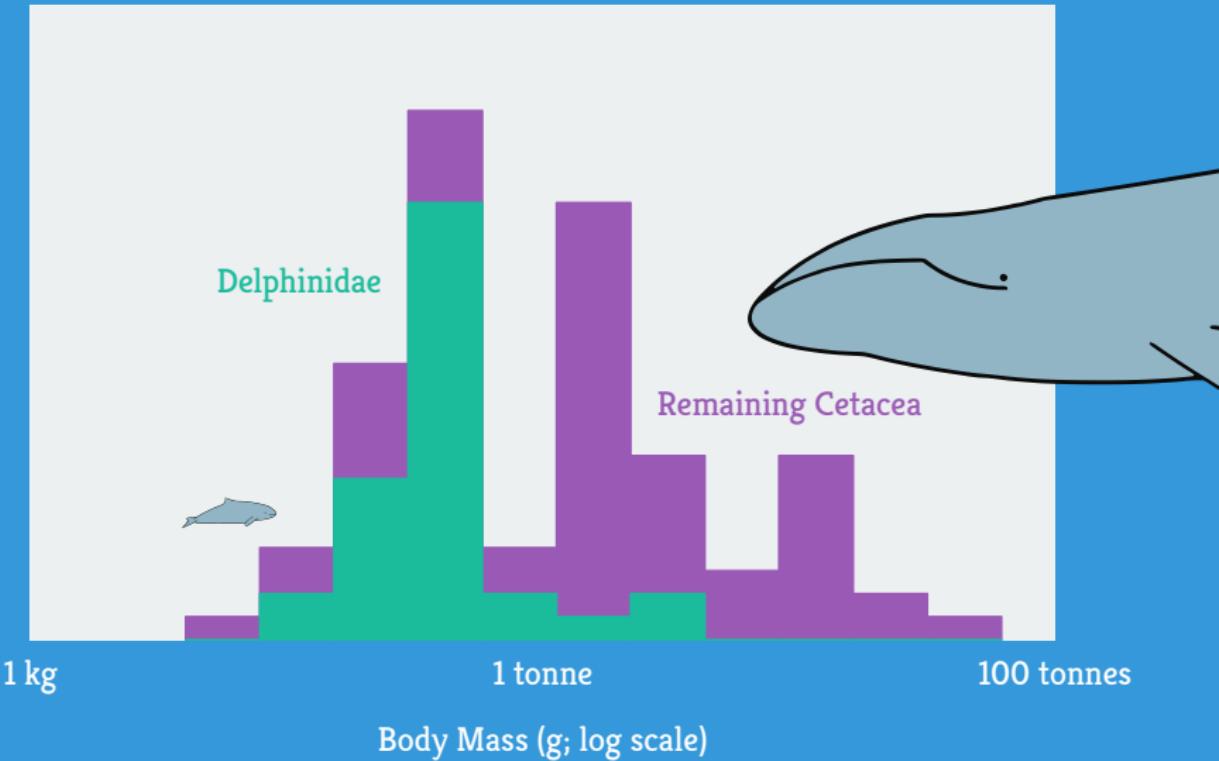
LIMITATIONS OF THESE METHODS

**SIGNIFICANT
VS
REPLICATED**

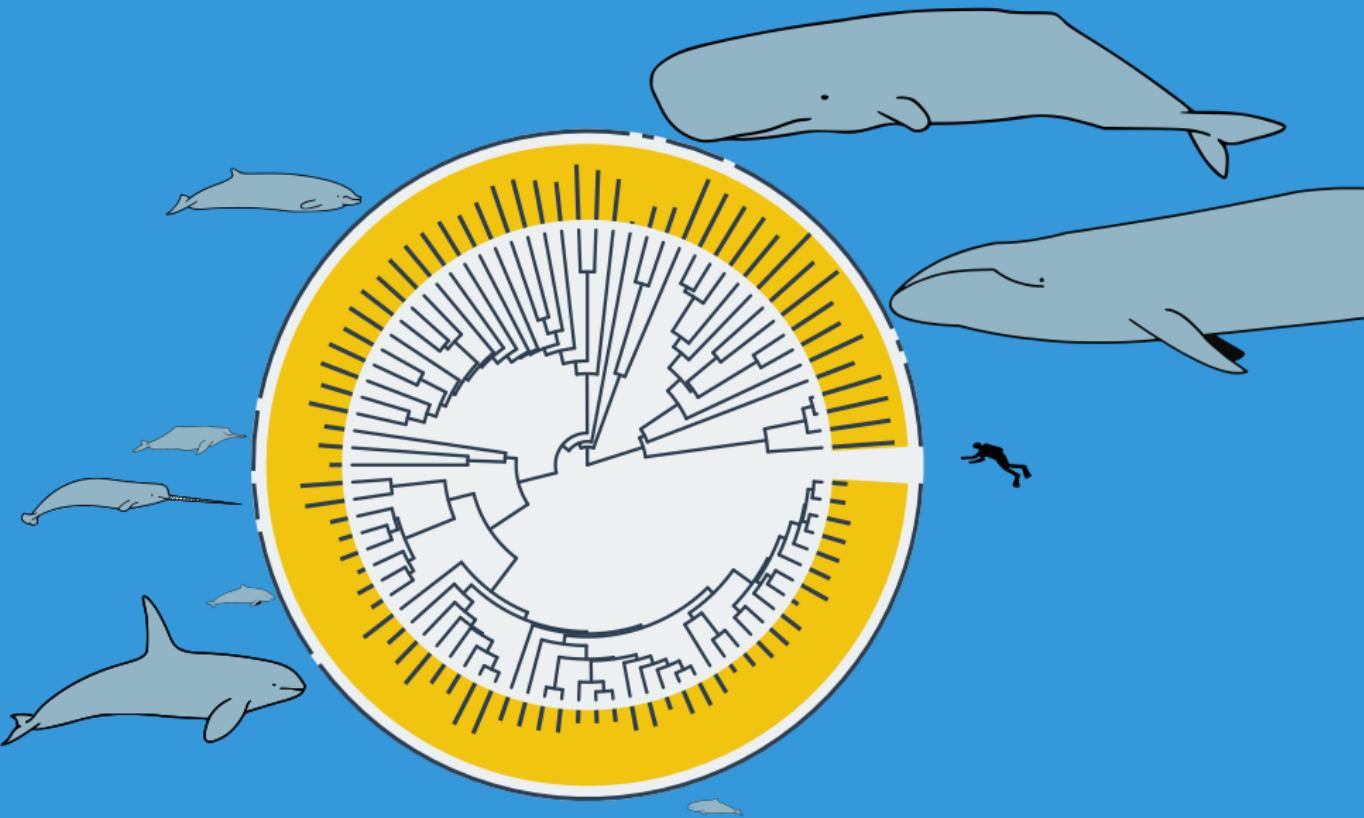


Tree: Steeman et al. 2009

Data: Nick Pyenson, unpublished



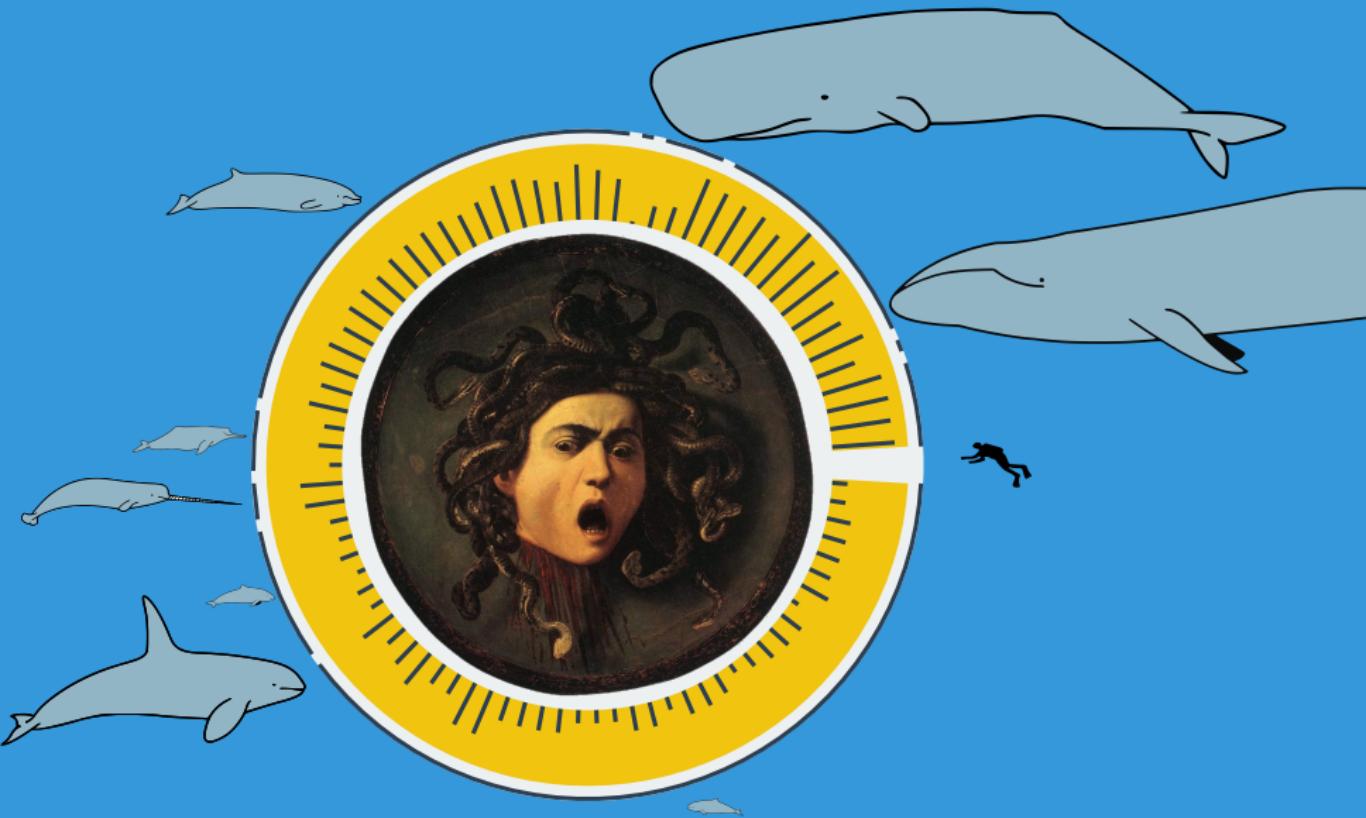
Data: Nick Pyenson, unpublished

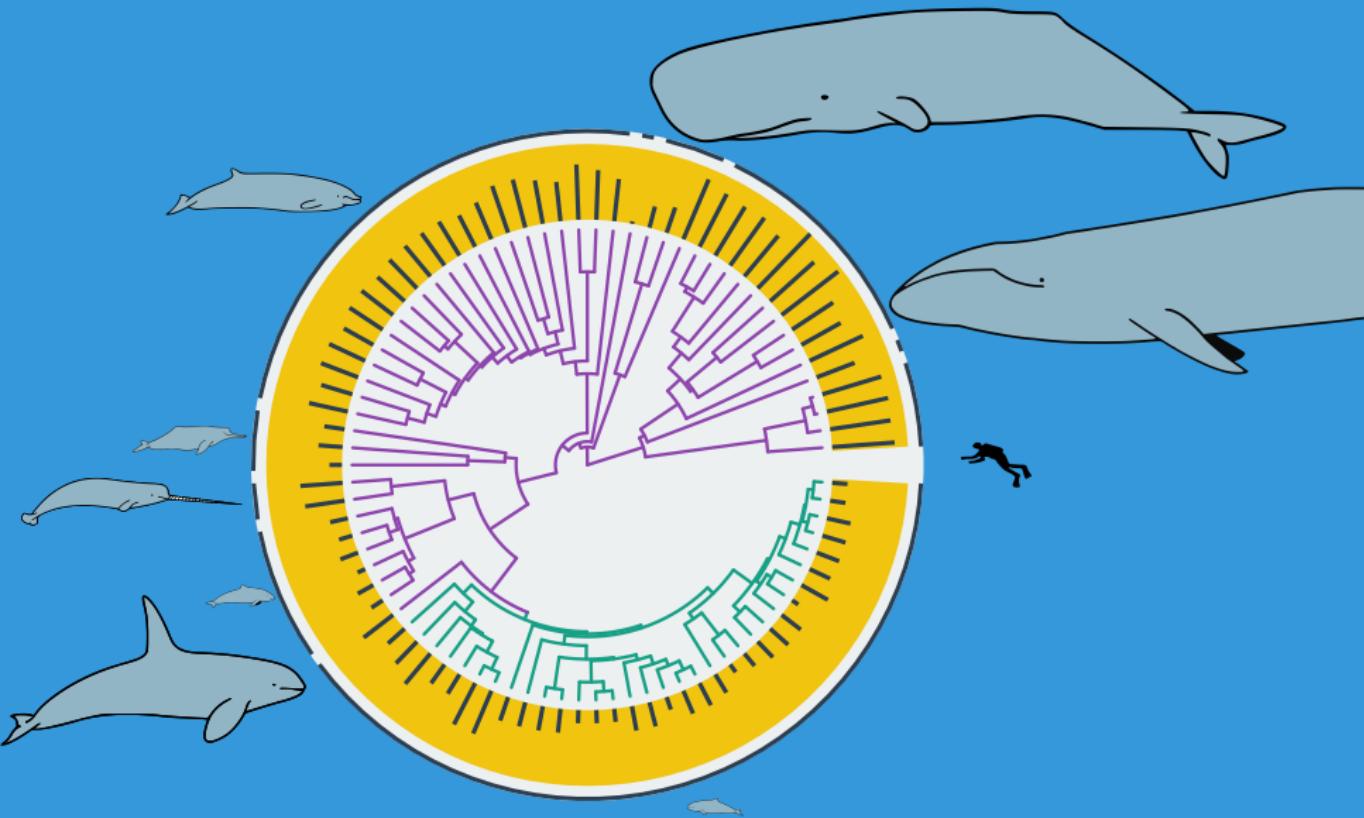


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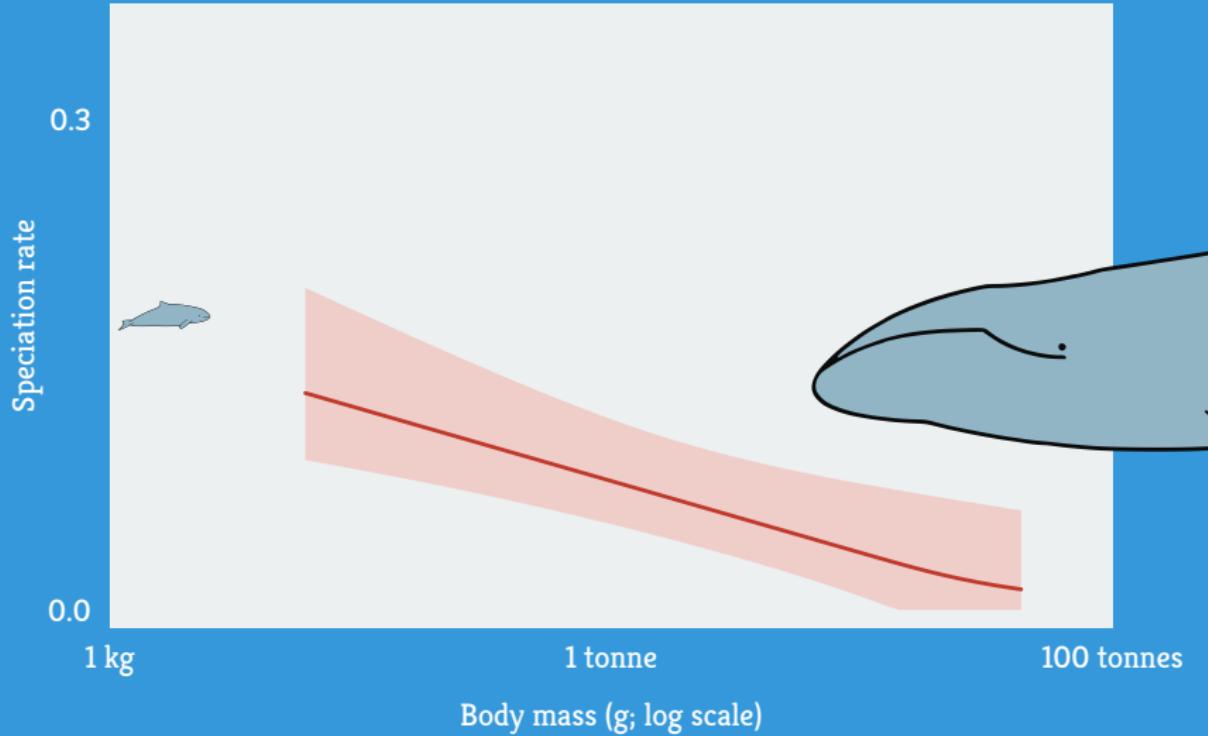
MEDUSA

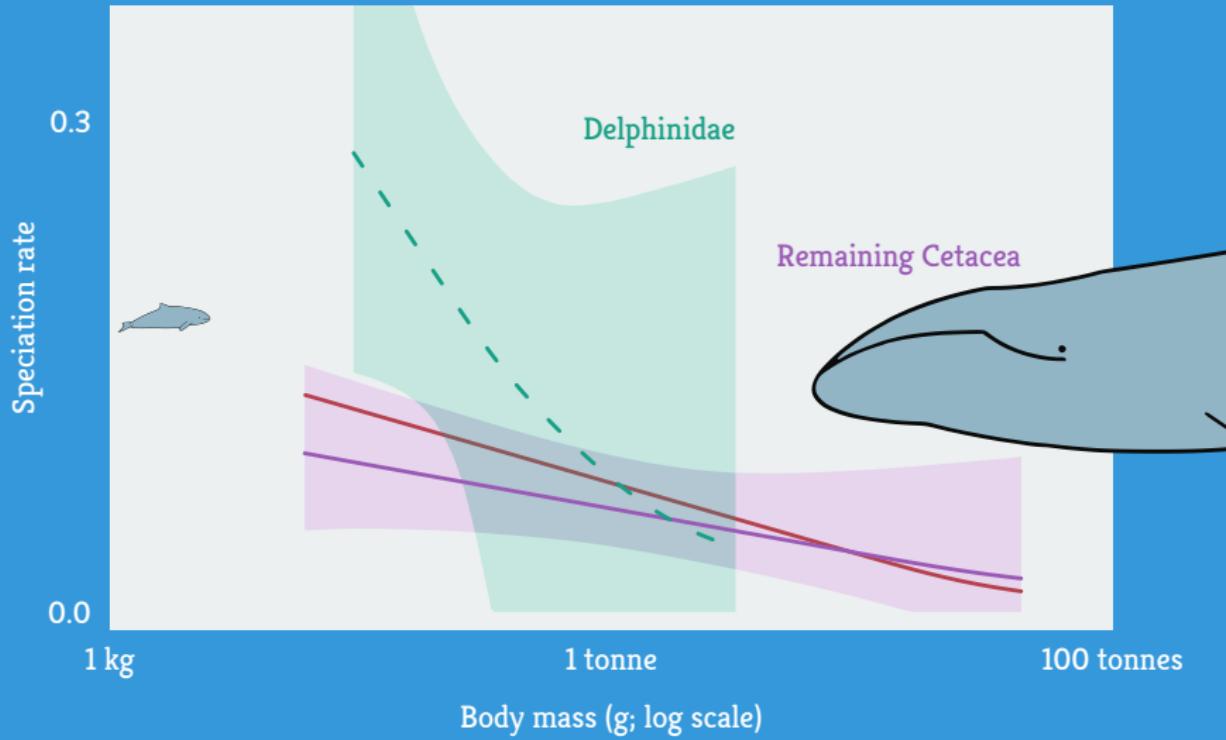


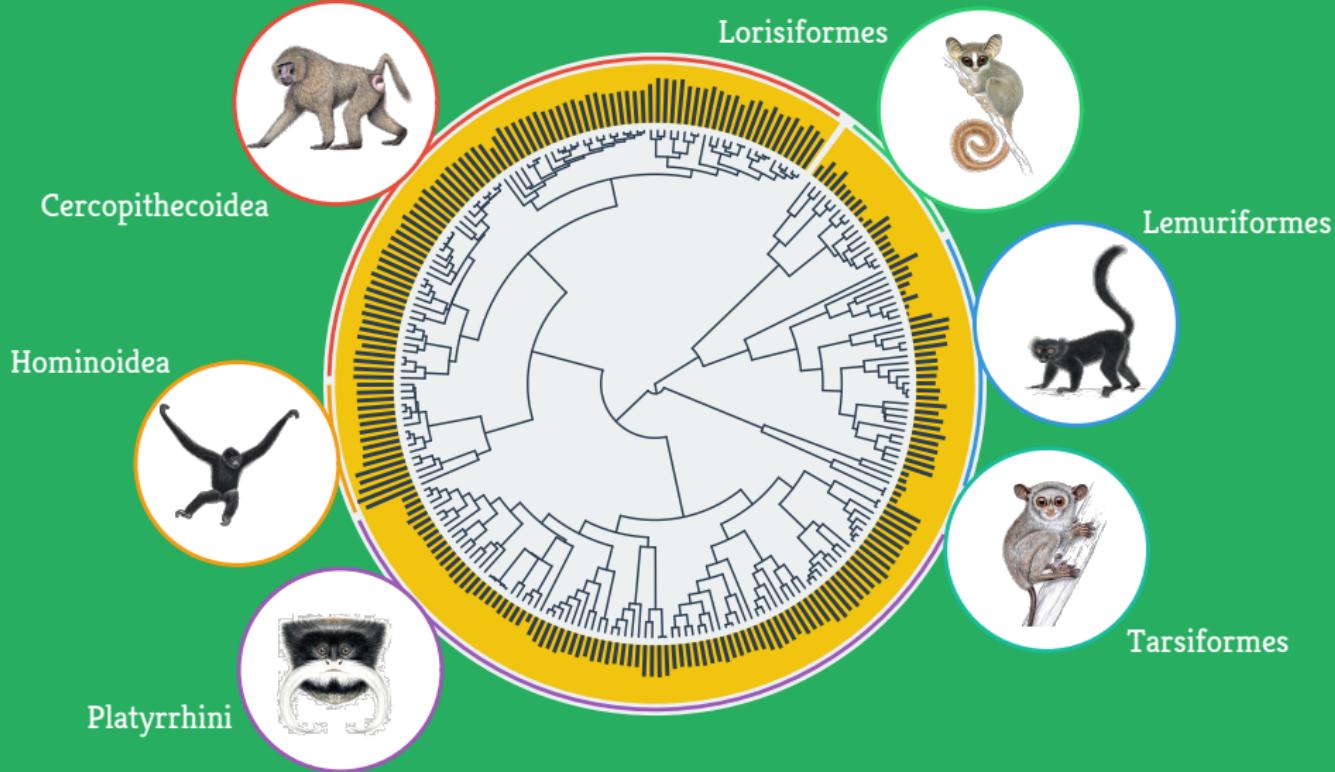


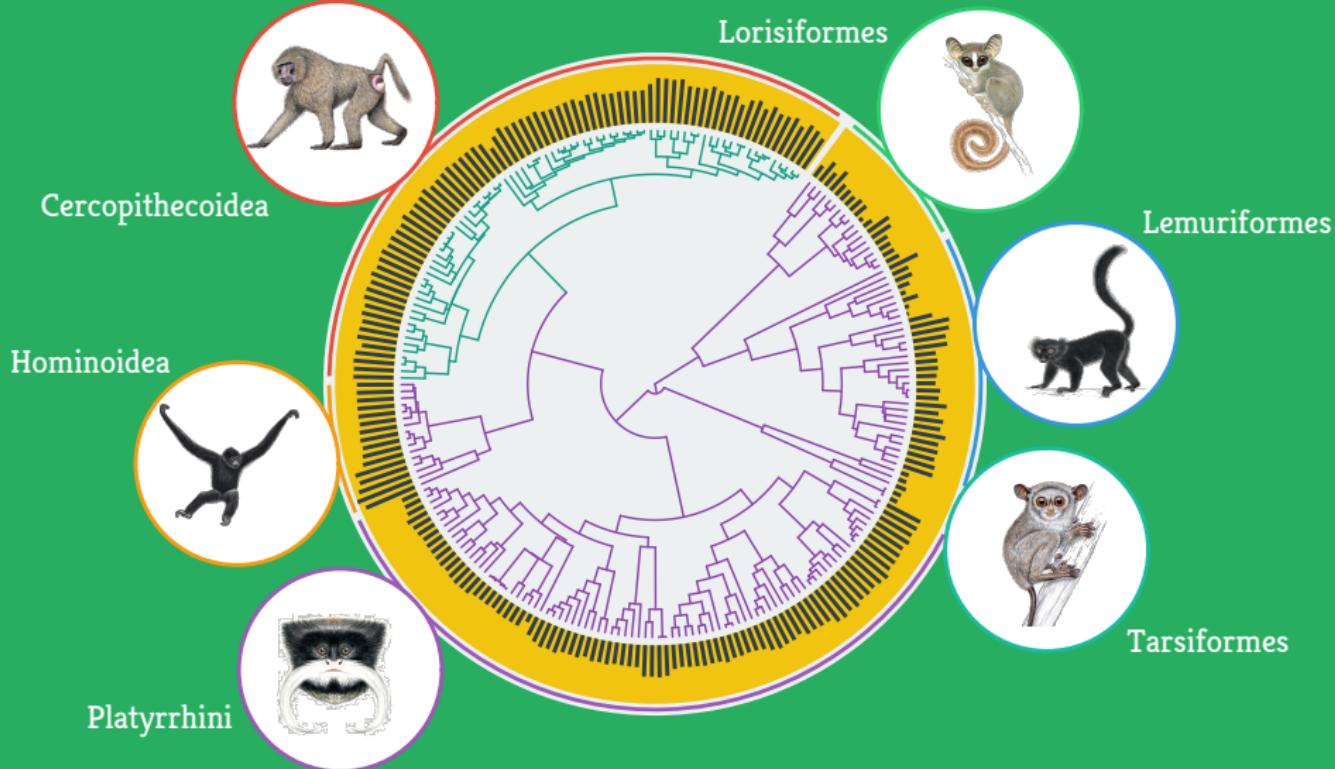
Tree: Steeman et al. 2009

Data: Nick Pyenson, unpublished









Speciation rate

0.5

0.0

1 kg

1 tonne

Body mass (g; log scale)



Speciation rate

0.5

Cercopithecidae



0.0

1 kg

Body mass (g; log scale)

Remaining Primates

1 tonne

Number of groups

30

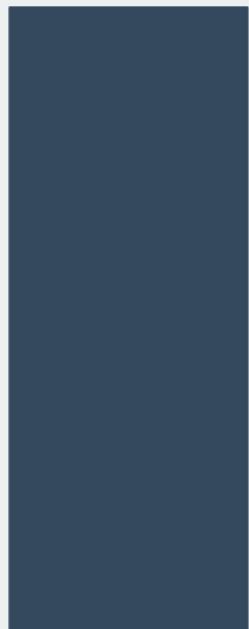
0

Negative

None

Positive

Speciation rate vs. body size relationship



Number of groups

30

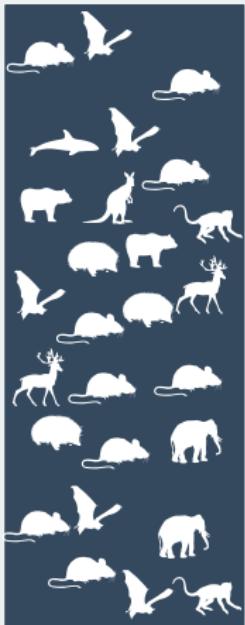
0

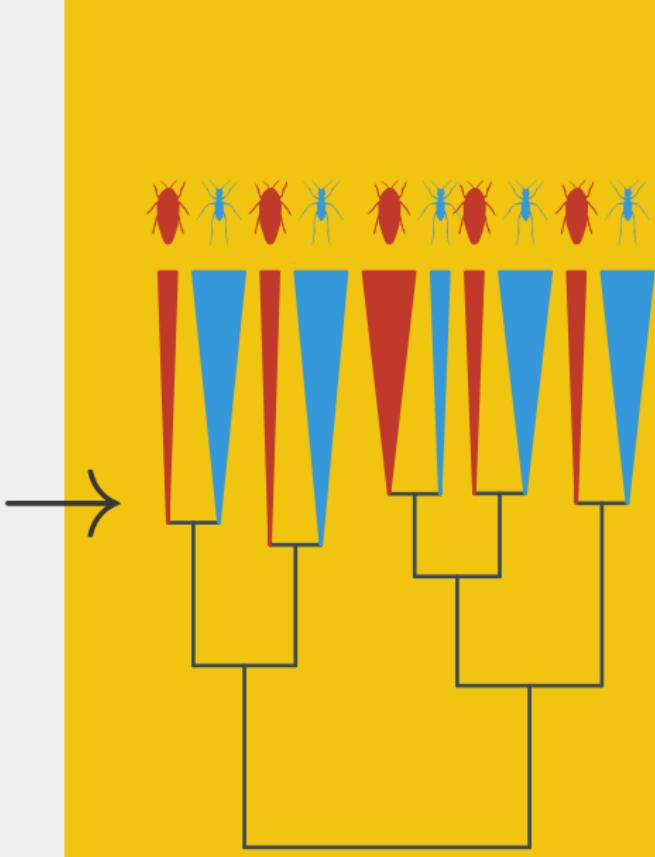
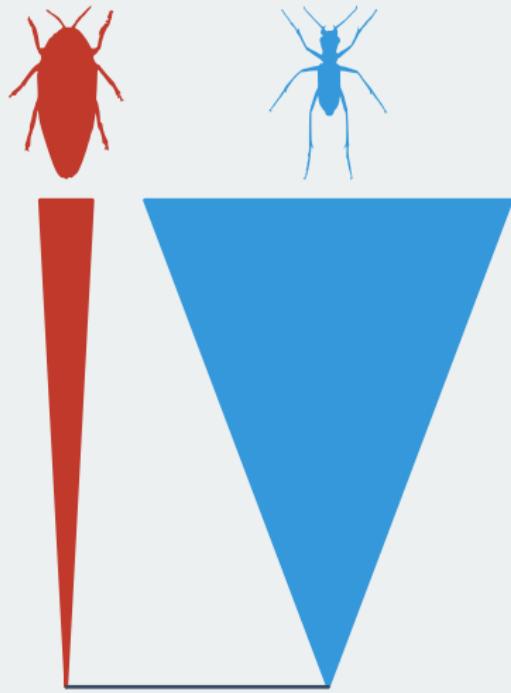
Negative

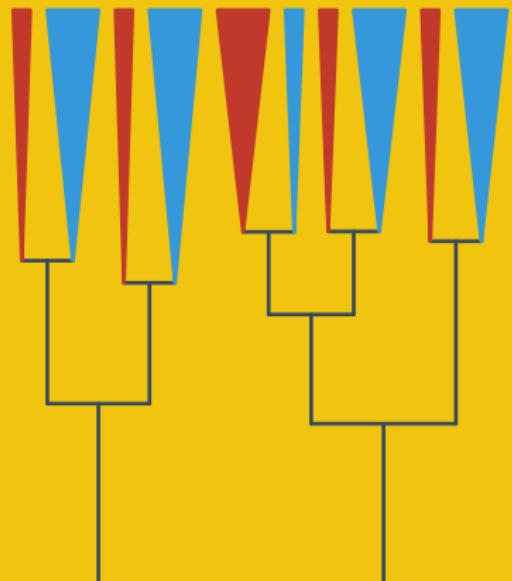
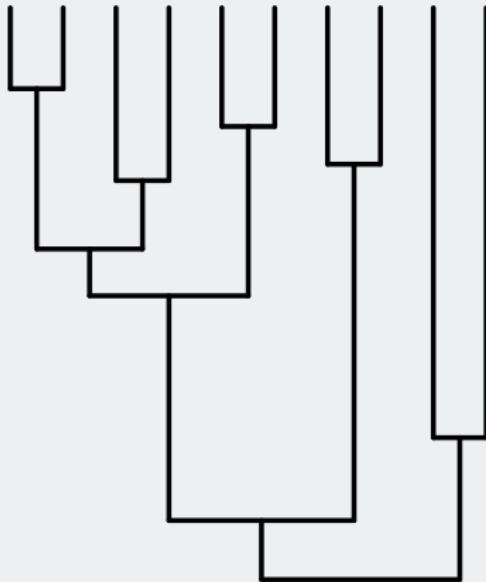
None

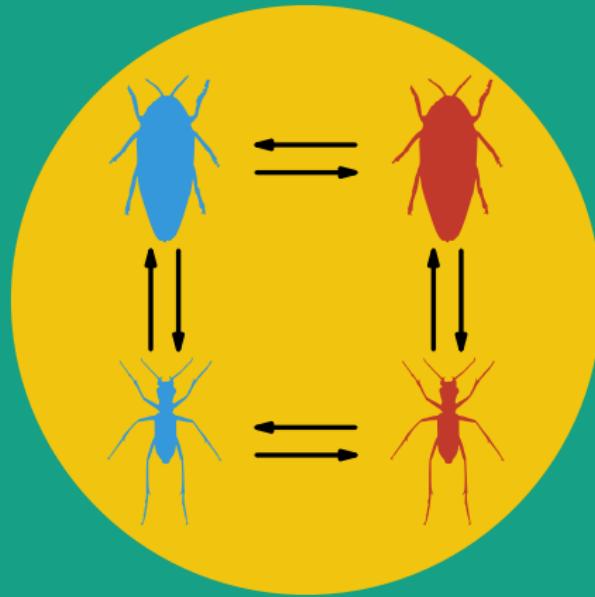
Positive

Speciation rate vs. body size relationship



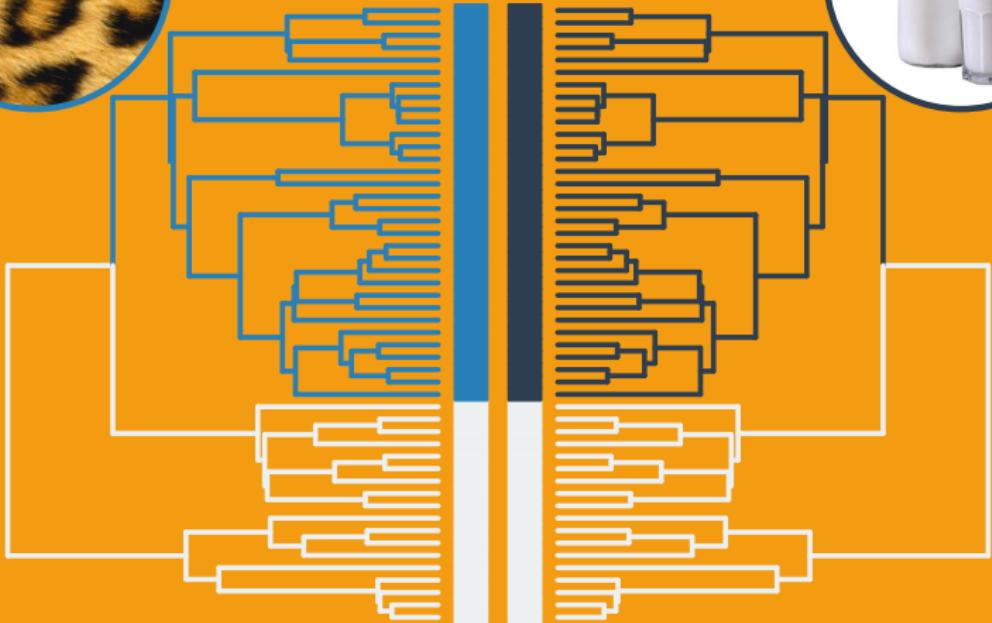


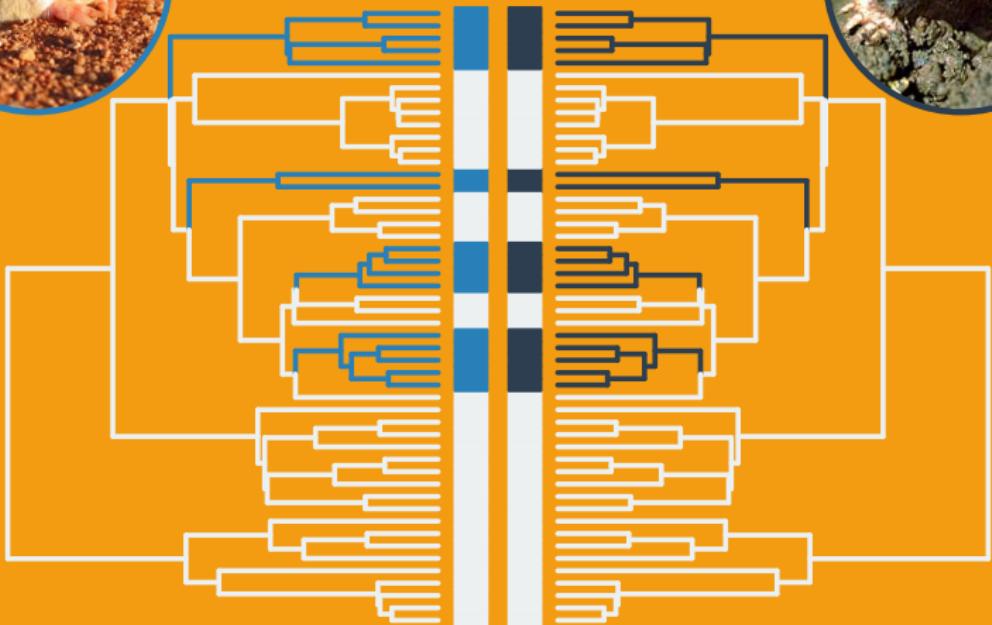


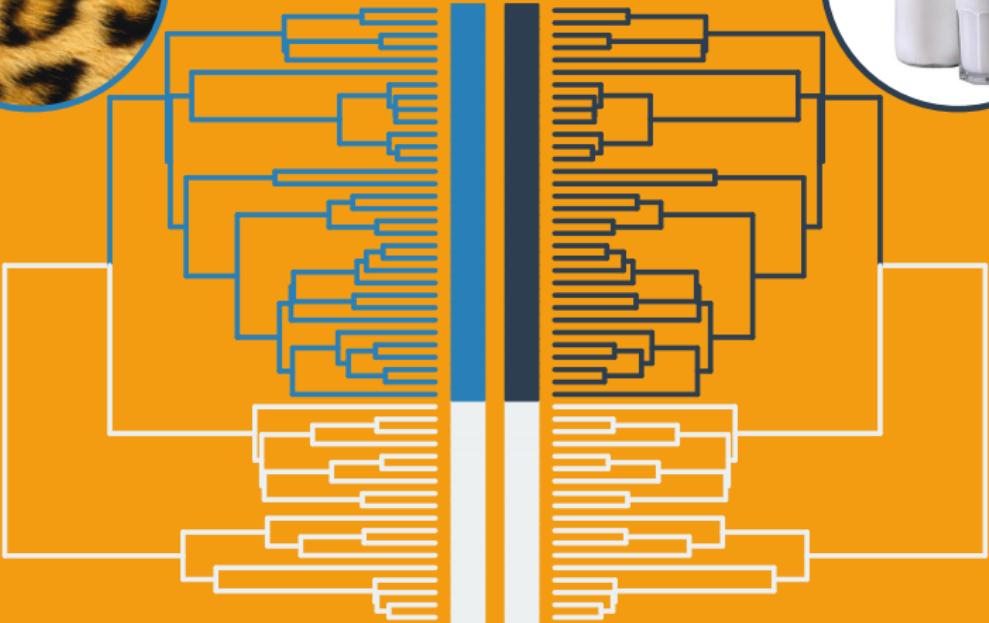


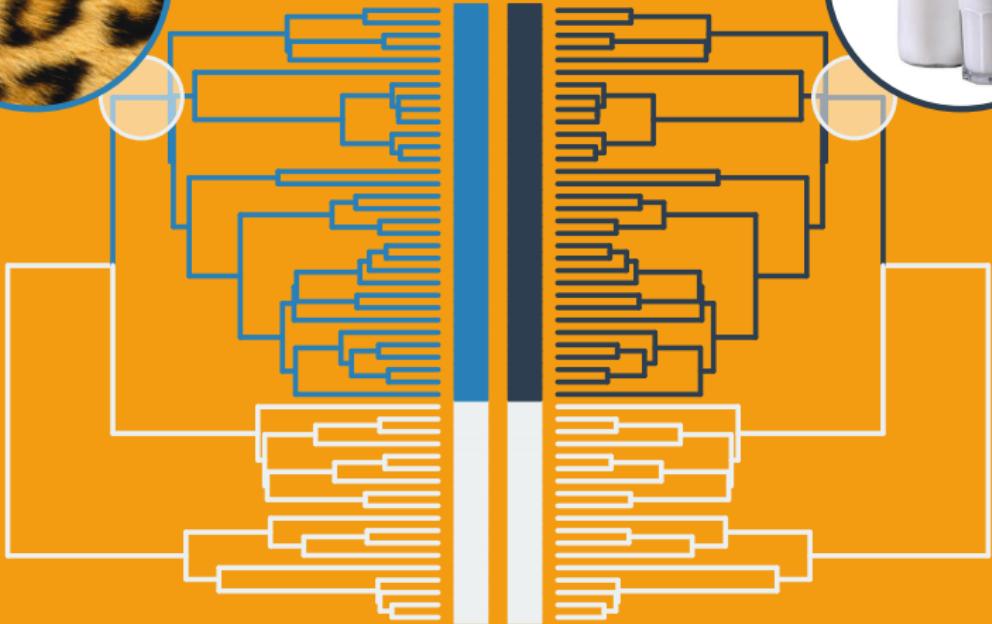
with Wayne Maddison





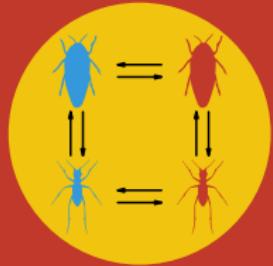
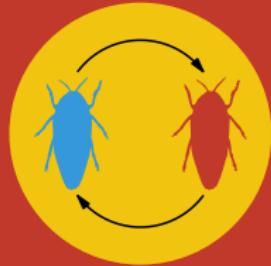




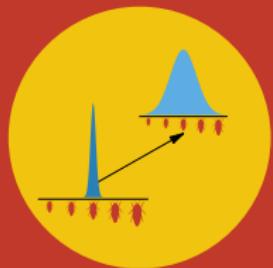


**WE HAVE
VERY SIMPLE MODELS**

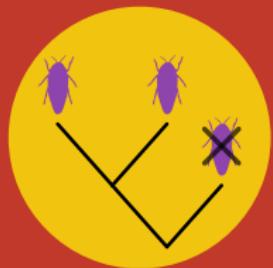
Morphological traits
do not evolve like
DNA



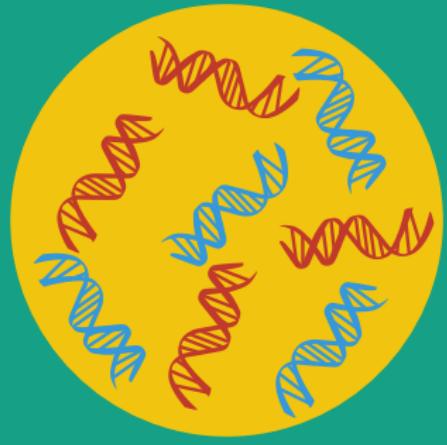
Quantitative traits
do not follow
Brownian motion

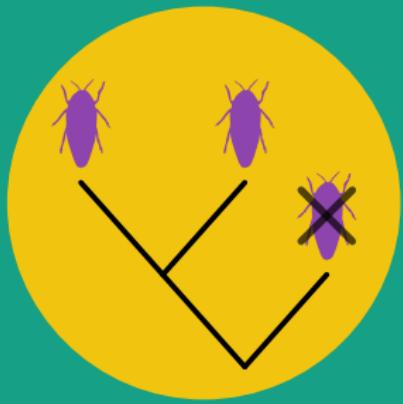


Diversification
does not follow
a birth death process

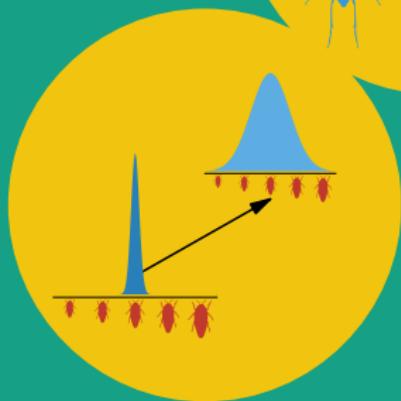
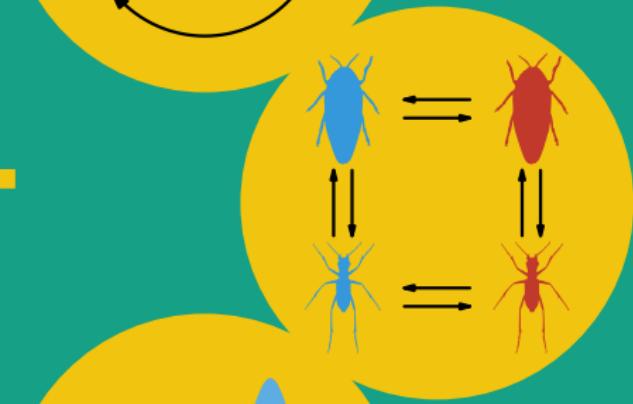
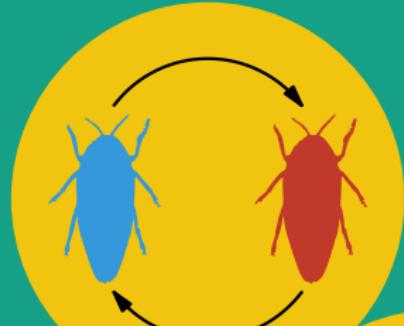


SUMMARY





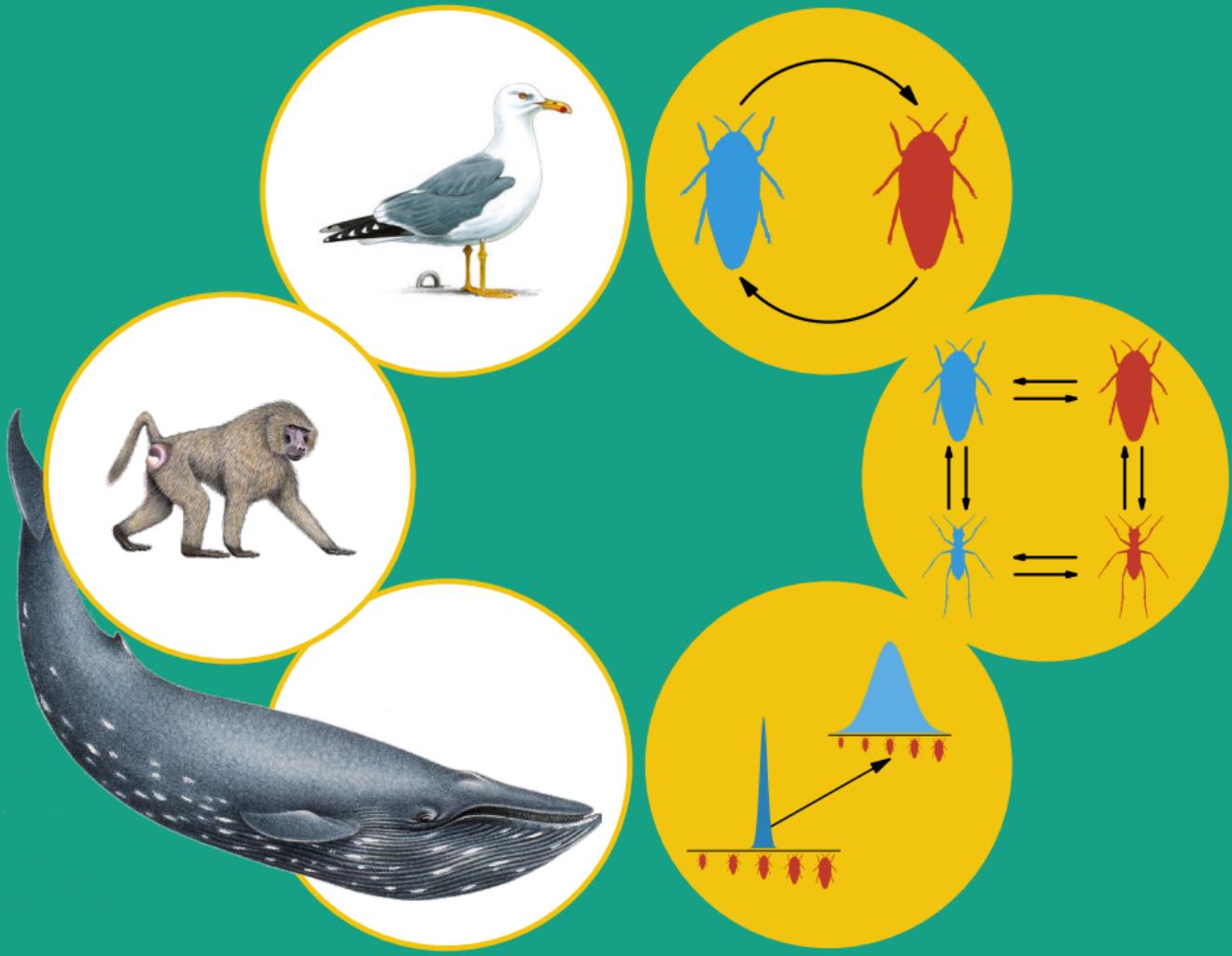
+





diversitree

www.zoology.ubc.ca/prog/diversitree
github.com/richfitz/diversitree



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Rutger Vos, David Redding, Arne Mooers (primates)

Olaf Bininda-Emonds, Kate Jones, Nick Isaac (all mammals)

Mette Steeman, Nick Pyenson (Cetacea)



**NSERC
CRSNG**



Images: Biodiversity Heritage Library: flickr.com/photos/biodivlibrary

Mean speciation rate

0.6



Canidae
(Carnivora)



Rhinolophus
(Chiroptera)



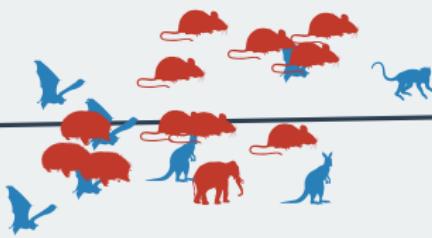
Microgale
(Afrotheria)



Marmotini
(Rodentia)



Cercopithecidae
(Primates)



1 g

1 kg

1 tonne

Median body mass (g; log scale)