Macroevolutionary perspectives to environmental change

Fabien Condamine

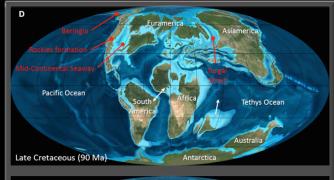
CNRS, UMR 7641 Centre de Mathématiques Appliquées (Ecole Polytechnique)

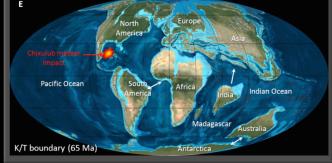












Human activities are generating major environmental changes

REVIEW

Introduction

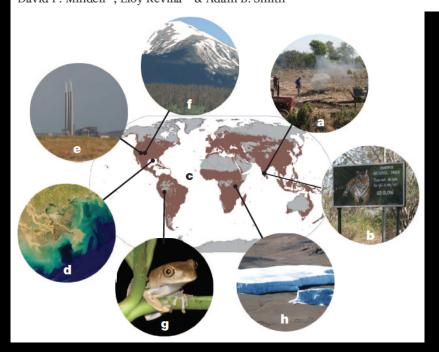
doi:10.1038/nature11018

Approaching a state shift in Earth's biosphere

Anthony D. Barnosky^{1,2,3}, Elizabeth A. Hadly⁴, Jordi Bascompte⁵, Eric L. Berlow⁶, James H. Brown⁷, Mikael Fortelius⁸, Wayne M. Getz⁹, John Harte^{9,10}, Alan Hastings¹¹, Pablo A. Marquet^{12,13,14,15}, Neo D. Martinez¹⁶, Arne Mooers¹⁷, Peter Roopnarine¹⁸, Geerat Vermeij¹⁹, John W. Williams²⁰, Rosemary Gillespie⁹, Justin Kitzes⁹, Charles Marshall^{1,2}, Nicholas Matzke¹, David P. Mindell²¹, Eloy Revilla²² & Adam B. Smith²³

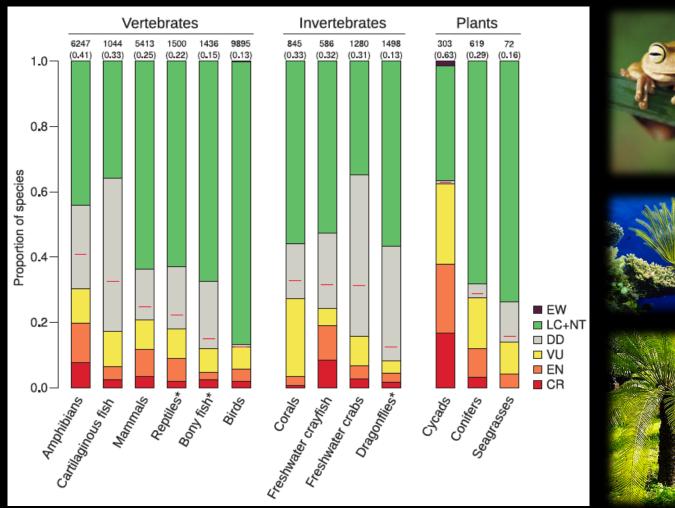
Ex.

- global warming
- habitat loss
- hunting
- increased UV-radiation
- overexploitation
- pollution





This exerts high pressure on ecosystems and biodiversity





Amphibians = 41%



Corals = 33%



Cycads = 63%

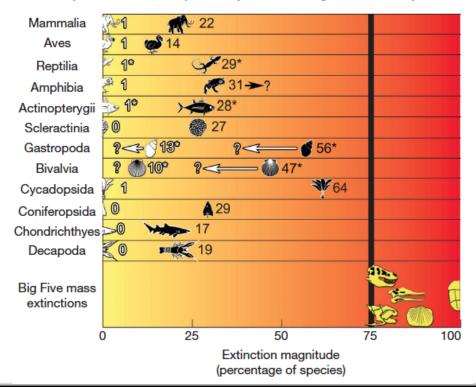
Paleontologists think we are entering a 'mass extinction'

REVIEW

doi:10.1038/nature09678

Has the Earth's sixth mass extinction already arrived?

Anthony D. Barnosky^{1,2,3}, Nicholas Matzke¹, Susumu Tomiya^{1,2,3}, Guinevere O. U. Wogan^{1,3}, Brian Swartz^{1,2}, Tiago B. Quental^{1,2}†, Charles Marshall^{1,2}, Jenny L. McGuire^{1,2,3}†, Emily L. Lindsey^{1,2}, Kaitlin C. Maguire^{1,2}, Ben Mersey^{1,4} & Elizabeth A. Ferrer^{1,2}

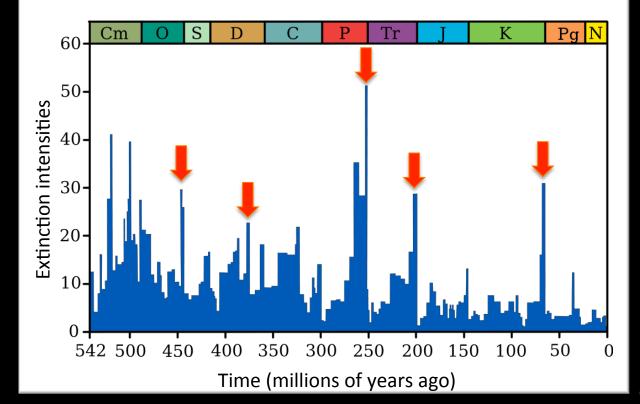


✓ Mass extinctions = times when the Earth loses 75% of its species in geologically short interval

Mass Extinctions in the Marine Fossil Record

Introduction

Abstract. A new compilation of fossil data on invertebrate and vertebrate families indicates that four mass extinctions in the marine realm are statistically distinct from background extinction levels. These four occurred late in the Ordovician, Permian, Triassic, and Cretaceous periods. A fifth extinction event in the Devonian stands out from the background but is not statistically significant in these data. Background extinction rates appear to have declined since Cambrian time, which is consistent with the prediction that optimization of fitness should increase through evolutionary time.



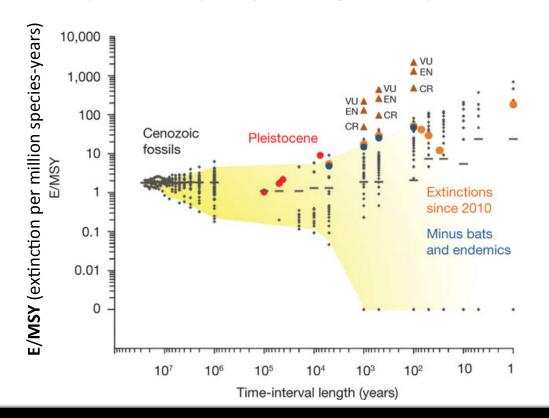
 ✓ Occurred only 5 times in the past 542 million years ('Big Five')

Ex. Most drastic event was 252 Ma (end-Permian) with 80-96% species loss

Current biodiversity crisis results from a decoupling between speciation and extinction

Has the Earth's sixth mass extinction already arrived?

Anthony D. Barnosky^{1,2,3}, Nicholas Matzke¹, Susumu Tomiya^{1,2,3}, Guinevere O. U. Wogan^{1,3}, Brian Swartz^{1,2}, Tiago B. Quental^{1,2}†, Charles Marshall^{1,2}, Jenny L. McGuire^{1,2,3}†, Emily L. Lindsey^{1,2}, Kaitlin C. Maguire^{1,2}, Ben Mersey^{1,4} & Elizabeth A. Ferrer^{1,2}



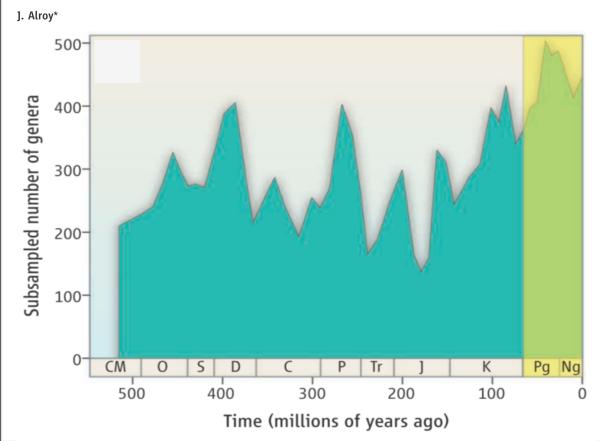
- ✓ Current extinction rates are higher than in the fossil record
- ✓ Species go extinct 1000 times faster

To understand the current biodiversity crisis, we need to understand its past long-term dynamics

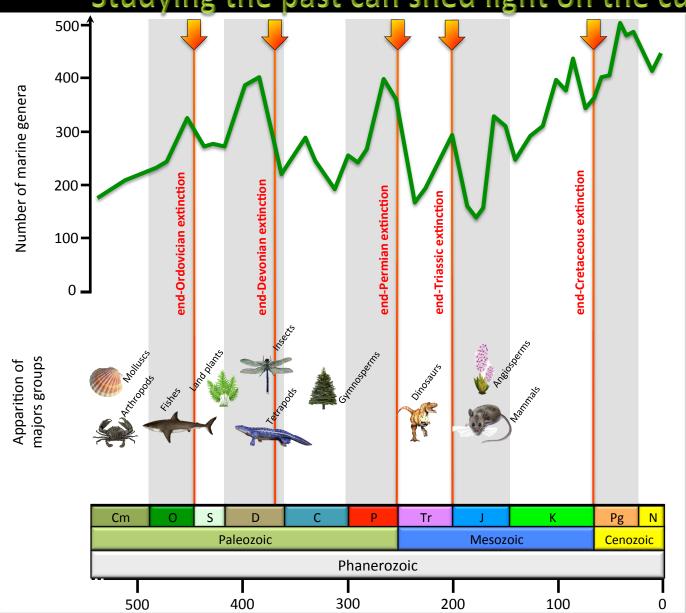
The Shifting Balance of Diversity Among Major Marine Animal Groups

Mass extinctions &

recovery



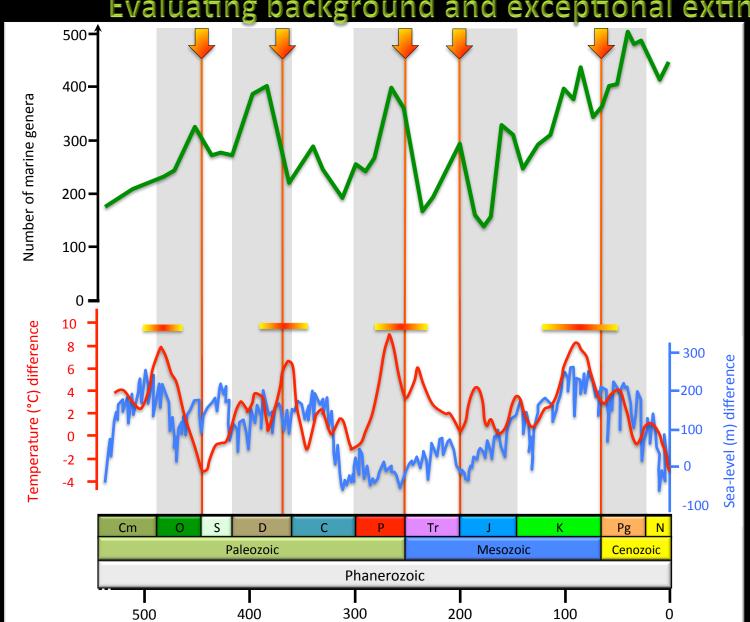
- ✓ Historically done with the marine fossil record
- ✓ **Direct evidence** of diversity dynamics through time



✓ Originations but also extinctions are part of the history of life

Condamine et al. 2013 – Ecol. Letters



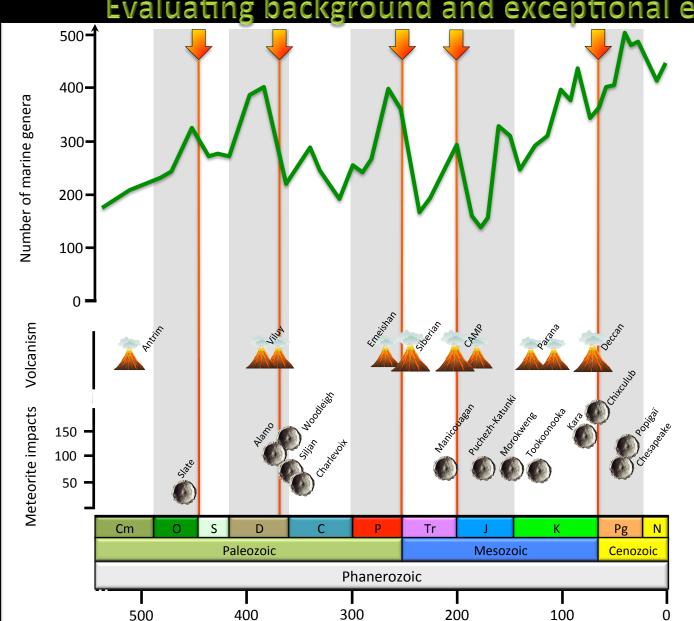


To better:

Understand causes of extinction due to climatic changes

Condamine et al. 2013 – Ecol. Letters





To better:

Understand causes of extinction due to geological changes

Condamine et al. 2013 – Ecol. Letters Background speciation & extinction

Vulnerability & evolutionary potential

Limitations & Perspectives

Profusion of evidence from the marine fossil record

Michael Benton

Introduction



Shanan Peters



ARTICLE

Diversification and Extinction in the History of Life

M. J. Benton

Speciation

REVIEW

The Red Queen and the Court Jester: Species Diversity and the Role of Biotic and Abiotic Factors Through Time

Michael 1. Benton

nature geoscience

REVIEW ARTICLE
PUBLISHED ONLINE: 27 MAY 2012 | DOI: 10.1038/NGE01475

The timing and pattern of biotic recovery following the end-Permian mass extinction

Zhong-Qiang Chen¹ and Michael J. Benton^{2*}

Benton 1995 – Science Benton 2009 – Science Chen & Benton 2012 – Nature Geosci. Geologic constraints on the macroevolutionary history of marine animals

Shanan E. Peters*

Department of Geological Sciences and Museum of Paleontology, University of Michigan, 1109 South Geddes Road, Ann Arbor, MI 48104

Edited by W. A. Berggren, Woods Hole Oceanographic Institution, Woods Hole, MA, and approved July 15, 2005 (received for review March 30, 2005)

nature

Vol 454 31 July 2008 doi:10.1038/nature07032

LETTERS

Environmental determinants of extinction selectivity in the fossil record

Shanan E. Peters1



doi:10.1038/nature11815

Oceanographic controls on the diversity and extinction of planktonic foraminifera

Shanan E. Peters1, Daniel C. Kelly1 & Andrew J. Fraass1†

Peters 2005 – PNAS Peters 2008 – Nature Peters et al. 2013 – Nature

But the picture from the fossil record is not exhaustive



Fossil record is:

- Biased
- Incomplete
- Uneven

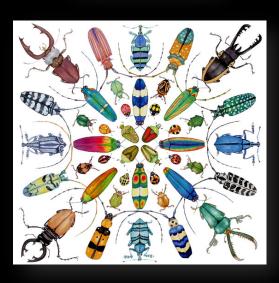


Many groups lack a suitable fossil record

e.g., insects, plants, birds...

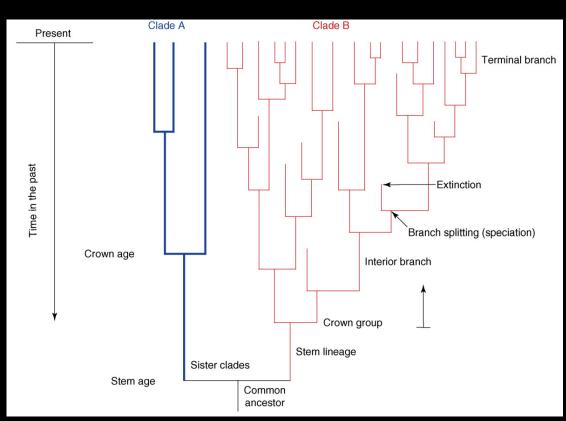
We need **more data** to better understand diversity dynamics in relation with environmental changes

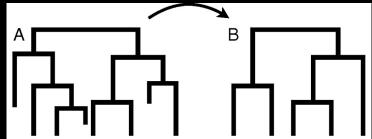






Using relationships between extant species and their divergence times

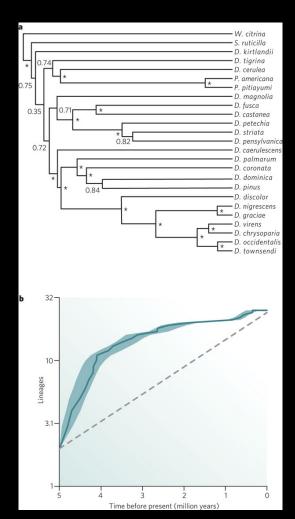


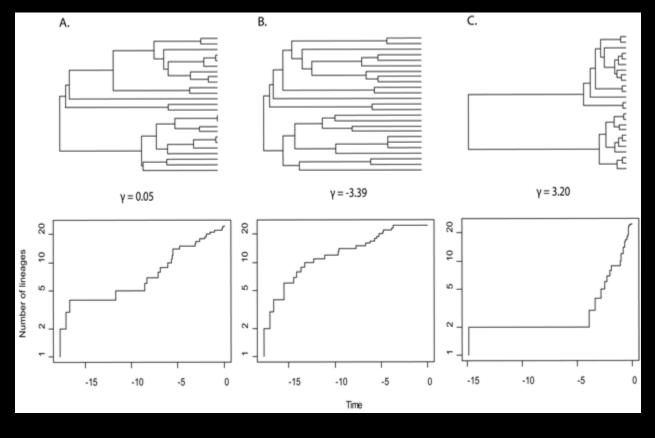


Phylogenetic approaches to diversification



Phylogenies can be converted into lineages-through-time (LTT) plot





ECOLOGY LETTERS

Ecology Letters, (2013)

doi: 10.1111/ele.12062

IDEA AND PERSPECTIVE

Introduction

Macroevolutionary perspectives to environmental change

Fabien L. Condamine¹ Jonathan Rolland¹ and Hélène Morlon¹

Abstract

Predicting how biodiversity will be affected and will respond to human-induced environmental changes is one of the most critical challenges facing ecologists today. Here, we put current environmental changes and their effects on biodiversity in a macroevolutionary perspective. We build on research in palaeontology and recent developments in phylogenetic approaches to ask how macroevolution can help us understand how environmental changes have affected biodiversity in the past, and how they will affect biodiversity in the future. More and more paleontological and phylogenetic data are accumulated, and we argue that much of the potential these data have for understanding environmental changes remains to be explored.

Keywords

Biodiversity, birth-death models, diversification rates, extinction, fossils, global change, mass extinctions, paleoenvironment, speciation.

Ecology Letters (2013)

Macroevolutionary perspectives to environmental change

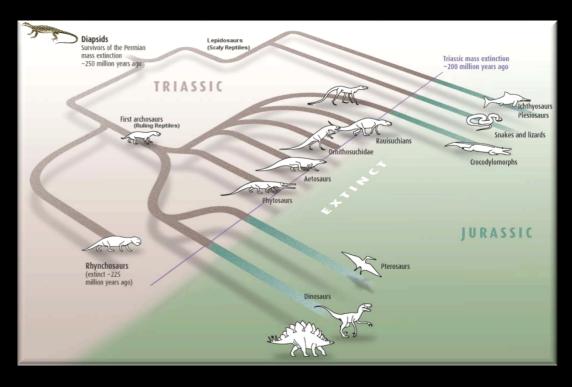
1 – Mass extinctions and recovery in relation to environmental change

2 – Background speciation and extinction in relation to environmental change

3 – Vulnerability and evolutionary potential

4 – Limitations and Perspectives

Mass extinctions and recovery in relation to environmental change





Why studying mass extinctions?



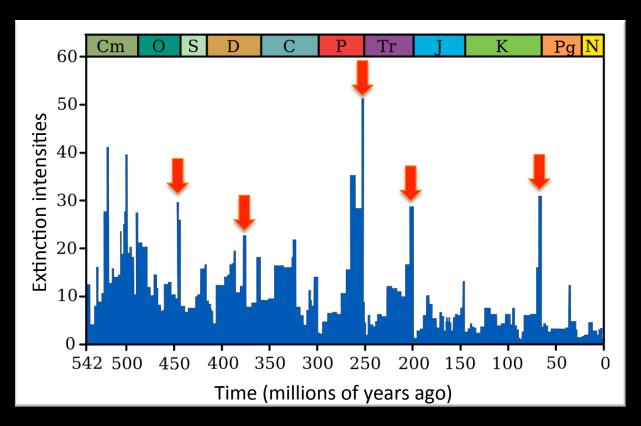


- 1. Useful to estimate when mass extinctions occurred
- 2. Useful to estimate how many species were lost (extinction intensity)
- 3. Useful to estimate which clades were impacted and what traits were associated with extinction (extinction selectivity)
- 4. Useful to estimate at which level of extinction biodiversity was able to recover

To find the causes of mass extinctions

Detecting mass extinctions in the fossil record

Paleontologists idenitified five mass extinctions:

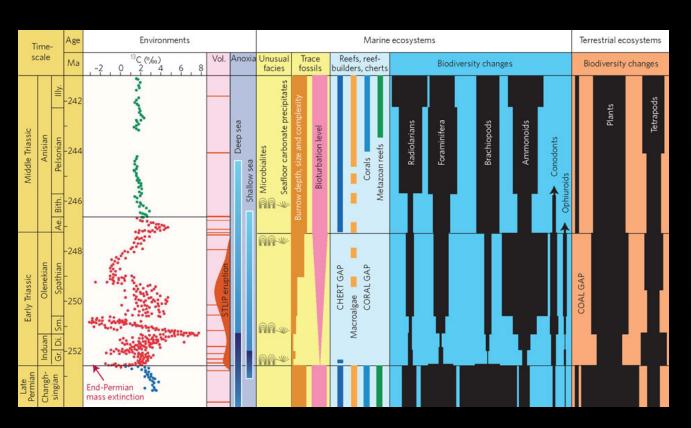


- 443 Ma with 86% species loss
- 359 Ma with 75% species loss
- 252 Ma with 95% species loss
- 201 Ma with 80% species loss
- 66 Ma with 76% species loss

Detecting mass extinctions in the fossil record



Paleontologists can study the causes of mass extinctions



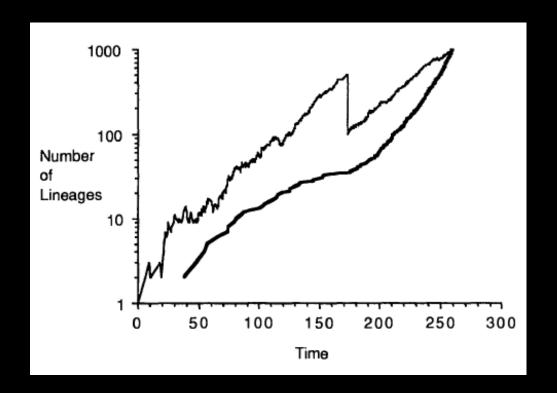








Theoretically possible to detect mass extinction in phylogenies



- ✓ Birth-death model can model the process of tree growth
- ✓ Adding a punctual event that remove a part of the species give a LTT interrupted by a plateau

Detecting mass extinctions in phylogenies



Few studies tested whether mass extinction can be found in phylogenies

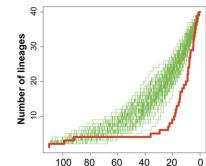
EXPLOSIVE RADIATION OR CRYPTIC MASS EXTINCTION? INTERPRETING SIGNATURES IN MOLECULAR PHYLOGENIES

Michael D. Crisp^{1,2} and Lyn G. Cook^{3,4}

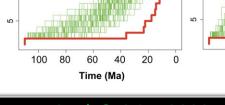
Mass Extinction, Gradual Cooling, or Rapid Radiation? Reconstructing the Spatiotemporal Evolution of the Ancient Angiosperm Genus Hedyosmum (Chloranthaceae) Using Empirical and Simulated Approaches

ALEXANDRE ANTONELLI^{1,2,*} AND ISABEL SANMARTÍN³

Number of Lineages 0.8 Relative time 10 30 10 100 20 Time (Ma)



10



Detecting mass extinctions in phylogenies

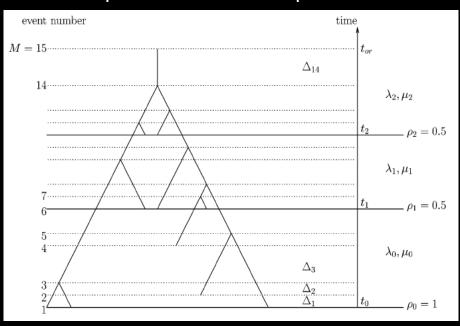


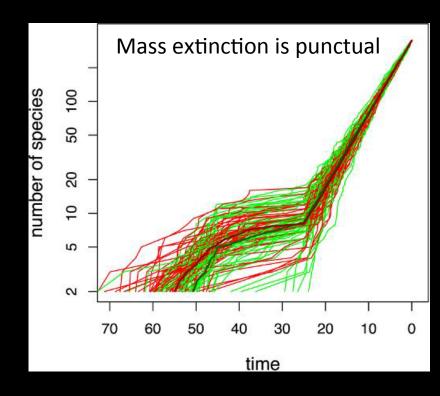
New methods (TreePar) have been developped to address this issue

Episodic birth-death process

Mass extinctions &

recovery







Case study: the Cycads, an ancient group of seed plants



300 palm-like species, distributed in all tropical regions



Appeared ca. 270 Ma, survived to three mass extinctions



63% of threatened species (IUCN's Red List)

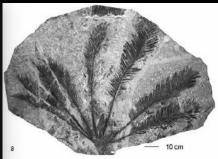


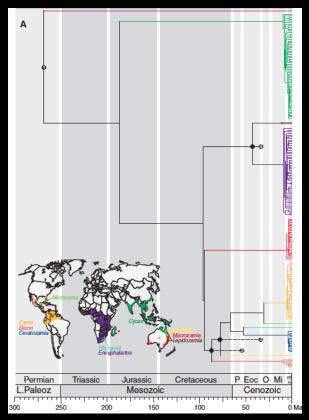
Cycad's distribution

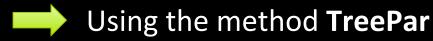
EW: Encephalartos woodii

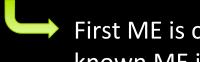


Triassic cycad





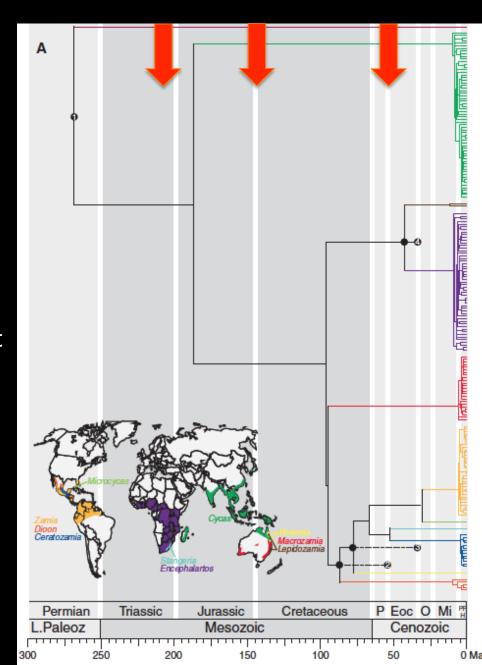




First ME is congruent with a known ME in the fossil record at 205 Ma (Triassic-Jurassic)

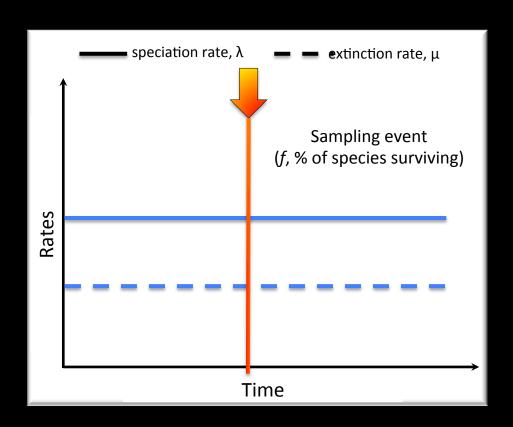
Second is a lesser ectinction event known from the fossil record at 135 Ma (Jurassic-Cretaceous)

Third is in the Cenozoic at 58 Ma and congruent with a global warming of climate





Speciation model with mass extinction and model with a stasis phase = same LTT plots

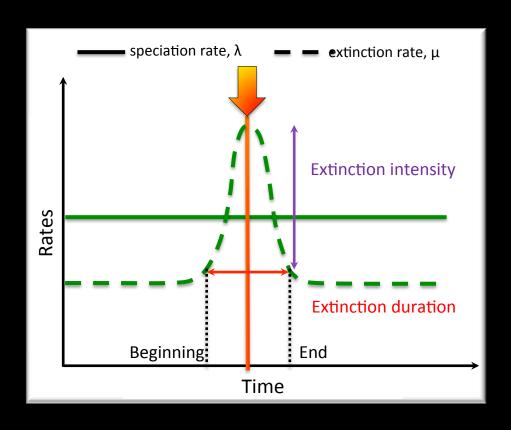


- ✓ Mass extinction modelled as an instantaneous event
- ✓ Classic rate shifts are instantaneous too
- ✓ Hard to distinguish between the two scenarios
- ✓ Same rates before and after the event

Problem with detecting mass extinctions in phylogenies



Mass extinctions have not necessary a short time duration



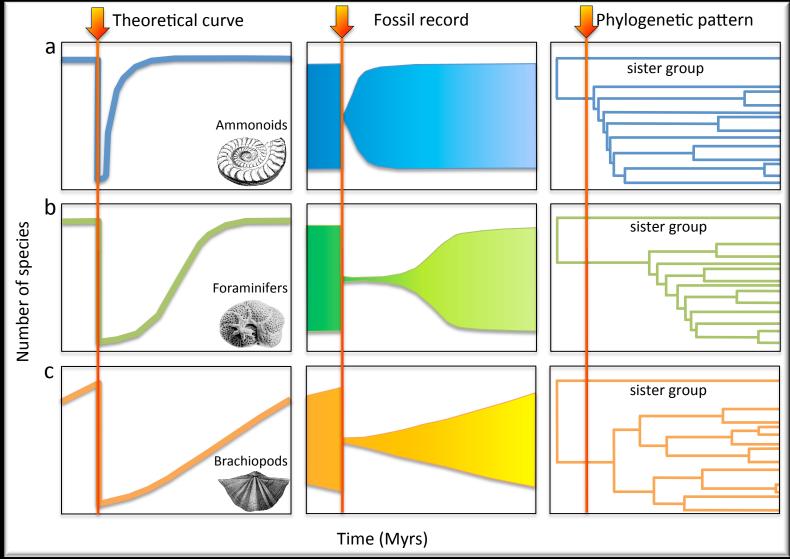
Mass extinctions &

recovery

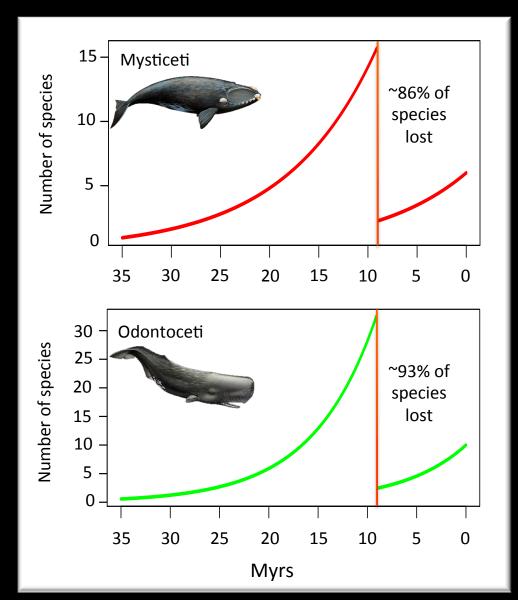
- Taking into account duration of the event
- Modelling extinction with continuous forms

Or:

- Background and mass extinction events modelled within the same framework
- Mass extinctions = extremes of a background continuum of intensities and duration



Recovery after mass extinctions

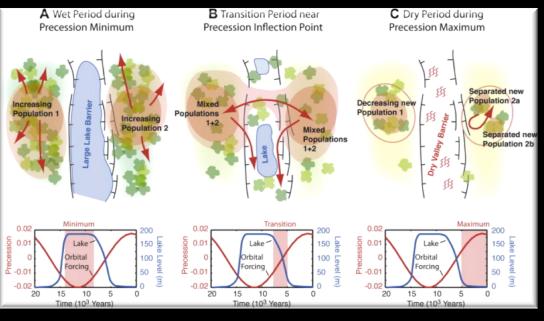


Mass extinctions &

recovery

- Mass extinctions (9 Ma) associated with high species loss (86-93%)
- Clades did not recover after suggesting a 'time-forspeciation' recovery

Background speciation and extinction in relation to environmental change





What does the fossil record teach us?

Review

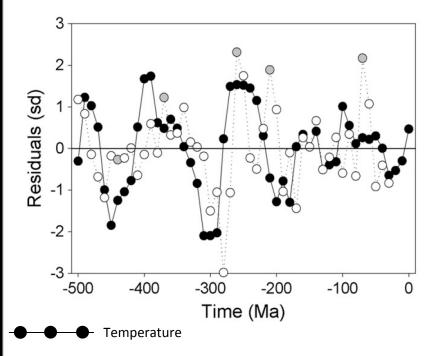


Biodiversity tracks temperature over time

Peter J. Mayhewa, Mark A. Bellb, Timothy G. Bentonc, and Alistair J. McGowanb

*Department of Biology, University of York, York YO10 5DD, United Kingdom; bSchool of Geographical and Earth Sciences, University Glasgow G12 8QQ, United Kingdom; and Faculty of Biological Sciences, University of Leeds, Leeds LS2 9JT, United Kingdom

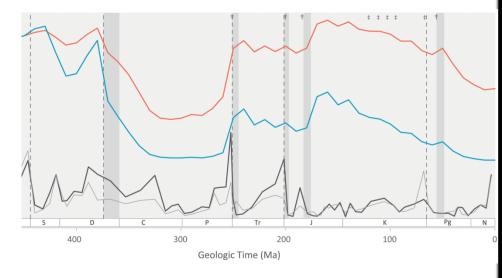
Edited by David Jablonski, The University of Chicago, Chicago, IL, and approved July 30, 2012 (received for review January 18, 2012)



Extinctions in ancient and modern seas

Paul G. Harnik¹, Heike K. Lotze², Sean C. Anderson³, Zoe V. Finkel⁴, Seth Finnegan⁵, David R. Lindberg^{5,6}, Lee Hsiang Liow⁷, Rowan Lockwood⁸, Craig R. McClain¹, Jenny L. McGuire^{1,9}, Aaron O'Dea¹⁰, John M. Pandolfi¹¹, Carl Simpson¹², and

Derek P. Tittensor^{2,13,14}



Temperature

CO₂

Extinction

O- Extinction

O- Speciation

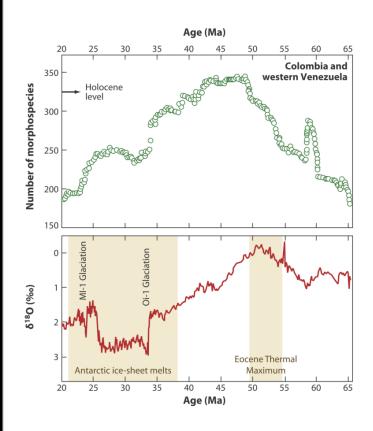
Mayhew et al. 2012 – PNAS

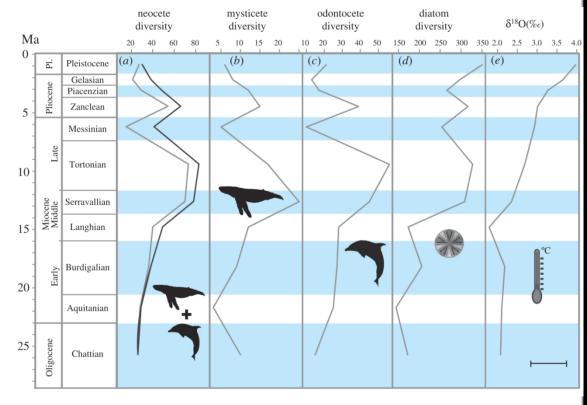
Harnik et al. 2012 – TREE

recovery

What does the fossil record teach us?

"Good correlation between diversity fluctuations and changes in global temperature, suggesting that climate change may be directly driving the diversity pattern."





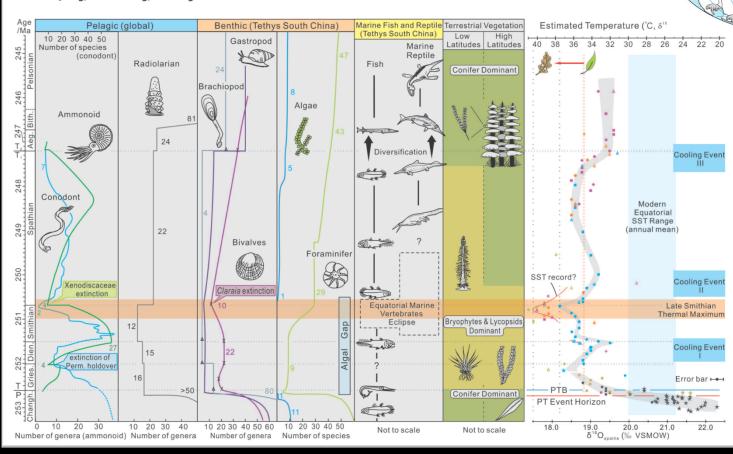
Panthalassa

'DEAD ZONE'

Lethally Hot Temperatures During the Early Triassic Greenhouse

Introduction

Yadong Sun,^{1,2}* Michael M. Joachimski,³ Paul B. Wignall,² Chunbo Yan,¹ Yanlong Chen,⁴ Haishui Jiang,¹ Lina Wang,¹ Xulong Lai¹



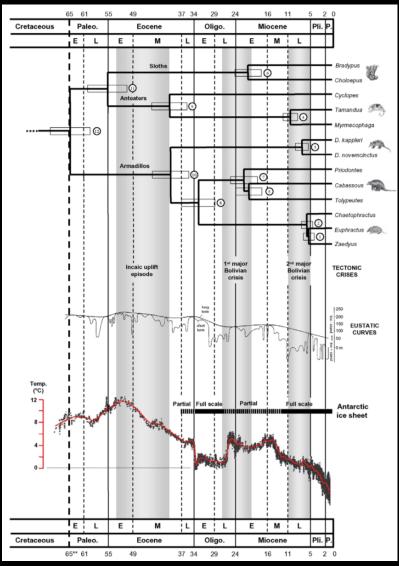
It was simply too hot to survive...

Sun et al. 2012 – Science

Phylogenetic approaches

- Several approaches exist to assess the impact of climate
 - 1. Comparing phylogenies with paleoclimatic curves

- 2. Fitting birth-death model at specific climatic event
- 3. Using trait-dependent diversification models



- ✓ <u>Idea:</u> Visually searched for correlations between phylogenetic events and specific climate changes
- <u>Drawback:</u> highly correlative and speculative approach

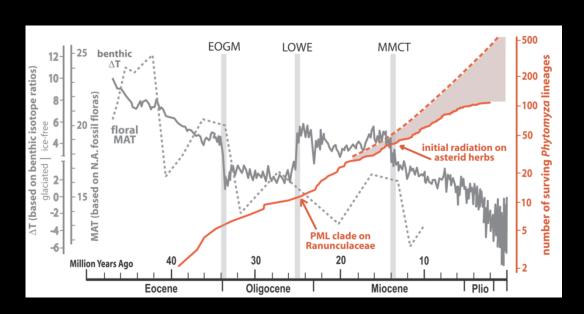
2. Fitting birth-death model at specific climatic event

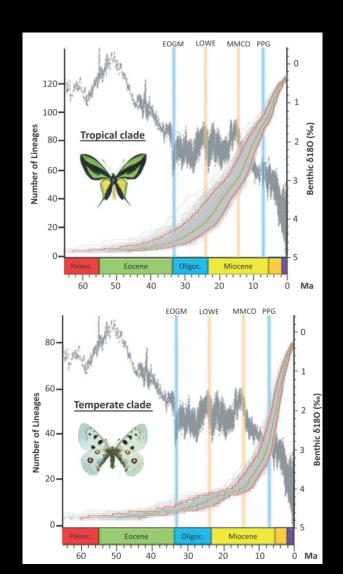
We know **periods of climate changes** in the history a.

Mass extinctions &

recovery

- b. **Compare** a one-rate BD model vs. 2-rates BD model
- **Shift time** is the climatic event (e.g. cooling event)





character

a. Inferring functional dependence of speciation and extinction rates on a

 b. Characters may be either binary (presence/absence) or quantitative (temperature)

BiSSE (Maddison et al. 2007 – Syst. Biol.) QuaSSE (FitzJohn 2010 – Syst. Biol.)

c. Compare several functional dependences (sigmoid, hump-shaped)

JOURNAL OF Evolutionary Biology

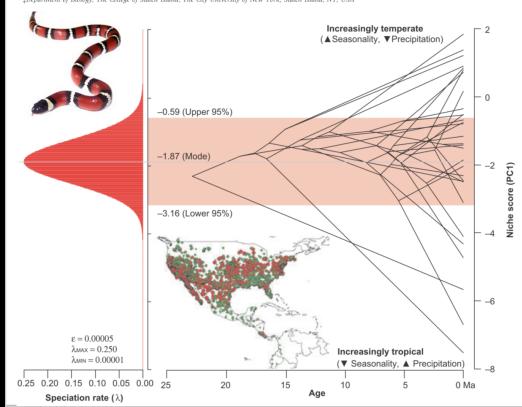


doi: 10.1111/j.1420-9101.2011.02440.x

Trait-dependent diversification and the impact of palaeontological data on evolutionary hypothesis testing in New World ratsnakes (tribe Lampropeltini)

R. A. PYRON* & F. T. BURBRINK†:

- *Department of Biological Sciences, The George Washington University, Washington, DC, USA
- †Department of Biology, The Graduate School and University Center, The City University of New York, New York, NY, USA
- Department of Biology, The College of Staten Island, The City University of New York, Staten Island, NY, USA



Pyron & Burbrink 2012 – J. Evol. Biol.

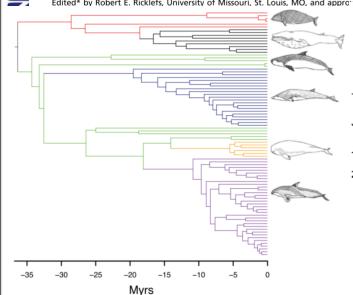
- 1 Approach based on a time-dependent diversification model
- Speciation and extinction rates can vary trough time
- Extinction rate can exceed that of speciation
- Only a fraction of extant species are sampled

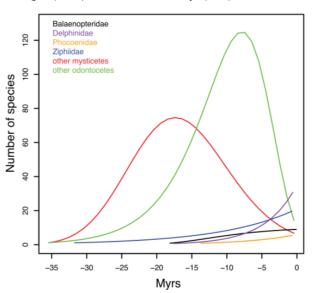
Reconciling molecular phylogenies with the fossil record

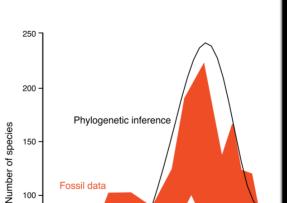
Hélène Morlon^{a,b,1}, Todd L. Parsons^b, and Joshua B. Plotkin^b

^aCenter for Applied Mathematics, Ecole Polytechnique, 91128 Palaiseau, France; and ^bBiology Department, University of Pennsylvania, Philadelphia, PA 19104

Edited* by Robert E. Ricklefs, University of Missouri, St. Louis, MO, and appro: -- ^a. ^ugust 1, 2011 (received for review February 14, 2011)







50

Morlon et al. 2011 – PNAS

- 1 Approach based on a time-dependent diversification model
- Speciation and extinction rates can vary trough time
- Extinction rate can exceed that of speciation
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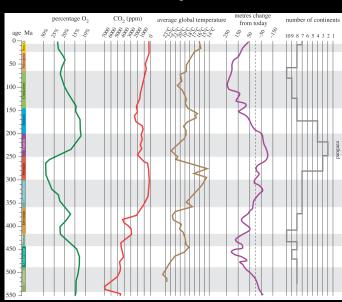
In addition

Morlon et al. 2011 - PNAS

2 – It allows speciation and extinction rates to also depend on an external

variable, itself depending on time:

- Temperature
- Sea level
- Atmospheric carbon concentration
- Number of continents
- ... any external variable you want



Equations of speciation and extinction rates depending on time and environmental variable (E(t))

$$\tilde{\lambda}(t) = \lambda(t, E_1(t), E_2(t), \dots, E_k(t))$$

$$\widetilde{\mu}(t) = \mu(t, E_1(t), E_2(t), \dots, E_k(t))$$

The new exact Likelihood expression of the model with time-varying speciation and extinction rates and incomplete sampling

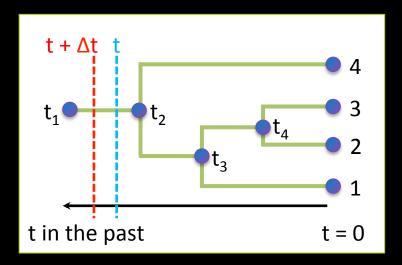
$$L(t_1,...,t_n) = \frac{f^n \Psi(t_2,t_1) \prod_{i=2}^n \tilde{\lambda}(t_i) \Psi(s_{i,1},t_i) \Psi(s_{i,2},t_i)}{1 - \Phi(t_1)}$$

Time is **measured from the present to the past**

recovery

Introduction

- **t = 0** denotes the present, and t increases into the past
- t₁ denotes the first time at which the ancestral species came into existence
- t, is the time of the most recent common ancestor of the sampled species
- $\{t_2, t_3, ..., t_n\}$ denote the times of branching events in the phylogeny, with $t_1 > t_2 > ... > t_n$



$$L(t_1,...,t_n) = \frac{\int_{i=2}^{n} \Psi(t_2,t_1) \prod_{i=2}^{n} \tilde{\lambda}(t_i) \Psi(s_{i,1},t_i) \Psi(s_{i,2},t_i)}{1 - \Phi(t_1)}$$

probability that *n* species are sampled today

$$L(t_1,...,t_n) = \frac{f^n \Psi(t_2,t_1) \prod_{i=2}^n \tilde{\lambda}(t_i) \Psi(s_{i,1},t_i) \Psi(s_{i,2},t_i)}{1 - \Phi(t_1)}$$

 $\Psi(t_2,t_1)$, the probability that a lineage survives from t_1 to t_2 and leaves one descendant lineage at time t_2 :

$$\Psi(t_2,t_1) = e^{\int_{t_2}^{t_1} \tilde{\lambda}(u) - \tilde{\mu}(u) du} \left[1 + \frac{\int_{s}^{t_1} e^{\int_{0}^{\tau} \tilde{\lambda}(\sigma) d\sigma} \tilde{\lambda}(\tau) d\tau}{\frac{1}{f} + \int_{0}^{t_2} e^{\int_{0}^{\tau} \tilde{\lambda}(\sigma) d\sigma} \tilde{\lambda}(\tau) d\tau} \right]^{-2}$$

$$L(t_1,...,t_n) = \frac{f^n \Psi(t_2,t_1) \prod_{i=2}^n \tilde{\lambda}(t_i) \Psi(s_{i,1},t_i) \Psi(s_{i,2},t_i)}{-\Phi(t_1)}$$

probability of a speciation event at time t_i

In which, $\mathbf{s}_{\mathbf{i},\mathbf{1}}$ and $\mathbf{s}_{\mathbf{i},\mathbf{2}}$ denote the times at which the descendant lineages introduced themselves branch at time $t_{\mathbf{i}}$

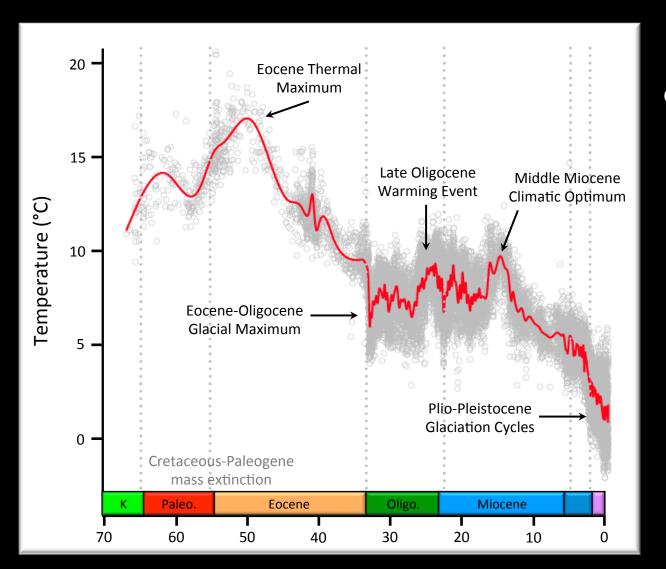
$$L(t_1,...,t_n) = \frac{f^n \Psi(t_2,t_1) \prod_{i=2}^n \tilde{\lambda}(t_i) \Psi(s_{i,1},t_i) \Psi(s_{i,2},t_i)}{1 - \Phi(t_1)}$$

1 - Φ(t₁), the probability that the subclade did not go extinct

 $\Phi(t)$, the probability that a lineage alive at time t has no descendant in the sample:

$$\Phi(t) = 1 - \frac{e^{\int_0^t \widetilde{\lambda}(u) - \widetilde{\mu}(u)du}}{\frac{1}{f} + \int_0^t e^{\int_0^s \widetilde{\lambda}(u) - \widetilde{\mu}(u)du} \widetilde{\lambda}(s)ds}$$

Introduction



Tested with an **exponential dependence** of speciation on temperature given by:

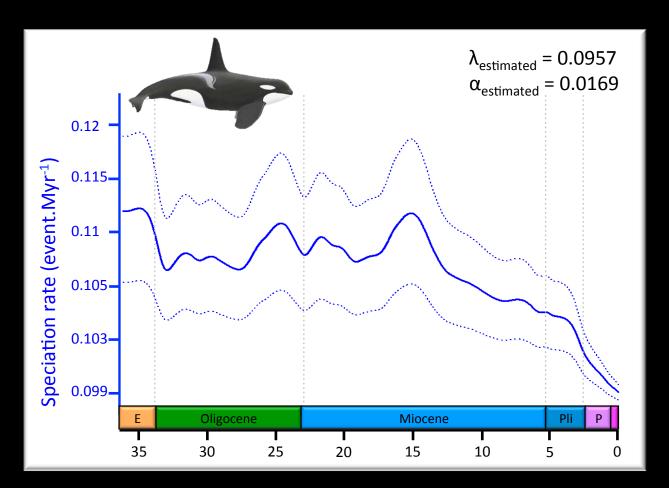
$$\widetilde{\lambda}(t) = \lambda_0 e^{\alpha T(t)}$$

Positive α indicate that higher temperatures enhance speciation

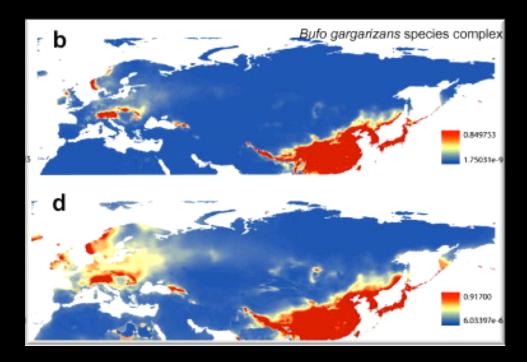
Negative α indicate that higher temperatures hamper speciation

Case study with Cetacea

The relationship between speciation rate and temperature estimated with the approach is $\lambda(T) = 0.0957e^{0.0169T}$, suggesting a positive dependence of speciation rates on temperature



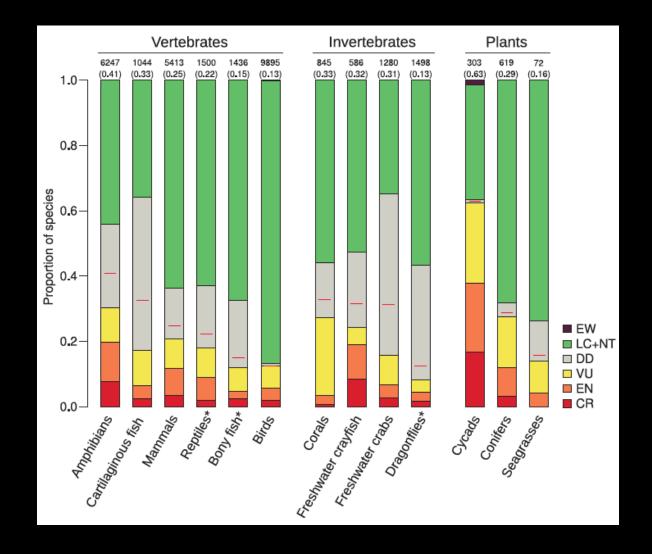
Vulnerability and evolutionary potential



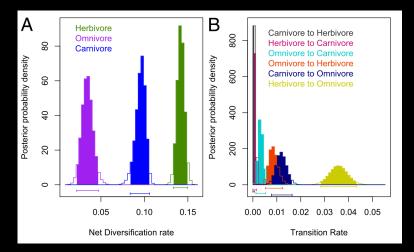


Introduction

Clades are more threatened by extinctions than others

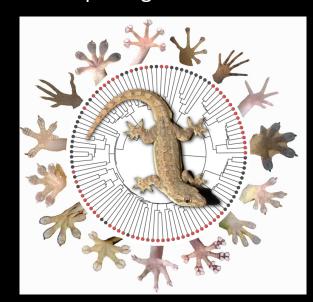


Evaluating vulnerability and evolutionary potential of lineages



Trophic strategy

Morphological character

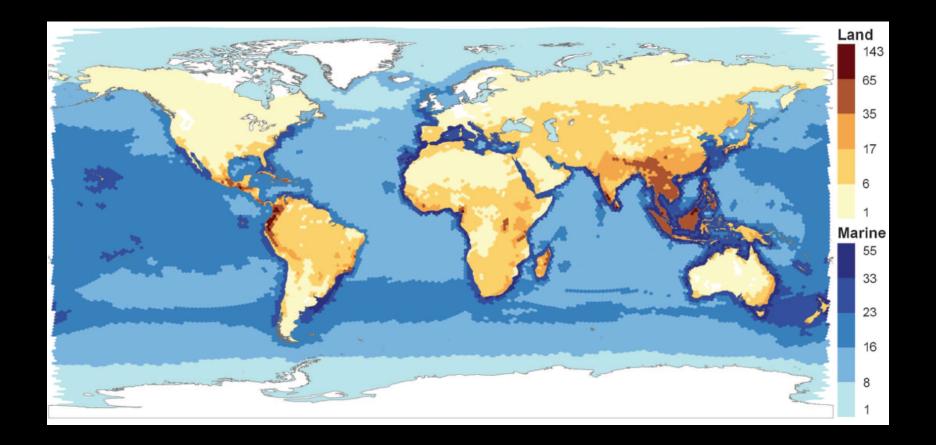


A satio of speciation rates and special spe

Colour polymorphism

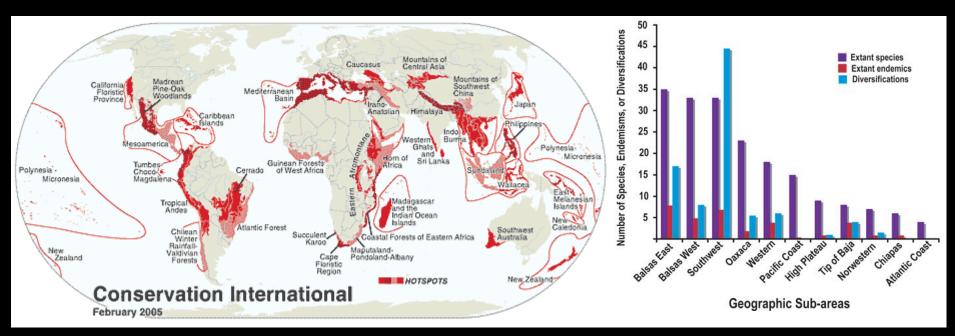
Goldberg et al. 2010 – Science Gamble et al. 2012 – PLoS One Hugall & Stuart-Fox 2012 – Nature Price et al. 2012 – PNAS Introduction

Regions are more threatened by extinctions than others



Conservation policies focus on biodiversity hotspots

... But areas of high diversity are not necessarily areas of high diversification

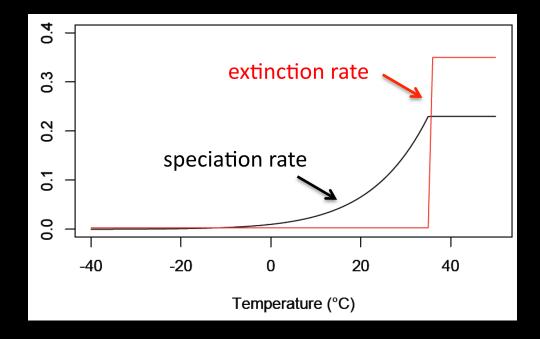


Evaluating vulnerability and evolutionary potential of regions



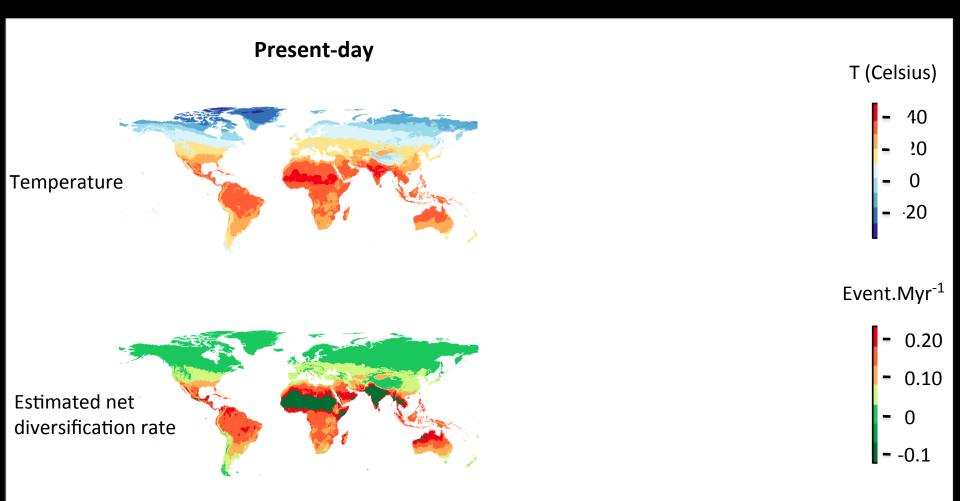
Using phylogenetic approaches (trait-dependent diversification model like QuaSSE)

Possible to estimate the functional dependence of diversification rates on environmental variables:

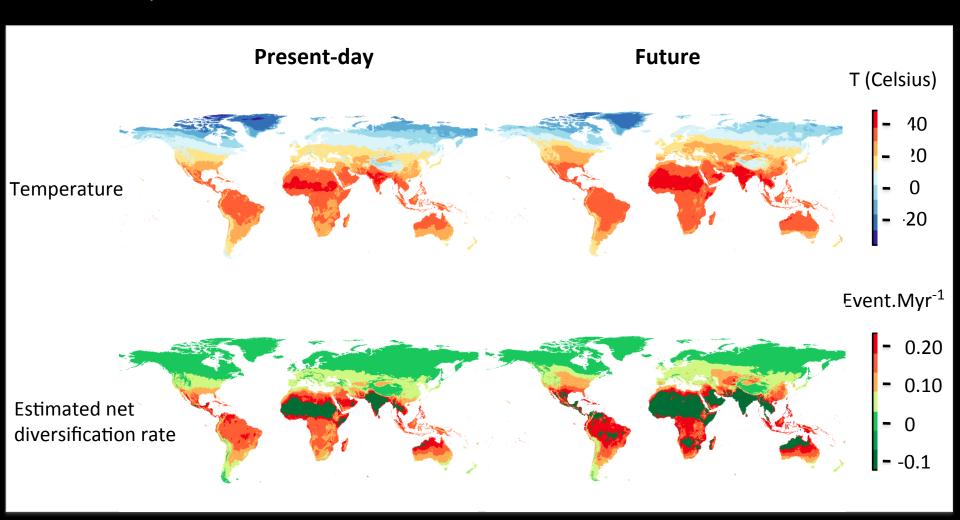


Evaluating vulnerability and evolutionary potential of regions

Map current environmental variables and associated diversification rate

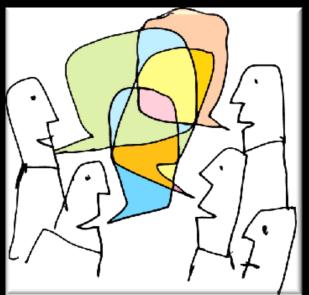


Map future (2080) environmental variables and associated diversification rate





Introduction



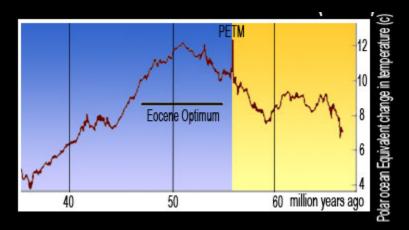


Past versus current environmental changes

- ✓ Keep in mind that what happened in the past is different from what happen today
- ✓ Environmental changes are faster than in the past:

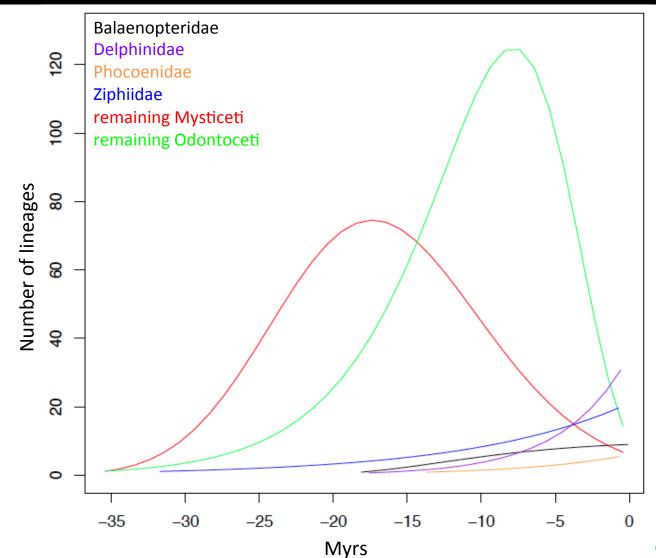
Ex.

- Habitat loss (bolide impacts) were instantaneous => similar to deforestation today?
- Climate changes were usually on long-term period but ... see the PETM at 56 Ma



Global temperatures have increased by 0.0074 °C per year, which is much faster than the 0.0003 °C per year increase within 20 000 years during one of the PETM

Introduction



Comparing current extinction risks to past extinction rates

Clades with negative diversification rates (diversity decline) have high current extinction risk

Clades	Net diversification rates at present		% of threatened	
Balaenopteridae	0.02		25	
Delphinidae	0.224	0.119	13.9	20.4
Phocoenidae	0.141	(±0.085)	42.9	(± 0.18)
Ziphiidae	0.093		0.0	
other mysticetes	-0.528	-0.703	33.3	50
other odontocetes	-0.877	(±0.247)	66.7	(±0.23)
	ı			



Suggest a phylogenetic conservatism of extinction

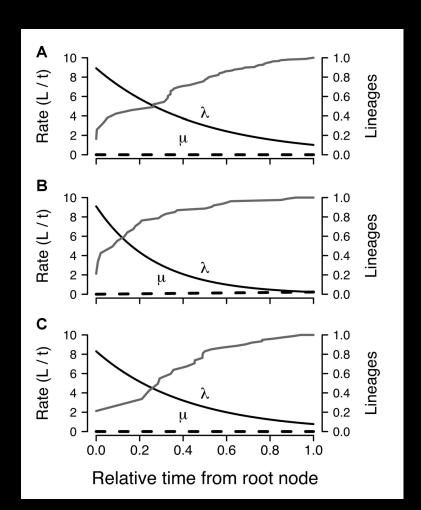


Some biological attributes (body size) that confer risk are conserved

recovery

Estimating extinction rates using phylogenies

For many groups there are low extinction rates which is unrealistic given the fossil record



Introduction

- Integrating phylogenies and the fossil **record** in time-dependent model (Morlon et al. 2011 - PNAS)
- Work in collaboration with Tiago Quental (U. Sao Paulo, Brazil)

Rabosky & Lovette 2008 – Evolution Quental & Marshall 2010 - TREE Didier et al. 2012 – JTB

Conclusion

- 1. Phylogenies can be used to understand past diversity dynamics and how they were influenced by environmental
- 2. Phylogenies provide an additional tool to predict how diversity dynamics may be influenced by environmental change in the future
- 3. The **causes** of current extinctions **may be different** from what happened in the past and they are orders of magnitude higher
- 4. We have **little power to understand the past**, and we will have even **less to predict the future**
- 5. There are many caveats but the **full potential of phylogenies** in global change biology and conservation has **yet to be explored**

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Thanks for your attention