

Phylogenetic diversity does not capture body size variation at risk in the world's mammals

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Mammals contribute to important ecosystem processes and services, but many mammalian species are threatened with extinction. We compare how global patterns in three measures of mammalian diversity—species richness, phylogenetic diversity (PD) and body mass variance (BMV)—would change if all currently threatened species were lost. Given that many facets of species' ecology and life history scale predictably with body mass, the BMV in a region roughly reflects the diversity of species' roles within ecosystems and so is a simple proxy for functional diversity (FD). PD is also often considered to be a proxy for FD, but our results suggest that BMV losses within ecoregions would be much more severe than losses of PD or species richness, and that its congruence with the latter two measures is low. Because of the disproportionate loss of large mammals, 65 per cent of ecoregions would lose significantly more BMV than under random extinction, while only 11 per cent would lose significantly more PD. Ecosystem consequences of these selective losses may be profound, especially throughout the tropics, but are not captured by PD. This low surrogacy stresses a need for conservation prioritization based on threatened trait diversity, and for conservation efforts to take an ecosystem perspective.

Keywords: body mass; extinction; functional diversity; mammals; phylogenetic diversity

1. INTRODUCTION

The recently published Global Mammal Assessment emphasized that human actions jeopardize many mammalian species: globally, 1141 species (21% of all mammals) are now threatened with extinction (IUCN 2008). These assessments count each species' loss equally, but species differ in many ways, such as biological traits, ecosystem roles and evolutionary age. Conservation prioritization on the basis of species numbers alone does not capture all aspects of evolutionary history, phylogenetic diversity (PD) or ecosystem roles (Vane-Wright *et al.* 1991; Millennium Ecosystem Assessment 2005). While it is clear that some species are more ecologically important because ecosystem processes depend on their biological traits, the magnitude of functional-trait loss may not be easily predicted from either species or PD loss (Jernvall & Wright 1998; Gross & Cardinale 2005; Hooper *et al.* 2005). Our study therefore aims to map the global loss of evolutionary history as well as changes of the variance in a key species trait, body size, if all currently threatened mammalian species were to go extinct.

Their comparatively large range and body sizes mean that mammals have impacts on ecosystems over large spatial scales. Mammalian seed dispersers, predators and herbivores have been shown to directly or indirectly influence invertebrate and plant community structure, primary productivity and nutrient cycling, suggesting

that species loss would impact ecosystem properties and functioning (Asquith *et al.* 1997; Terborgh *et al.* 2001; Pringle *et al.* 2007; Johnson 2009). Mammals are of special conservation concern because they are charismatic, they provide recreational value and their populations are declining rapidly (Collen *et al.* 2009). They are also of direct economic importance in most areas of the world as sources of food and income from meat, fur and tourism (Milner-Gulland *et al.* 2003).

Large mammals are disproportionately at risk, especially in the tropics (Cardillo *et al.* 2005; Fritz *et al.* 2009). Size or body mass is an important predictor of many ecological traits in mammals, and an indicator of a species' ecological niche (Western 1979; Eisenberg 1981). Species of different sizes generally fulfil different ecosystem functions: for example, the largest mammals are wide-ranging herbivores or carnivores, whereas small mammals are often insectivores or seed dispersers. Preferential extirpation of large species can lead to disproportionate fast loss of functional diversity (FD; Petchey & Gaston 2002). We therefore investigate the impacts of selectivity in the current extinction risk in terms of the variance in log₁₀-transformed body mass (BMV), using ecoregions as spatial units. Given the links between body size and ecology in mammals, the change in BMV for these large-scale ecosystem units can be interpreted as a simple and practicable, if rough, indicator for possible changes in FD (see Mason *et al.* 2003 for a similar index).

Measuring FD accurately within an area is extremely difficult, and comparing FD across different ecosystems is even harder, because functional group definitions or complex ecological distance measures are problematic

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at the global scale and require vast trait databases (Petchey & Gaston 2006). Evolutionary history or PD is sometimes used as a proxy for FD (Faith 1992; Forest *et al.* 2007), because it is an indirect measure of phenotypic diversity if traits evolve along the branches of phylogeny. However, PD and FD of a community can be uncorrelated, so PD loss cannot always reliably predict FD loss (Jernvall & Wright 1998; Hooper *et al.* 2005). Our aim was to compare possible losses of mammalian diversity using three diversity measures: species richness, PD and BMV, where both PD and BMV can be seen as tentative measures of mammalian FD.

Our analyses combined a large dataset of 4230 mammalian species body mass values (Jones *et al.* 2009) with geographical range maps (IUCN 2008) and a species-level phylogeny (Bininda-Emonds *et al.* 2007; Fritz *et al.* 2009), and used phylogenetically explicit methods to account for bias caused by terminal polytomies and by the lack of body-mass estimates for many small species. We compared current levels of diversity with those projected if either all threatened species on the IUCN Red List were to go extinct, or all of these plus species classified as Near Threatened (IUCN 2008). We also simulated random species loss within ecoregions to identify the areas where disproportionately high losses of mammalian PD and BMV would be incurred because of the selectivity of anthropogenic threats.

2. MATERIAL AND METHODS

The spatial units of our analyses, the World Wildlife Fund (WWF) terrestrial ecoregions, are biogeographically coherent areas defined on the basis of natural assemblages (Olson *et al.* 2001). They are widely used in conservation planning (but see Jepson & Whittaker (2002) for a critique). Their large spatial scale ameliorates errors in species distribution maps (Jetz *et al.* 2008), and is more appropriate for investigations of ecosystem functioning than, for example, a fine global grid. We overlaid the ecoregion shapefiles (www.worldwildlife.org/science/ecoregions/item1267.html, accessed August 2006) with mammal range maps from the Global Mammal Assessment (GMA; IUCN 2008), and extracted ecoregion occurrences for each species.

We excluded domesticated species and range parts labelled as 'historical', 'presence uncertain', 'introduced origin' or 'extinct'. GMA maps and risk data were matched to the taxonomy of the body mass data and phylogeny (Wilson & Reeder 2005); hence, 130 species that were absent from this taxonomy were excluded. A total of 56 species ranges were added to the set by splitting existing GMA maps, and 40 species ranges missing from the GMA set were added from Jones *et al.* (2009). Excluding species classified as Data Deficient, Extinct or Extinct in the Wild, our final range dataset contained 4230 species: 2939 of these were ranked as Least Concern, 306 as Near Threatened and 985 species were in one of the three threatened categories (Vulnerable, Endangered and Critically Endangered; IUCN 2008).

The software R (v. 2.7.2) was used for all analyses (R Development Core Team 2008). We used \log_{10} -transformed body mass throughout, because its distribution better approximates the normal distribution; also, the logarithmic transformation reflects the fact that the biological impact of a given difference in body mass (e.g. 10 g) will be much

greater for small species than for large ones. Body mass values for 3382 of the 4230 species (80%) came from the PanTHERIA dataset (Jones *et al.* 2009). As large species are better known, the dataset is biased towards them; we therefore interpolated body mass values for the remaining species as the value of their closest relative (or the mean of several equally close relatives), on the basis of a supertree of 5020 extant mammalian species (Bininda-Emonds *et al.* 2007; Fritz *et al.* 2009). The mean of the resulting frequency distribution of \log_{10} -transformed body mass values was indeed significantly lower than the original mean, confirming the need to interpolate missing data to avoid bias (electronic supplementary material, figure S1; mean \log_{10} -transformed body mass with interpolated values = 2.15; mean without interpolated values = 2.22; *t*-test: $t = -2.64$, d.f. = 7155, $p < 0.01$). We excluded three species-poor ecoregions from all analyses because 50 per cent or more of their body mass values were interpolated. There was strong spatial pattern in the number and proportion of species for which body mass data were interpolated (see the electronic supplementary material, figure S2).

Ecoregion PD was calculated using the total sum of branch lengths in an ecoregion phylogeny (Faith 1992), initially directly derived from the species-level supertree (Bininda-Emonds *et al.* 2007; Fritz *et al.* 2009). PD measured on this tree is inflated because of its polytomies, which mostly reflect the lack of resolution rather than the real speciation pattern; we corrected for this following Davies *et al.* (2008) (see electronic supplementary material for details). We mapped each of the three diversity measures within ecoregions (species richness, PD and BMV) for all species, Least Concern and Near Threatened species (corresponding to the loss of currently threatened species), and just Least Concern species (for the loss of all species whose status is at least Near Threatened).

Congruence of estimated proportional losses was assessed with Pearson correlation coefficients. Because of spatial autocorrelation in our ecoregion dataset, we did not test these for significance: spatial non-independence inflates degrees of freedom in statistical testing, but the coefficients themselves are thought to be unbiased measures of the correlation strength (Legendre 1993). To simulate PD and BMV losses expected if threatened species were a strictly random subset of an ecoregion's species, we used 1000 shuffles of threat status data within each ecoregion; these randomizations therefore preserved the spatial non-randomness in extinction risk prevalence, while simulated risk was random with respect to body mass and phylogeny within each ecoregion. Significance tests were one-tailed (proportion of random values that were smaller than the observed value).

3. RESULTS

Our results suggest that extinction of all but currently Least Concern species would reduce BMV more dramatically than either species richness or PD. Some current 'hotspots' of BMV would completely vanish, while areas of high current species richness or PD would retain comparatively high values (figure 1). Maps of mean body mass within ecoregions suggested that most of the large reduction in variance was owing to selective losses of large species (figure 1*d*). Geographical patterns of losses for all diversity measures were similar whether species classified as Near Threatened were also lost or not (see

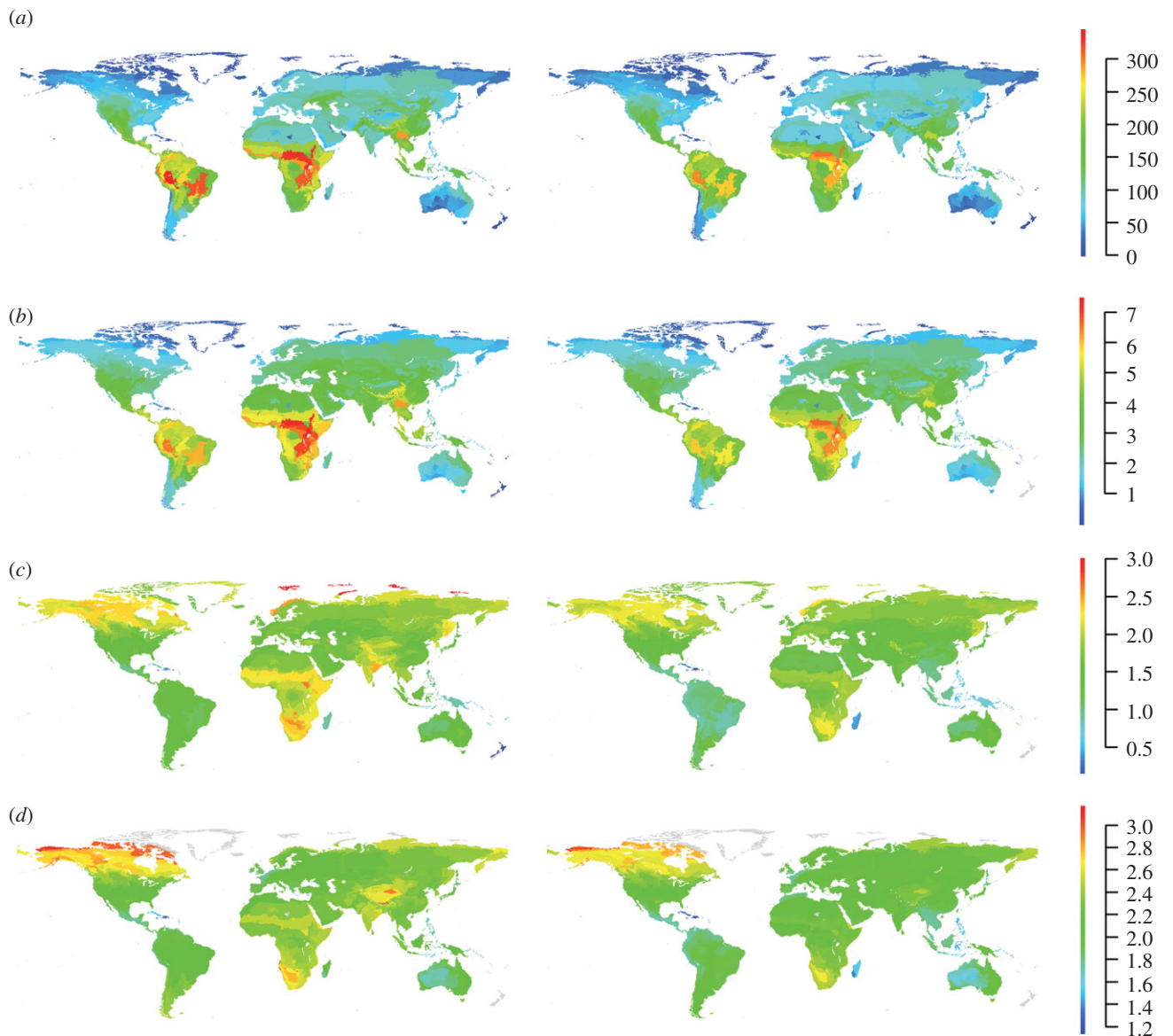


Figure 1. Mammalian diversity within ecoregions for all species (left column) and just species currently rated as Least Concern (right column): (a) species richness, (b) phylogenetic diversity (PD), (c) variance in \log_{10} -transformed body mass (BMV) and (d) mean \log_{10} -transformed body mass. PD (in billions of years) was the total sum of branch lengths for the ecoregion phylogeny modified to account for terminal polytomies (see §2). Mean \log_{10} -transformed body mass within ecoregions is shown only for ecoregions with greater than or equal to 20 species.

the electronic supplementary material, figure S3), so we concentrated on the more severe scenario. Absolute losses for all three diversity measures would be highest in the Indo-Malayan realm, with the Afrotropics and Neotropics also strongly affected (figure 1; table 1). If all currently threatened and Near Threatened species were to go extinct, each ecoregion would lose a global average of 14 species (on an average 15% of its current species), 283 million years of evolutionary history (9%), and 14% of its current body-mass variety (table 1).

The magnitude of proportional losses would differ among the three diversity measures and regionally: they would be high for all three measures in the tropical realms apart from Australasia, but BMV losses would be greater there than losses in richness or PD (figure 2a–c; table 1). Histograms of proportional losses showed that the frequency of larger losses was higher for BMV than for richness and PD (figure 3a–c). In

particular, random species-loss simulations for BMV within ecoregions differed hugely from proportional losses as predicted from Red List risk status, especially in the tropical realms (figure 2c,e).

The often severe reductions in within-ecoregion BMV contrast particularly strongly with the expectation under random species loss (figure 3c,f). Conversely, random species loss led to very similar histograms and maps of PD loss to those projected by losing at-risk species (figures 2b,d and 3b,e). Globally, 65 per cent of ecoregions would lose significantly more BMV than in a random species loss scenario, whereas only 11 per cent of ecoregions would lose significantly more PD (table 1). Much of the northern Holarctic and most tropical areas apart from Australia are predicted to lose significantly more BMV than under random species loss; only some of these areas, most of them tropical, also experienced significantly higher PD losses in our

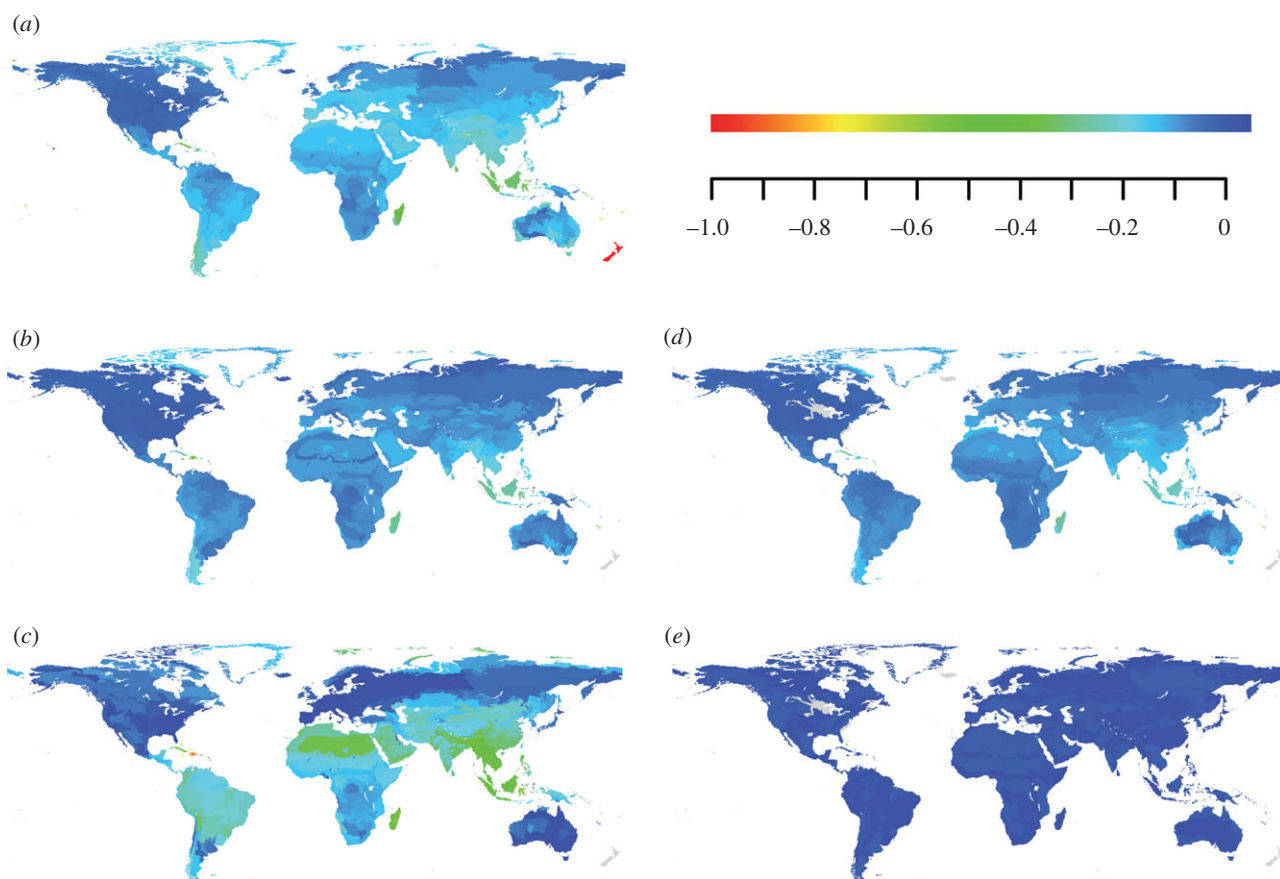


Figure 2. Projected and simulated proportional losses of mammalian diversity within ecoregions: (a–c) projected proportional change if all mammalian species currently ranked as threatened or Near Threatened were lost, for (a) species richness, (b) phylogenetic diversity (PD) and (c) variance in \log_{10} -transformed body mass (BMV). Simulated proportional change for (d) PD and (e) BMV. Negative proportions indicate losses. In (c) and (e), all positive proportions (gains in BMV) were set to dark blue and do not reflect the magnitude of proportional gains. Simulated losses were the mean values of 1000 randomizations for the projected species loss within each ecoregion.

will still impact local ecosystems. Our results provide the first global sketch of how the mammalian trait variety currently present in ecosystems may be reduced in the future, and the emerging picture is not encouraging.

(b) *Selective losses of large mammals*

Reductions in BMV as measured in this study seemed to be mostly owing to the selective loss of large mammals. Large mammals face disproportionate threats, because they are often exploited by humans for meat and fur (Bodmer *et al.* 1997; Fa *et al.* 2002); because their larger home ranges lead to increased exposure to habitat loss and other threats (Woodroffe & Ginsberg 1998); and because life-history traits that increase species' susceptibility to anthropogenic threats scale with body size (Cardillo *et al.* 2005). Our results also agree with previous findings showing that selective loss of large mammalian, bee and beetle species from natural assemblages leads to faster declines in FD than random species losses do (Petchey & Gaston 2002; Larsen *et al.* 2005).

If all but Least Concern species were to disappear, nearly all ecoregions in the large tropical realms (Afrotropic, Neotropic and Indo-Malay) would experience disproportionately large reductions in BMV. These are places where larger species are particularly likely to be declining or at risk of extinction (Collen *et al.* 2009; Fritz *et al.* 2009), apparently because of high recent and ongoing rates of agricultural land conversion (Millennium

Ecosystem Assessment 2005). Historic agricultural impacts have probably already caused local declines and extinctions in temperate regions and Australia in the past, such that non-tropical large mammals tend to be either locally extinct or not perceived as currently under threat (Fritz *et al.* 2009). A recent meta-analysis of mammalian and bird communities also found that previous land conversion was linked to FD being currently lower than expected from species richness (Flynn *et al.* 2009). Our results here imply that boreal forests in North America and most of the Siberian tundra stand to experience strong reductions of mammalian BMV, highlighting the high intrinsic susceptibility of these species-poor but currently relatively pristine areas (Cardillo *et al.* 2006). It seems that human actions have affected large mammals disproportionately at least since the industrial revolution: our figure 1 vividly illustrates how the world might look if these drivers continue unchecked.

(c) *Implications for conservation planning*

Only 11 per cent of ecoregions worldwide stand to lose a significantly higher amount of their current mammalian PD than expected under random species loss. The high redundancy of phylogenetic trees (Nee & May 1997) means that while a global average of 15 per cent of the species in an ecoregion are at risk, representing 14 per cent of current regional BMV on average, only 9 per cent of PD would be lost on average within ecoregions.

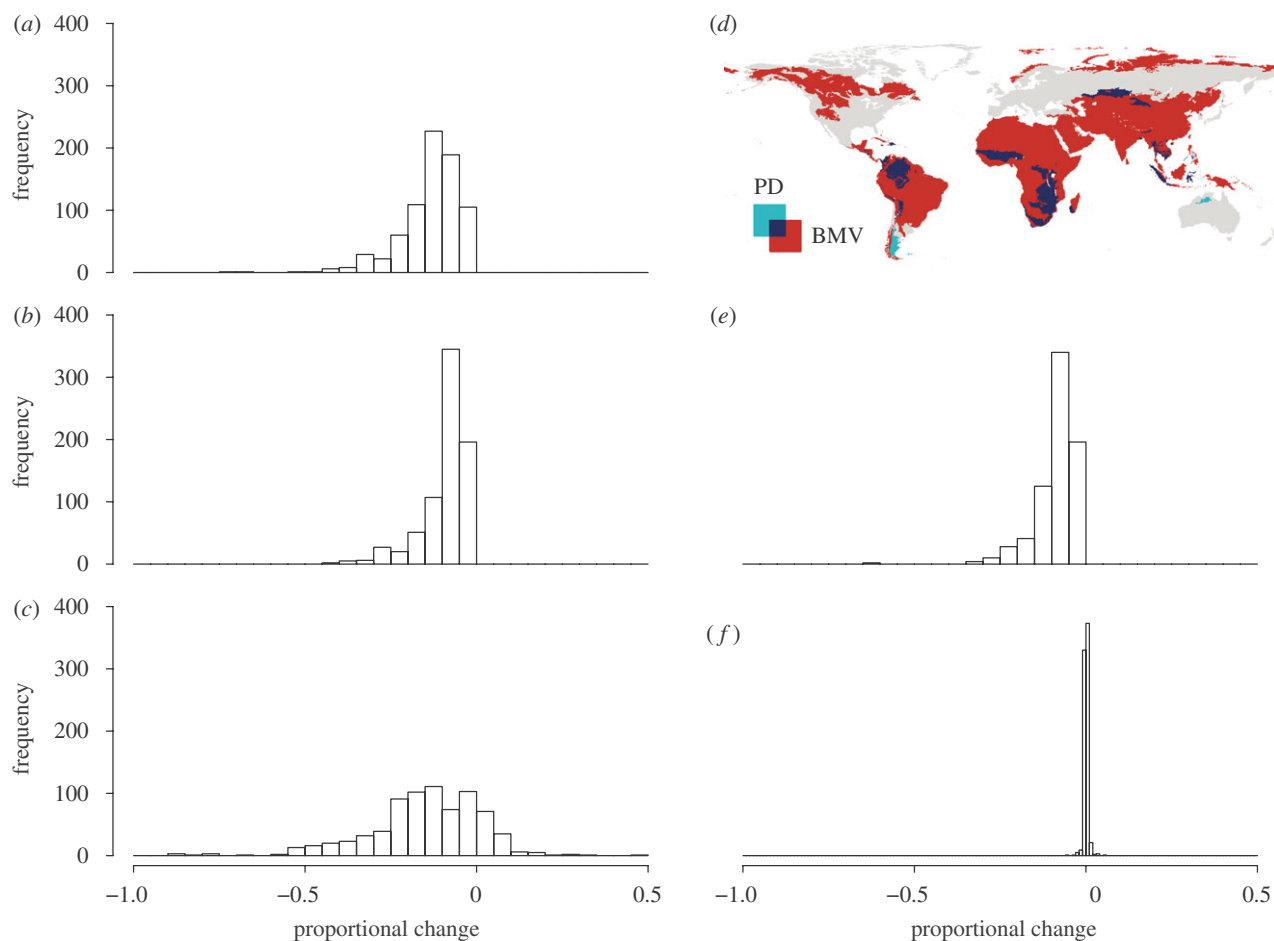


Figure 3. Histograms of (*a–c*) projected and (*e,f*) simulated proportional changes of mammalian diversity within ecoregions, and (*d*) map showing areas projected to lose significantly more-than-random phylogenetic diversity (PD; light blue), variance in \log_{10} -transformed body mass (BMV; red), or both (dark blue). Projected proportional changes were for (*a*) species richness, (*b*) PD and (*c*) BMV; simulated proportional changes were for (*e*) PD and (*f*) BMV. Negative proportions indicate losses; all histograms were scaled to the same axes to facilitate comparison. Simulated losses were the mean values of 1000 randomizations for the projected species loss within each ecoregion; significance of losses within each ecoregion in (*d*) was assessed using a one-tailed test based on the permutations for random species loss.

Congruence with proportional losses in BMV was higher for PD than for species richness, but still relatively low (0.39). Clearly, BMV captures a very different aspect of diversity when compared with PD, and it is unclear which is the better indicator for FD.

PD is still relevant for conservation planning because it acts as a surrogate of diversity for features not correlated with extinction risk (Vane-Wright *et al.* 1991; Faith 1992) and because lost evolutionary heritage is irretrievable. Congruence between predicted losses of species richness and PD was high in this study, which is expected to be a common outcome (Rodrigues *et al.* 2005). This result suggests that ‘classic’ global conservation schemes trying to capture high numbers of threatened species could perform well for the preservation of mammalian evolutionary history, at least at a large spatial scale. However, the low surrogacy between prospective losses of PD and BMV indicates an urgent need for more work on global FD indicators. For conservation planning to consider ecosystem processes and services, we need to understand which species traits underpin them.

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