Project summary

Overview

Intellectual merit

Broader impacts

Introduction

Why certain species go extinct while others do not is one of the most fundamental questions in paleobiology. It is expected that for the majority of geological time, extinction is biologically non-random [64, 85, 87, 95, 96, 104, 105]. Determining which biological factors or traits influence extinction risk and how is vital for understanding the differential diversification of life during the Phanerozoic. Periods of background extinction also represent the majority of geologic time, remain relatively predictable and change slowly, and thus providing a better opportunity to study how traits are related to survival than periods of mass extinction [87, 111]. The Law of Constant Extinction [122] posits that extinction risk of taxa within a given adaptive zone is age independent (memoryless), however the generality of this statement is possibly suspect [77, 78, 109, 114]. By analyzing survival patterns within adaptive zones during periods of background extinction, it should be possible to determine if extinction is best modeled as age independent or dependent.

A simple expectation based on purely stochastic grounds, where extinction is not selective, is that abundant and widespread taxa are less likely to go extinct that rare and restricted taxa [?]. For example, species with larger geographic ranges tend to have lower extinction rates than species with smaller geographic ranges [84, 87, 90, 104, 113]. However, this common pattern does not explain why certain taxa may be less prone to extinction than others. In the example, how range size is formed varies between clades and thus remains a black box for most taxa [86] and so determining if differential extinction is a purely stochastic process or is actually the product of selection is impossible. Instead, by focusing on traits related to environmental preference, the traits which may underly why a taxon is abundant or widespread, the process of selection underlying differential extinction may be elucidated.

In addition to understanding patterns of survival, how community composition changes over time is extremely important for understanding how trophic structure changes or is maintained over time. Additionally, community connectedness is important for understanding the degree to which global, regional, or local scale processes are important for shaping the environment, or the set of all possible biotic and abiotic interactors. In addition to total community connectedness, the dynamics of connectedness of taxa within various ecological categories are important for understanding whether different adaptive conditions are differently affected by global, regional, or local scale processes. The Law of Constant extinction is theorized to hold during periods of environmental stability and is thus considered extremely difficult/impossible to test [99]. However, if environmental shifts are incorporated into the analysis of survival distributions, it may be possible to actually test the relationship between taxon age and extinction risk in the context of their adaptive zone

and environment. Additionally, this may allow for illumination of what actual processes underly extinction during the majority of geologic time.

It is under this framework that I propose to study how ecological traits associated with environmental preference have affected both differential survival and cosmopolitan-endemism dynamics. I will be studying two distantly related and biotically different groups: Permian brachiopods and Cenozoic mammals. Both of these groups are considered to have very good fossil records able to reflect long term evolutionary patterns [102]. These two time periods were chosen because they represent periods of approximately the same length (47 My and 65 My) and of climatic change, global warming and global cooling respectively. Also, these two groups are a marine and terrestrial system respectively and the traits associated with environmental preference and range size (described below) are fundamentally very different. Both patterns of survival and community connectedness will be measured for both of these groups. The differences between these two groups in terms of life-habit and environmental preference, along with global climatic context, provides a fantastic scenario to understand how long-term, large-scale processes away from mass extinctions proceed.

Mammals are motile organisms which can track their preferred environmental context over time and space. However, if a taxon requires rare or fragile environmental conditions, or is a poor disperser, this would limit the availability of suitable environments or ability to track the preferred environment. Three important traits that describe the relationship between mammals and their environmental context are body size, dietary category, and locomotor category [74, 75, 94, 100, 101, 117, 118]. Each of these traits describe different aspects of a taxon's adaptive zone such as energetic cost, population density, expected home range size, set of potential prey items, and dispersal ability among others.

Environmental availability, along with stability, is crucial for both the establishment and persistence of a species. During the Cenozoic, primarily between the Paleogene–Neogene, there was a shift from a predominately closed environment to a predominately open environment [70, 92, 112]. This environmental shift was differently timed between continents [119, 120]. Because of the differential timing of environmental shift, along with the different biotic context, the survival and community patterns are expected to vary between continents.

Dietary categories are coarse groupings of similar dietary ecologies: carnivores, herbivores, omnivores, and insectivores. Each of these categories is composed of taxa with a variety of ecologies. For example, herbivores include both browsers and grazers which are known to have had different diversification dynamics during the Cenozoic [91]. Dietary categories are roughly linked with position in trophic hierarchy, with decreasing stability away from the "base." Stability here meaning trophic "distance" from primary productivity, with herbivores having greater stability than carnivores because of the increased likelihood of prey item occurrence. Additionally, with increased likelihood of prey item occurrence, abundance can increase [72, 74, 91, 115, 124] which can effect both survival and increase occupancy [71, 82, 93, 94].

Locomotor categories describe the motility of a taxon, the plausibility of occurrence, and dispersal ability. For example, an obligate arboreal taxon can only occur in locations with a minimum of tree cover and can most likely only disperse to other environments with suitable tree cover. Locomotor categories are similar to dietary categories in that they represent coarse groupings of taxa with similar life habits. Here, the categories are arboreal, ground dwelling, and scansorial. Similar to dietary category, this trait is considered constant at the specific level. Dispersal ability is important

for determining the extent of a taxon's geographic range [69, 83, 89] and affects both the taxon's extinction risk and regional community evenness. With the transition from primarily closed to open environments, there is an expected shift in stability associated with arboreal and ground dwelling taxa.

An organisms body size, here defined as (estimated) mass, has an associated energetic cost in order to maintain homeostasis which in turn necessitates a supply of prey items. Many life history traits are associated with body size: reproductive rate, metabolic rate, home range size, among others [72, 74, 106, 117]. While studies of body size dynamics are very common [67, 73, 75, 95, 97], the interactions or processes that are correlated with body size might better explain the observed diversity pattern more than body size itself. By combining analysis of body size and both dietary and locomotor categories, it should be possible to better understand what processes underly the patterns of survival and community connectedness.

Because dietary category describes, roughly, the trophic position of a taxon and its related stability, it is predicted that more stable categories will have longer durations than less stable categories. Stability here being "distance" from primary productivity, thus it is expected that herbivores will have greater duration than carnivores. Omnivorous taxa are expected to have average taxon durations compared to the other two categories. If dietary category is not found to be important for modeling survival it may mean that trophic category is not a major factor for determining species level survival and that other factors, such as body size, may dominate.

Mammalian herbivores and carnivores have been found to have a greater diversification rate than omnivores [107] which may indicate that these traits are better for survival. However diversification can be caused either by an increase in origination relative to extinction or a decrease in extinction relative to origination. Which scenario occurred, however, is (currently) impossible to determine from a phylogeny of only extant organisms [108] which means that analysis of the fossil record is required. If survival is found to be similar between all dietary categories, this may mean that the differential diversification patterns observed by Price et al. [107] are due to differences in speciation and not extinction.

It is expected that arboreal taxa during the Paleogene will have a greater expected duration than Neogene taxa while the opposite will be true for ground dwelling taxa. In comparison, taxon duration of scansorial taxa is expected to remain relatively similar between the two time periods because it represents a mixed environmental preference that may be viable in either closed or open environments. If locomotor category is not included in the best model of survival this may mean that it is either a poor descriptor of dispersal ability, which may or may not affect mammalian survival. It may also be the case that other factors, measured or unmeasured, may be of greater importance in determining differential survival. The difficulty of a Paleogene–Neogene comparison, which is potentially underminded by heterogeneous preservation, will be explored in simulation.

Body size can possibly scale up to affect species level patterns because, for example, as body size increases, home range size increases [74]. If individual home range size scales up to reflect minimum total species geographic range, we would expect that taxa with larger body sizes would have lower extinction rates than species with smaller body sizes. This expectation, however, may not be right. As body size increases, reproductive rate decreases [95], populations get smaller [125], and generations get longer [103] all of which can increase extinction risk, as has been observed

[76, 97]. However, the relationship between body size and extinction rate at the generic level has been found to vary between continents [97, 121]. By expanding to include a third continent, South America, and analyzing specific level data I hope to elucidate how differences in taxonomic diversity at a continental level might affect body size mediated extinction rate. If body size is found to be unimportant for modeling survival, as in the generic level analysis of Tomiya [121], this means that other biotic or abiotic factors may dominate. This may also mean that individual level home range size does not scale to increased species level range size, and there is therefore no correlated decrease in extinction rate. If increase in body size increases extinction risk, this may be due to traits correlated with body size and not necessarily body size itself [95].

The interaction of body size, locomotor category, and dietary category is also extremely important. For example, a small bodied arboreal taxon of any trophic category during the heavily forested and warm time of the Paleogene would be expected at once to have both a small body size determined range, a large potential geographic range determined by locomotion, as well as an increased availability of resources. Together this would mean that relative survival would be expected to be less than, greater than, and greater than average respectively. Determining which factors dominate during the Paleogene, as well as other parts of the Cenozoic, must be done empirically.

To analyze differential mammalian survival, I propose a survival analysis approach similar to that described above for Permian brachiopods. Mammalian occurrence data will be collected primarily through a combination of the PBDB, Neogene Old World Database (NOW; http://www.helsink.fi/science/now/and museum collections. North American fossil mammal data are well represented in the PBDB because of the extensive work of Alroy [65–67]. European fossil mammal data is also well represented between the PBDB and NOW. South American fossil mammal data is available through the PBDB, but has poor overall coverage. Because of this, South American fossil mammal data will be gathered via various museums such as the Field Museum of Natural History and the American Museum of Natural History as well as published occurrence compilations. With the South American taxa, taxonomy and sampling may not be as well resolved as for North and South America and it may be necessary to restrict analysis to the most taxonomically resolved and sampled groups such as Notoungulata, Marsupials, Carnivora, and Primates.

As described above, duration will be measured as the difference between the observed FAD and LAD of every taxon. Taxa which originated prior to the Cenozoic and all taxa that are either extant or went extinct within 2 My of the present will be censored. This threshold is to limit the effect of the improved record of the Recent.

Dietary category, locomotor category, and body size will be considered constant throughout the duration of a taxon and will be modeled as time-independent covariates of survival. While body size is actually a distribution of values, it is quite common to use a single estimate of mean body size as an aggregate trait in studies of clade-wise dynamics [88]. While all three of these traits may have evolved over a taxon's duration, this will not be considered as part of this study.

While many analyses of survivorship are done using generic data [78, 80, 84, 97, 121], there are potential biases in accurately modeling a specific level process using generic level data [109, 110, 114, 116, 123]. In order to assess some of the differences between generic and specific level survival, I will estimate specific and generic level survival models. Using an approach similar to previous work on estimating specific level origination and extinction rates from generic level survival curves [79], I

will measure the deviance between extinction rate directly estimated from the specific survivorship and the specific level extinction rates estimated from the generic level survival data. In addition to empirical comparison between generic and specific level survival, simulations of diversification with varying levels of cryptic speciation (anagenesis). This may also act as a proxy for generic level diversification because a lineage having a long duration because it is not correctly broken up can be considered analogous to a genus persisting because it continues to speciate.

In order to account for environmental shifts, two different time-dependent covariates will be used. δO^{18} isotope information for the whole Cenozoic [126] will be used as a global climate proxy. Additionally, the Paleogene–Neogene divide, which may reflect global environmental shift, will be modeled as a time-dependent step-function.

Preliminary results and proposed work

Intellectual merit

Broader impacts

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Biographical sketch

Peter D. Smits Committee on Evolutionary Biology, University of Chicago

Professional preparation

University of Washington Biology (Ecology and Evolution) B.S. 2010 Monash University Biological Sciences M.Sc. 2012

University of Chicago Evolutionary Biology Ph.D. 2017 (expected)

Appointments

September 2012–present, Ph.D Candidate, Committee on Evolutionary Biology, University of Chicago.

July 2010–August 2010, Lab Assistant, Vertebrate Paleontology, Burke Museum of Natural History, University of Washington.

July 2009–August 2009, REU Intern, Mammalogy, American Museum of Natural History.

September 2006–September 2008, Lab and Field Assistant, Mammalogy, Burke Museum of Natural History and Culture, University of Washington.

Products

Related publications

• .

Unrelated publications

- .
- .
- .

Synergistic activities

Reviewer for Systematic Biology, PLOS One.

Member of student mental health panel GSMAB, University of Chicago, 2014–present.

Volunteer instructor for peer led programming class, University of Chicago, 2012–2013.

Volunteer expert at "Dino-Day" at the Burke Museum of Natural History and Culture, 2010.

Volunteer expert at "Meet the Mammals" at the Burke Museum of Natural History and Culture, 2007, 2008, 2009.

Collaborators and other affiliations

Collaborators and co-editors

Philip D. Clausen (University of Newcastle), Liliana M. Davalos (Stony Brook University), Alistair R. Evans (Monash University), Richard H. Madden (University of Chicago), Matthew R. McCurry (Monash University), Colin R. McHenry (Monash University), Christopher C. Oldfield (University of Newcastle), P. David Polly (University of Indiana—Bloomington), Michelle R. Quayle (Monash University), Richard H. Ree (Field Museum of Natural History), Heather S. Richards (University of Newcastle), Nancy B. Simmons (American Museum of Natural History), Christopher W. Wamsley (Monash University), Omar M. Warsi (Stony Brook University), Stephen Wroe (University of New England), Paul M. Velazco (American Museum of Natural History).

Graduate advisors and postdoctoral sponsors

Kenneth D. Angielczyk (Field Museum of Natural History), Michael J. Foote (University of Chicago).

Facilities, equipment, & other resources

Major equipment

Calipers, both small (12") and large (24"), are avaliable through the Committee on Evolutionary Biology.

Laboratory

NA

Clinical

NA

Animal

NA

Computer

All data collected from both databases and personal measurements will be stored on a personal computer owned by the Co-PI as well as in both cloud storage and external hard dives. Computer expertise is avaliable at the University of Chicago and the Co-PI has access to large-scale computer clusters via the University of Chicago. All analysis software is free and open-source and is installed on the Co-PI's personal computer and can be installed on all additionally necessary computing clusters.

Office

The PI has dedicated office space at the Field Museum of Natural History. The Co-PI has dedicated office space provided by both the Committee on Evolutionary Biology and the Department of Geophysical Sciences.

Other

NA

Data management plan

The major data and analytical products of the proposed project are 1) anatomical measurements of specimens from museum collections, 2) organized and updated ecological information, and 3) statistical analysis code. All information gathered will be stored indefinitely on both the PI's and Co-PI's personal computers. Additionally, they will be archived on an external hard drive indefinitely in case of the loss of either personal computer.

All anatomical measurements and ecological information used in the proposed study will be provided as supplementary material for all papers produced from this research. These data will also be archived using the data storage service Dryad (http://datadryad.org). Finally, all measurements and ecological information will be available through the Co-PI's personal website (http://home.uchicago.edu/psmits/home.html).

Museums and other institutions where specimens will be measured will be named in all subsequent presentations and papers. These institutions will also be provided with all measurements made to housed specimens, as well as reprints of all related papers.

Anatomical measures, body mass estimates, and updated ecological information will be sent to the Paleobiology Database (http://paleobiodb.org) which is the largest repository of palentological taxonomic, occurrence, and ecological information.

All code used in the proposed analyses will be archived using Dryad, along with the relevant data as discussed above. Code will also be made available through the Co-PI's GitHub page (http://github.com/psmits), a free code sharing and archiving service, as well as through the Co-PI's website (http://home.uchicago.edu/psmits/home.html).

The Co-PI will present the results of the proposed research at conferences and publish said results in peer-reviewed journals in the fields of evolution, paleontology, evolutionary ecology, and global conservation.