

Deglaciation explains bat extinction in the Caribbean

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Abstract

Ecological factors such as changing climate on land and interspecific competition have been debated as possible causes of postglacial Caribbean extinction. These hypotheses, however, have not been tested against a null model of climate-driven postglacial area loss. Here, we use a new Quaternary mammal database and deep-sea bathymetry to estimate species–area relationships (SARs) at present and during the Last Glacial Maximum (LGM) for bats of the Caribbean, and to model species loss as a function of area loss from rising sea level. Island area was a significant predictor of species richness in the Bahamas, Greater Antilles, and Lesser Antilles at all time periods, except for the Lesser Antilles during the LGM. Parameters of LGM and current SARs were similar in the Bahamas and Greater Antilles, but not the Lesser Antilles, which had fewer estimated species during the LGM than expected given their size. Estimated postglacial species losses in the Bahamas and Greater Antilles were largely explained by inferred area loss from rising sea level in the Holocene. However, there were more species in the Bahamas at present, and fewer species in the smaller Greater Antilles, than expected given island size and the end-Pleistocene/early Holocene SARs. Poor fossil sampling and ecological factors may explain these departures from the null. Our analyses illustrate the importance of changes in area in explaining patterns of species richness through time and emphasize the role of the SAR as a null hypothesis in explorations of the impact of novel ecological interactions on extinction.

Introduction

The terrestrial mammal fauna of the West Indies once comprised sloths, primates, rodents, insectivores, and bats (Morgan and Woods 1986; Dávalos 2004). During the late Pleistocene and early Holocene waves of extinction nearly obliterated this biota, but the majority of the bats survived (Dávalos and Turvey 2012). Bats were not traditionally hunted for food in the Caribbean, and many species have proven resilient in the face of introduced predators (although see Tejedor et al. 2005). Although habitat changes (Pregill and Olson 1981) and competition (Koopman and Williams 1951; Williams 1952) have been proposed to explain extirpations of Caribbean bats since the Last Glacial Maximum (LGM), sea-level rise caused by nonanthropogenic climate change may be a more important driver of extinction in this fauna (Morgan 2001; Dávalos and Turvey 2012).

The most drastic climatic change since the late Pleistocene was the shift from the conditions of the LGM – from ~22,000 to ~19,000 years before present (yBP; Yokoyama et al. 2000) – to the interglacial climate prevalent since the mid-Holocene. In the terrestrial ecosystems of the West Indies, deglaciation replaced xerophytic environments with mesic habitats (Higuera-Gundy et al. 1998; White et al. 1998; Pajón et al. 2001; McFarlane et al. 2002). One key consequence of climate change was sea-level rise. From 15,000 to 7000 yBP, sea level rose from 100 to 10 m below current level in three bursts marking the collapse of ice sheets, the reorganization of ocean–atmosphere circulation, and the release of glacial meltwater (Blanchon and Shaw 1995). This period corresponds to the inferred last occurrences of many bats, as well as birds and lizards, on many islands (Pregill and Olson 1981; Morgan and Woods 1986; Morgan 1989, 1994, 2001; McFarlane et al. 2002). There are no direct fossil

dates for extinct bat populations, and the 22,000- to 7000-yBP interval corresponding to dramatic rises in sea level overlaps with all indirect radiometric dates for extinct bat populations (Suárez and Díaz-Franco 2003; Jiménez Vázquez et al. 2005). Here, the considerable island area loss caused by deglaciation during the end-Pleistocene/early Holocene serves as an abiotic null hypothesis to explain extinction patterns in the absence of more recent ecological changes, including anthropogenic species introductions, habitat, and climate change.

We combine analyses of bathymetry and estimates of bat species richness across three Caribbean archipelagos to estimate land area and species richness at the LGM (before the end-Pleistocene/early Holocene area loss) and quantify the impact of area declines on bat species richness. The bat biota of the Caribbean is uniquely suited to evaluate the species–area relationship (SAR) across time: the land area experienced significant changes since the LGM, and numerous bat fossils in cave sediments enable reasonable estimates of species richness at the end of the Pleistocene (Fig. 1). In addition, the Caribbean has experienced the highest level of recent species loss of any mammal fauna in the world (MacPhee and Flemming 1999; Morgan 2001; MacPhee 2009; Turvey 2009), so we expect these data will retain considerable power to examine the effects of recent extinction.

Material and Methods

At the LGM, sea levels were 120–135 m below current level (Hearty 1998; Clark et al. 2003). To estimate the area of the islands at the LGM, we decreased sea level by 125 m on the global 1-km grid topography and bathymetry of Becker and Sandwell (2008) in Lambert cylindrical equal-area projection. We investigated the sensitivity of the LGM area estimate for the Bahamas to coral accretion by estimating the effect of a linear growth rate of 1 cm/year over the last 20,000 yBP (Johnson and Pérez 2006). The resulting linear change (200 m) was subtracted from the radius of individual Bahamian banks, and the corresponding areas were recalculated. Current areas were calculated based on current sea level, or compiled from the United Nations Environment Program Earthwatch Database (<http://islands.unep.ch/Tiarea.htm>).

To obtain species richness, we used the extant and extinct mammalian distribution database for the islands of the Caribbean (Willig et al. 2010; Dávalos and Turvey 2012). Species richness at the LGM was calculated as the sum of current and extinct species richness. Stratigraphic and indirect radiometric analyses of fossil faunas including bats have found Late Wisconsinan or Early Holocene dates for the remains (Koopman and Williams 1951; Morgan 2001; McFarlane et al. 2002; Suárez and Díaz-Franco 2003; Mancina and Garcia-Rivera 2005; Steadman



Figure 1. Representative subfossils (Chiroptera: Mormoopidae) from a cave deposit in the Dominican Republic. From left: *Mormoops blainvillei*, *Pteronotus parnellii*, and *P. quadridens*. White bar indicates 1 cm. Quaternary fossils and subfossils on many islands of the West Indies enable estimates of species richness at the Last Glacial Maximum, before sea-level rise drastically reduced the area of most islands.

et al. 2007), indicating most fossil populations would have been extant at the LGM. The ~7000 yBP date for a Cuban fauna of Jiménez Vázquez et al. (2005) coincides with the date at which sea level reached ~10 m below present levels (Blanchon and Shaw 1995). Stratigraphic and radiometric analyses support end-Pleistocene/early Holocene dates for included fossil species, and modern faunal surveys strongly support our designation of species as extinct. The only species in the current fauna thought to have immigrated so recently that it may not have been part of the end-Pleistocene/early Holocene fauna is *Artibeus jamaicensis* (Koopman and Williams 1951; Williams 1952; Morgan 1994), so we estimated SARs with and without this species to assess its effect on results.

Based on biogeographic and geological similarities, we subdivided analyses into three archipelagos: the Bahamas, the Greater Antilles, and the Lesser Antilles (Willig et al. 2010). The fauna of Trinidad, Tobago, Margarita, Aruba, Bonaire, and Curaçao were excluded because these islands are characterized by a South American bat biota (Morgan and Woods 1986; Koopman 1989; Morgan 2001) and are likely subject to fundamentally different biogeographic processes.

To estimate the parameters of the SARs, we fitted separate linear models of species as a function of area for the LGM and the present. The slope of the SAR is expected to become steeper with increasing isolation (MacArthur and Wilson 1967); therefore, higher sea levels since the LGM may have shifted the slope of the current curve relative to the past. Comparisons between the predictions based on the SAR at the LGM and current observations would not be valid if that were the case. To test for homogeneity of slopes (z), we fitted analysis of covariance (ANCOVA) models of species as a function of area (both log-transformed) with LGM or current islands as the factor. These models also tested the homogeneity of the intercept term of SARs – $\log(c)$ – through time.

Since:

$$\log(S_{\text{present}}) = \log(c) + z \log(A_{\text{present}}),$$

and

$$\log(S_{\text{LGM}}) = \log(c) + z \log(A_{\text{LGM}}),$$

assuming c and z remain constant – tested as above – then:

$$\log\left(\frac{S_{\text{present}}}{S_{\text{LGM}}}\right) = z \log\left(\frac{A_{\text{present}}}{A_{\text{LGM}}}\right).$$

Based on this relationship between changes in richness and area, we modeled log-transformed ratios of present/

LGM richness as a function of the ratio of areas without an intercept term.

Finally, we compared the predicted species diversity of each island based on the LGM SAR to the observed current species diversity. If the LGM-based SAR correctly estimated current richness, then islands should fall along a curve of slope = 1 in a plot of predicted versus observed richness. The area below the expected line would indicate underestimated species richness at the LGM and/or more species today than predicted. Conversely, the area above the line would indicate fewer species observed today than expected given the LGM SAR. All analyses were conducted in the R v.1.14.2 statistical environment (R Development Core Team 2010).

Results

Island area was a significant predictor of species richness for all archipelagos and time periods, excluding the Lesser Antilles at the LGM (Table 1, Fig. 2). Species–area curves for the Bahamas and the Greater Antilles had similar slopes for the LGM and present (Table 2). In contrast, the species–area curves fitted for the two time periods for the Lesser Antilles had significantly different slopes, with LGM area explaining a very small portion of the variation in richness at the LGM compared with the present relationship (Tables 1 and 2). We excluded this archipelago from estimates of species loss as a function of area loss, and from comparisons of LGM SARs to present richness because of the heterogeneity of slopes of LGM and current SARs (Table 2).

Island size change since the LGM explained most, but not all, of the decline in species richness on the Bahamas and Greater Antilles (Table 2, Fig. 3). To examine the relationship between LGM and current SARs, we used LGM SARs to predict current species richness from current island area (Fig. 4). If SARs have not changed since the Pleistocene, then LGM SARs should predict observed species richness, and a plot of observed and predicted

Table 1. Slopes and significance of species–area relationships for Caribbean archipelagos.

Archipelago	Period	Slope $\pm$ standard error	R^2	P -value
Bahamas	Last Glacial Maximum (LGM)	0.33 $\pm$ 0.04	0.88	0.0003
	Present	0.24 $\pm$ 0.06	0.40	0.0007
	Present/LGM	0.27 $\pm$ 0.02	0.83	0.0000
Greater Antilles	LGM	0.32 $\pm$ 0.06	0.77	0.0012
	Present	0.28 $\pm$ 0.04	0.69	0.0000
	Present/LGM	0.28 $\pm$ 0.04	0.85	0.0000
Lesser Antilles	LGM	0.08 $\pm$ 0.04	0.15	0.1076
	Present	0.33 $\pm$ 0.07	0.44	0.0003

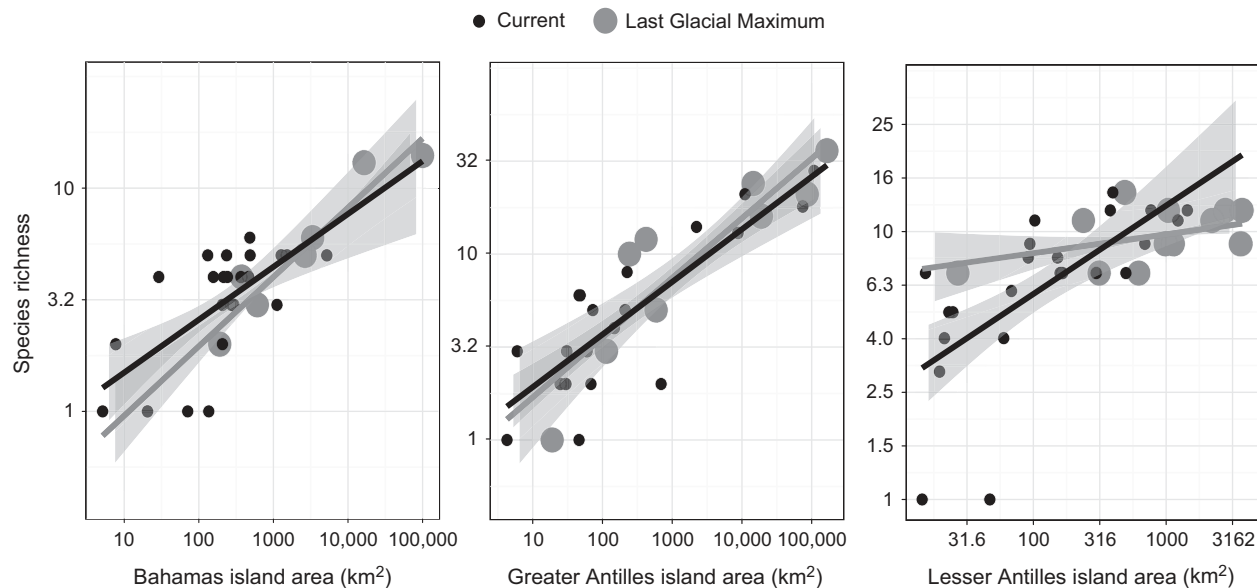


Figure 2. Species–area curves for three Caribbean archipelagos at the Last Glacial Maximum (LGM) and present. Shaded areas indicate the 95% confidence interval around the mean of the curves. LGM species–area relationships (SARs) were highly significant for the Bahamas and the Greater Antilles, but not the Lesser Antilles (Table 1). Current SARs were highly significant for all archipelagos (Table 1). The slopes of the curves fitted for each time period were not statistically different in the Bahamas or Greater Antilles, but were significantly different in the Lesser Antilles (Table 2).

Table 2. Analyses of covariance (ANCOVA) testing for the homogeneity of intercepts and slopes of species–area relationships at present and Last Glacial Maximum.

Archipelago	Time period as factor	P-value	Interaction log area and time period	P-value
Bahamas	0.267 ± 0.300	0.381	-0.074 ± 0.094	0.441
Greater Antilles	0.093 ± 0.243	0.705	-0.038 ± 0.074	0.611
Lesser Antilles	-0.672 ± 0.308	0.037	0.260 ± 0.112	0.027

species richness should show islands roughly falling along an expected line of slope = 1. In the majority of islands in the Bahamas, the LGM SAR predicted fewer species at present than have been observed. The opposite was true for the Greater Antilles, where most of the significant deviations from the expected relationship involved smaller islands with lower-than-expected current species richness.

Species richness on all archipelagos may have changed because of colonization, and island area in the Bahamas may have increased from coral accretion. Widespread species shared with the continent and lacking fossil records are the most likely recent colonizers. Only *Artibeus jamaicensis* meets these criteria: it may be a recent colonizer in the Bahamas. This species was inferred to be present in every island bank of the Greater and Lesser Antilles, so its exclusion

cannot change the slope of those SARs. We conducted analyses accounting for coral accretion and excluding *Artibeus jamaicensis* from the Bahamas (Supporting information). The area difference when accounting for coral deposition in Bahamian banks since the LGM ranged from 0.2% to 5.1% of the estimated LGM area, with a median of 1.3%, and a mean of 2.0%. Over the timespan considered here, colonization by new species has had minimal effect on species richness. Therefore, analyses presented in the main text ignored coral accretion and included *A. jamaicensis* in the LGM Bahamian fauna.

Discussion

We find that island size change is the greatest single predictor of species loss in the Bahamas and Greater Antilles. Although this abiotic change in island area explains most of the observed species loss, there are more species in the Bahamas, and fewer in the smaller Greater Antilles, than expected given current island sizes and predictions from LGM SARs. In the Lesser Antilles, however, there are fewer species known from the LGM than were expected given their size.

Species–area relationships in the Lesser Antilles

Island area was not a significant predictor of species richness at the LGM in the Lesser Antilles (Table 1). This

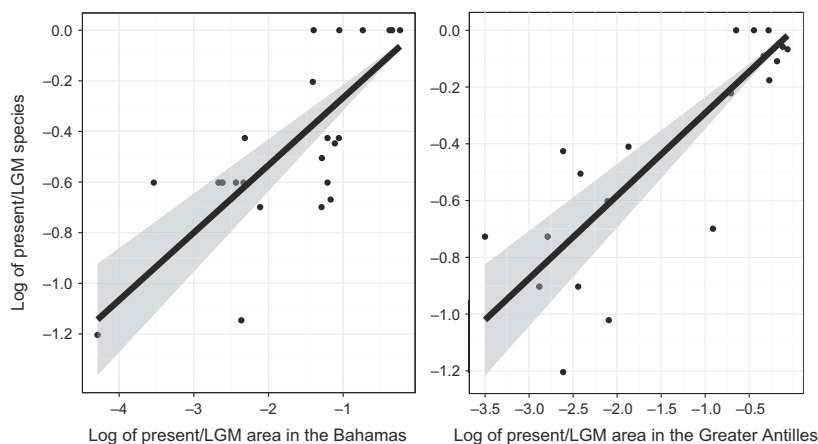


Figure 3. Curves for change in species richness from the Last Glacial Maximum (LGM) to the present as a function of change in area in two Caribbean archipelagos. Shaded areas indicate the 95% confidence interval around the mean of the curves. All relationships were highly significant (Table 1).

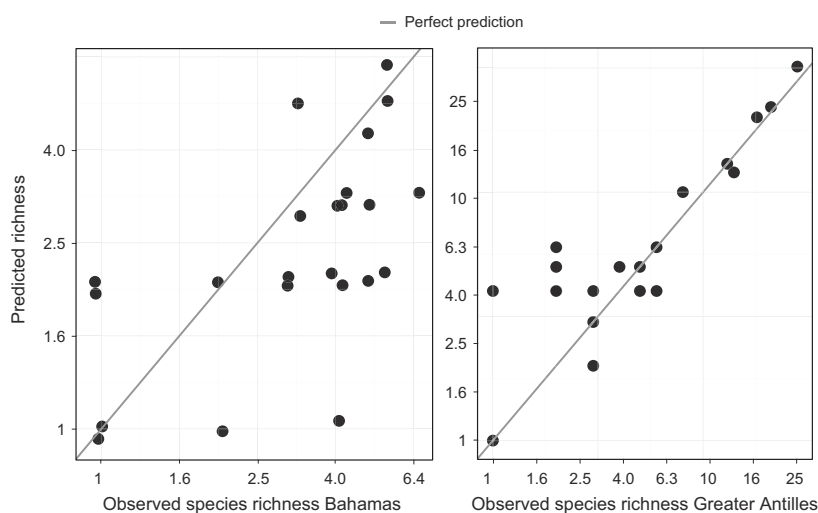


Figure 4. Predicted versus observed species richness in the Bahamas and Greater Antilles. The curve of slope = 1 indicates where the Last Glacial Maximum (LGM) species–area relationships (SAR) perfectly predicts current species richness. The LGM SAR underestimates current species richness in the area below the curve and overestimates current richness in the area above the curve.

result could arise by overestimating the LGM richness of smaller islands that were only recently colonized, or underestimating the richness of larger islands whose fossil records may be incomplete, or both. If the high richness of the smallest island bank (Saba) drove this result, then removing this point would result in a steeper, significant relationship, but it does not (recalculated slope 0.04 ± 0.06 , linear model P -value = 0.5210). Several island banks larger than 1500 km^2 share similar richness estimates of ~ 10 despite differences of hundreds of km^2 in area at the LGM. The expected species richness for these island banks is at least 16 species based on the current curve (Fig. 2). Despite their large size at the LGM, the estimated species richness of these banks is small, and it is likely underestimated because of the scant fossil record of this archipelago. Few fossil sites in the Lesser Antilles have been excavated, and only on Anguilla, and Antigua and Barbuda (these last two islands are part of the same bank; Morgan 2001). The small number of

documented fossil species explains the independence of richness from area in LGM estimates for this archipelago. Our results suggest that more fossil species remain to be discovered from the late Pleistocene/early Holocene of the Lesser Antilles.

Area loss explains most of the change in richness in the Bahamas and Greater Antilles

Five hypotheses other than overhunting and predators introduced by humans have been proposed to explain Caribbean mammal extinction events since the LGM: (1) postglacial sea-level rise reducing island area (Morgan 2001; Dávalos and Turvey 2012); (2) postglacial sea-level rise flooding caves (Morgan 2001); (3) postglacial climate change replacing xerophytic environments with mesic habitats (Pregill and Olson 1981); (4) competition from new colonizers leading to faunal replacement (Koopman

and Williams 1951; Williams 1952), and (5) habitat conversion for human agriculture over the last few thousand years (Gannon et al. 2005). Our estimates of the impact of sea-level change on this biota support the first hypothesis: area loss from postglacial sea-level rise was a major predictor of species loss (Table 1). These results held, even after accounting for sources of error such as coral accretion and the possible recent arrival of *Artibeus jamaicensis* onto the islands (Tables S1 and S2). This model of extinction caused by area loss associated with postglacial sea-level rise has been supported for other Caribbean mammals, such as the giant hutia *Amblyrhiza* in the Sangamonian (McFarlane et al. 1998). We propose extinction caused by area loss as the null hypothesis in investigating insular postglacial extinctions.

In most islands of the Bahamas, LGM SARs predict fewer species at present than are observed. These results could arise through underestimation of species richness at the LGM and suggest that our understanding of the fossil bat biota is incomplete for these banks. A similar analysis of the Greater Antilles showed that SARs for the most species-rich islands in this archipelago are largely unchanged from the LGM (Fig. 4). In smaller islands of the Greater Antilles, however, LGM SARs predict greater species richness than observed. This pattern may be caused by underestimation of current species richness on smaller banks, or because of drivers of richness beyond island area. If current richness at smaller banks were underestimated, then SARs would show a break between smaller and larger areas, with higher slopes at the lower end of the relationship. To evaluate this prediction, we fitted segmented regression models with a single breakpoint for each archipelago (Muggeo 2008), but found no significant breakpoints in the Greater Antillean SAR (P -value = 0.189).

Because underestimation on smaller Greater Antillean banks did not explain the lower-than-expected species richness at present, we suggest that alternative ecological explanations such as the collapse of specific habitats (caves), competition, or habitat loss need to be explored.

By accounting for the major effect of area loss on species declines across most of the Caribbean, and highlighting departures from SAR arising from a poor understanding of the fossil bat fauna in the Lesser Antilles and Bahamas, our analyses illuminate the potential scope of ecological constraints, species interactions, and anthropogenic change on the regional Caribbean fauna.

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Conflict of Interest

None declared.

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Supporting Information

Additional Supporting Information may be found in the online version of this article:

Figure S1. Species-area curves and observed versus predicted richness for the Bahamas at the LGM and present after excluding *Artibeus jamaicensis* and accounting for coral accretion since the LGM. Shaded areas indicate the 95% confidence interval around the mean of the curves. Left: SARs fitted to observed current and estimated LGM values. Right: predicted versus observed species richness. The curve of slope = 1 indicates where the LGM SAR perfectly predicts current species richness. The LGM SAR underestimates current species richness in the area below the curve, and overestimates current richness in the area above the curve.

Table S1. Caribbean bat species inventory by island and archipelago.

Table S2. Slopes and significance of SARs for the Bahamas after excluding *Artibeus jamaicensis* and accounting for coral accretion since the LGM.

Table S3. Analyses of covariance (ANCOVA) testing for the homogeneity of intercepts and slopes of SARs at LGM and present for the Bahamas after excluding *Artibeus jamaicensis* and accounting for coral accretion since the LGM.

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Electronic Supplementary Materials

Supplementary Table S1. Caribbean bat species inventory by island and archipelago.

Supplementary Table S2. Slopes and significance of SARs for the Bahamas after excluding *Artibeus jamaicensis* and accounting for coral accretion since the LGM.

Archipelago	Period	Slope $\pm$ standard error	R^2	P -value
Bahamas	LGM	0.35 ± 0.06	0.82	0.0011
	Present/LGM	0.26 ± 0.02	0.83	0.0000

Supplementary Table S3. Analyses of covariance (ANCOVA) testing for the homogeneity of intercepts and slopes of SARs at LGM and present for the Bahamas after excluding *Artibeus jamaicensis* and accounting for coral accretion since the LGM.

Archipelago	Time period as factor	P -value	Interaction Log Area & Time	P -value
Bahamas	0.412 ± 0.308	0.1923	-0.113 ± 0.097	0.2551

Supplementary Table 1

Species	Family	Endemic to																			Eleuthera Island	Great Exuma	Little Exuma	Long Island	New Providence	Great Inagua	Little Inagua	Grand Bahama
		Extinct in West Indies	West Indies	Acklins	Crooked Island	Fortune Island	East Caicos	Middle Caicos	North Caicos	Providenciales	Andros	Cat Island	Darby															
					Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Bahamas							
<i>Eumops auripendulus</i>	Molossidae																											
<i>Eumops glaucinus</i>	Molossidae																											
<i>Eumops perotis</i>	Molossidae																											
<i>Molossus molossus</i>	Molossidae																											
<i>Mormopterus minutus</i>	Molossidae		yes																									
<i>Nyctinomops laticaudatus</i>	Molossidae																											
<i>Nyctinomops macrotis</i>	Molossidae																											
<i>Tadarida brasiliensis</i>	Molossidae			extant	extant	extant		extinct						extant	extant	extant	extant	extinct										
<i>Mormoops blainvillei</i>	Mormoopidae		yes													extinct		extinct										
<i>Mormoops magna</i>	Mormoopidae	yes	yes																									
<i>Mormoops megalophylla</i>	Mormoopidae	yes									extinct																	
<i>Pteronotus davyi</i>	Mormoopidae																											
<i>Pteronotus macleayii</i>	Mormoopidae		yes																extinct									
<i>Pteronotus parnellii parnellii</i>	Mormoopidae		yes																extinct									
<i>Pteronotus parnellii portoricensis</i>	Mormoopidae		yes																									
<i>Pteronotus parnellii pusillus</i>	Mormoopidae		yes																									
<i>Pteronotus parnellii rubiginosus</i>	Mormoopidae																											
<i>Pteronotus pristinus</i>	Mormoopidae	yes	yes																									
<i>Pteronotus quadridens</i>	Mormoopidae		yes								extinct								extinct									
<i>Pteronotus sp. nov.</i>	Mormoopidae	yes	yes																									
<i>Chilonatalus micropus macer</i>	Natalidae		yes																									
<i>Chilonatalus micropus micropus</i>	Natalidae		yes																									
<i>Chilonatalus tumidifrons</i>	Natalidae		yes									extant	extinct			extinct			extinct									
<i>Natalus jamaicensis</i>	Natalidae		yes																									
<i>Natalus major</i>	Natalidae		yes					extinct																				
<i>Natalus primus</i>	Natalidae	yes	yes									extinct							extinct									
<i>Natalus stramineus</i>	Natalidae		yes																									
<i>Nyctiellus lepidus</i>	Natalidae		yes									extinct	extant		extant	extinct	extant	extant										
<i>Noctilio leporinus</i>	Noctilionidae																			extant								
<i>Ardops nicholli</i>	Phyllostomidae		yes																									
<i>Artibeus flavescens</i>	Phyllostomidae		yes																									
<i>Artibeus anthonyi</i>	Phyllostomidae	yes	yes																									
<i>Artibeus jamaicensis</i>	Phyllostomidae											extant							extant	extant								
<i>Artibeus lituratus</i>	Phyllostomidae																											
<i>Artibeus planirostris</i>	Phyllostomidae																											
<i>Artibeus schwartzi</i>	Phyllostomidae		yes																									
<i>Brachyphylla cavernarum</i>	Phyllostomidae		yes																									
<i>Brachyphylla nana nana</i>	Phyllostomidae		yes										extinct						extinct									
<i>Brachyphylla nana pumila</i>	Phyllostomidae		yes																									
<i>Chiroderma improvisum</i>	Phyllostomidae		yes																									
<i>Cubanyceris silvai</i>	Phyllostomidae	yes	yes																									
<i>Desmodus puntajudensis</i>	Phyllostomidae	yes	yes																									
<i>Erophylla bombifrons</i>	Phyllostomidae		yes																									
<i>Erophylla sezekorni</i>	Phyllostomidae		yes	extant	extant		extant	extant	extant	extant	extant	extant	extant	extant	extant	extant	extant	extant	extant		extant							
<i>Glossophaga longirostris</i>	Phyllostomidae																											
<i>Glossophaga soricina</i>	Phyllostomidae																											
<i>Macrotus waterhousii</i>	Phyllostomidae		yes	extant	extant		extant	extinct	extant	extant	extant	extant	extant	extant	extant	extant	extant	extant	extant									
<i>Monophyllus plethodon</i>	Phyllostomidae		yes																									
<i>Monophyllus redmani</i>	Phyllostomidae		yes	extant	extant			extant	extant	extant	extinct								extinct									
<i>Phyllonycteris aphylla</i>	Phyllostomidae		yes																									
<i>Phyllonycteris major</i>	Phyllostomidae	yes	yes																									
<i>Phyllonycteris poeyi</i>	Phyllostomidae		yes																extinct									
<i>Phyllops falcatus</i>	Phyllostomidae		yes																									
<i>Phyllops silvai</i>	Phyllostomidae	yes	yes																									
<i>Phyllops vetus</i>	Phyllostomidae	yes	yes																									
<i>Stenoderma rufum</i>	Phyllostomidae		yes																									

Supplementary Table 1

Species	Family	Endemic to																			
		Extinct in West Indies	West Indies	Acklins	Crooked Island	Fortune Island	East Caicos	Middle Caicos	North Caicos	Providenciales	Andros	Cat Island	Darby	Eleuthera Island	Great Exuma	Little Exuma	Long Island	New Providence	Great Inagua	Little Inagua	Grand Bahama
Bahamas																					
<i>Sturnira lilium</i>	Phyllostomidae																				
<i>Sturnira thomasi</i>	Phyllostomidae		yes																		
<i>Tonatia saurophila</i>	Phyllostomidae	yes																			
<i>Antrozous pallidus</i>	Vespertilionidae																				
<i>Eptesicus fuscus</i>	Vespertilionidae			extant	extant					extant				extant		extant	extant				extant
<i>Eptesicus guadeloupensis</i>	Vespertilionidae		yes																		
<i>Lasiurus degelidus</i>	Vespertilionidae		yes																		
<i>Lasiurus insularis</i>	Vespertilionidae		yes																		
<i>Lasiurus intermedius</i>	Vespertilionidae		yes																		
<i>Lasiurus minor</i>	Vespertilionidae		yes						extant	extant	extant					extant	extant	extant			extant
<i>Lasiurus pfeifferi</i>	Vespertilionidae		yes																		
<i>Myotis cf. M. austroriparius</i>	Vespertilionidae	yes	yes																		
<i>Myotis dominicensis</i>	Vespertilionidae		yes																		
<i>Myotis martiniquensis</i>	Vespertilionidae		yes																		
<i>Nycticeius cubanus</i>	Vespertilionidae		yes																		
Extant #																					
species				5	5	1	2	3	3	5	5	4	1	4	4	4	6	4	5	1	3
Extinct #																					
species				0	0	0	0	3	0	0	6	1	0	0	2	1	0	10	0	0	0
Total																					
species				5	5	1	2	6	3	5	11	5	1	4	6	5	6	14	5	1	3

Supplementary Table 1

Species	Family	Endemic to																Puerto Rico							
		Extinct in West Indies	West Indies	Great Abaco	Little Abaco	Mayaguana	East Cay	Plana Cay	San Salvador	Cayman Brac	Little Cayman	Cuba	Isle of Pines	Grand Cayman	Hispaniola	Ile de la Gonave	Ile de la Tortue	Jamaica	Mona	Navassa	Anegada	Culebra	Guana	Rico	St. John
				Bahamas	Bahamas	Bahamas	Bahamas	Bahamas		Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles
<i>Eumops auripendulus</i>	Molossidae																	extant							
<i>Eumops glaucinus</i>	Molossidae											extant						extant							
<i>Eumops perotis</i>	Molossidae											extant													
<i>Molossus molossus</i>	Molossidae									extant		extant	extant	extant	extant	extant		extant				extant	extant	extant	extant
<i>Mormopterus minutus</i>	Molossidae		yes									extant													
<i>Nyctinomops laticaudatus</i>	Molossidae											extant													
<i>Nyctinomops macrotis</i>	Molossidae											extant													
<i>Tadarida brasiliensis</i>	Molossidae			extant	extant							extant	extant	extant	extant	extinct		extant						extant	extant
<i>Mormoops blainvillei</i>	Mormoopidae		yes	extinct								extant						extant	extant					extant	
<i>Mormoops magna</i>	Mormoopidae	yes	yes									extinct													
<i>Mormoops megalophylla</i>	Mormoopidae	yes		extinct								extinct						extinct							
<i>Pteronotus davyi</i>	Mormoopidae																								
<i>Pteronotus macleanii</i>	Mormoopidae		yes									extant	extant	extinct				extant							
<i>Pteronotus parnellii parnellii</i>	Mormoopidae		yes	extinct								extant	extinct	extinct				extant							
<i>Pteronotus parnellii portoricensis</i>	Mormoopidae		yes																extant					extant	
<i>Pteronotus parnellii pusillus</i>	Mormoopidae		yes													extant	extinct								
<i>Pteronotus parnellii rubiginosus</i>	Mormoopidae																								
<i>Pteronotus pristinus</i>	Mormoopidae	yes	yes									extinct													
<i>Pteronotus quadridens</i>	Mormoopidae		yes	extinct								extant				extant		extant						extant	
<i>Pteronotus sp. nov.</i>	Mormoopidae	yes	yes													extinct									
<i>Chilonatalus micropus macer</i>	Natalidae		yes									extant	extant	extinct											
<i>Chilonatalus micropus micropus</i>	Natalidae		yes													extant		extant							
<i>Chilonatalus tumidifrons</i>	Natalidae		yes	extant					extant																
<i>Natalus jamaicensis</i>	Natalidae		yes															extant							
<i>Natalus major</i>	Natalidae		yes													extant									
<i>Natalus primus</i>	Natalidae	yes	yes	extinct								extant	extinct	extinct											
<i>Natalus stramineus</i>	Natalidae		yes																						
<i>Nyctiellus lepidus</i>	Natalidae		yes									extant	extant												
<i>Noctilio leporinus</i>	Noctilionidae											extant	extant			extant		extant	extant			extant		extant	extant
<i>Ardops nicholli</i>	Phyllostomidae		yes																						
<i>Artibeus flavescens</i>	Phyllostomidae		yes															extant							
<i>Artibeus thomasi</i>	Phyllostomidae	yes	yes									extinct													
<i>Artibeus jamaicensis</i>	Phyllostomidae					extant				extant	extant	extant	extant	extant	extant	extant		extant		extant	extant	extant	extant	extant	extant
<i>Artibeus lituratus</i>	Phyllostomidae																								
<i>Artibeus planirostris</i>	Phyllostomidae																								
<i>Artibeus schwartzi</i>	Phyllostomidae		yes																						
<i>Brachyphylla cavernarum</i>	Phyllostomidae		yes																				extant	extant	extant
<i>Brachyphylla nana nana</i>	Phyllostomidae		yes							extinct		extant	extant	extant											
<i>Brachyphylla nana pumila</i>	Phyllostomidae		yes							extinct						extant		extinct							
<i>Chiroderma improvisum</i>	Phyllostomidae		yes																						
<i>Cubanycotis silvai</i>	Phyllostomidae	yes	yes									extinct													
<i>Desmodus punctatus</i>	Phyllostomidae	yes	yes									extinct													
<i>Erophylla bombifrons</i>	Phyllostomidae		yes													extant								extant	
<i>Erophylla sezekorni</i>	Phyllostomidae		yes	extant		extant	extant	extant	extant	extant		extant	extant	extant				extant							
<i>Glossophaga longirostris</i>	Phyllostomidae																								
<i>Glossophaga soricina</i>	Phyllostomidae																								
<i>Macrotus waterhousii</i>	Phyllostomidae		yes	extant			extant	extant	extant	extant	extant	extant	extant	extant	extant	extinct		extant		extant				extinct	
<i>Monophyllus plethodon</i>	Phyllostomidae		yes																					extinct	
<i>Monophyllus redmani</i>	Phyllostomidae		yes	extinct						extinct		extant	extant	extinct	extant	extinct		extant						extant	
<i>Phyllonycteris aphylla</i>	Phyllostomidae		yes																						
<i>Phyllonycteris major</i>	Phyllostomidae	yes	yes																					extinct	
<i>Phyllonycteris poeyi</i>	Phyllostomidae		yes	extinct						extinct		extant	extant	extant	extant	extant									
<i>Phyllops falcatus</i>	Phyllostomidae		yes																						
<i>Phyllops silvai</i>	Phyllostomidae	yes	yes																						
<i>Phyllops vetus</i>	Phyllostomidae	yes	yes																						
<i>Stenoderma rufum</i>	Phyllostomidae		yes																					extant	extant

Supplementary Table 1

Species	Family	Endemic to																Puerto Rico							
		Extinct in West Indies	West Indies	Great Abaco	Little Abaco	Mayaguana	East Cay	Plana Cay	San Salvador	Cayman Brac	Little Cayman	Cuba	Isle of Pines	Grand Cayman	Hispaniola	Ile de la Gonave	Ile de la Tortue	Jamaica	Mona	Navassa	Anegada	Culebra	Guana	Rico	St. John
				Bahamas	Bahamas	Bahamas	Bahamas	Bahamas	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles
<i>Sturnira lilium</i>	Phyllostomidae		yes																						
<i>Sturnira thomasi</i>	Phyllostomidae	yes																extinct							
<i>Tonatia saurophila</i>	Phyllostomidae																								
<i>Antrozous pallidus</i>	Vespertilionidae											extant													
<i>Eptesicus fuscus</i>	Vespertilionidae			extant					extant	extant			extant	extant	extant			extant						extant	
<i>Eptesicus guadeloupensis</i>	Vespertilionidae		yes																						
<i>Lasiurus degelidus</i>	Vespertilionidae		yes															extant							
<i>Lasiurus insularis</i>	Vespertilionidae		yes										extant												
<i>Lasiurus intermedius</i>	Vespertilionidae		yes										extant	extant		extinct									
<i>Lasiurus minor</i>	Vespertilionidae		yes			extant										extant							extant		
<i>Lasiurus pfeifferi</i>	Vespertilionidae		yes										extant												
<i>Myotis cf. M. austroriparius</i>	Vespertilionidae	yes	yes	extinct																					
<i>Myotis dominicensis</i>	Vespertilionidae		yes																						
<i>Myotis martiniquensis</i>	Vespertilionidae		yes																						
<i>Nycticeius cubanus</i>	Vespertilionidae		yes									extant													
Extant #																									
species				5	1	3	2	4	6	2	28	14	8	18	2	0	21	3	1	1	3	3	13	6	
Extinct #																									
species				8	0	0	0	0	4	0	8	3	4	3	4	0	3	0	0	0	0	0	3	0	
Total																									
species				13	1	3	2	4	10	2	36	17	12	21	6	0	24	3	1	1	3	3	16	6	

Supplementary Table 1

Species	Family	Endemic to																					
		Extinct in West Indies	West Indies	St. Thomas	Tortola	Vieques	Virgin Gorda	St. Croix	Anguilla	St. Barthelemy	St. Eustatius	St. Martin	Tintamarre	Antigua	Barbuda	Barbados	Dominica	Bequia	Carriacou	Mustique	Union	Guadeloupe	La Desirade
				Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Greater Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles	Lesser Antilles
<i>Eumops auripendulus</i>	Molossidae																						
<i>Eumops glaucinus</i>	Molossidae																						
<i>Eumops perotis</i>	Molossidae																						
<i>Molossus molossus</i>	Molossidae			extant	extant	extant	extant	extant	extant	extant	extant	extant		extant	extant	extant	extant		extant		extant	extant	extant
<i>Mormopterus minutus</i>	Molossidae		yes																				
<i>Nyctinomops laticaudatus</i>	Molossidae																						
<i>Nyctinomops macrotis</i>	Molossidae																						
<i>Tadarida brasiliensis</i>	Molossidae								extant	extant	extant	extant		extant	extant		extant				extant	extant	
<i>Mormoops blainvillei</i>	Mormoopidae		yes						extinct					extinct	extinct								
<i>Mormoops magna</i>	Mormoopidae	yes	yes																				
<i>Mormoops megalophylla</i>	Mormoopidae	yes																					
<i>Pteronotus davyi</i>	Mormoopidae																extant						
<i>Pteronotus macleanii</i>	Mormoopidae		yes																				
<i>Pteronotus parnellii parnellii</i>	Mormoopidae		yes																				
<i>Pteronotus parnellii portoricensis</i>	Mormoopidae		yes																				
<i>Pteronotus parnellii pusillus</i>	Mormoopidae		yes											extinct									
<i>Pteronotus parnellii rubiginosus</i>	Mormoopidae																						
<i>Pteronotus pristinus</i>	Mormoopidae	yes	yes																				
<i>Pteronotus quadridens</i>	Mormoopidae		yes																				
<i>Pteronotus sp. nov.</i>	Mormoopidae	yes	yes																				
<i>Chilonatalus micropus macer</i>	Natalidae		yes																				
<i>Chilonatalus micropus micropus</i>	Natalidae		yes																				
<i>Chilonatalus tumidifrons</i>	Natalidae		yes																				
<i>Natalus jamaicensis</i>	Natalidae		yes																				
<i>Natalus major</i>	Natalidae		yes																				
<i>Natalus primus</i>	Natalidae	yes	yes																				
<i>Natalus stramineus</i>	Natalidae		yes						extant				extant		extant	extant		extant				extant	
<i>Nyctiellus lepidus</i>	Natalidae		yes																				
<i>Noctilio leporinus</i>	Noctilionidae			extant		extant		extant					extant		extant	extant	extant	extant		extant		extant	
<i>Ardops nicholli</i>	Phyllostomidae		yes										extant		extant		extant					extant	
<i>Arctus flavescens</i>	Phyllostomidae		yes																			extant	
<i>Artibeus thomasi</i>	Phyllostomidae	yes	yes																				
<i>Artibeus jamaicensis</i>	Phyllostomidae			extant	extant	extant	extant	extant	extant	extant	extant	extant		extant	extant	extant	extant	extant	extant	extant	extant	extant	extant
<i>Artibeus lituratus</i>	Phyllostomidae																						
<i>Artibeus planirostris</i>	Phyllostomidae																						
<i>Artibeus schwartzi</i>	Phyllostomidae		yes																				
<i>Brachyphylla cavernarum</i>	Phyllostomidae		yes	extant				extant	extant	extant	extant	extant		extant	extant	extant	extant				extant	extant	
<i>Brachyphylla nana nana</i>	Phyllostomidae		yes																				
<i>Brachyphylla nana pumila</i>	Phyllostomidae		yes																				
<i>Chiroderma improvisum</i>	Phyllostomidae		yes																			extant	
<i>Cubanactes silvai</i>	Phyllostomidae	yes	yes																				
<i>Desmodus puntajudensis</i>	Phyllostomidae	yes	yes																				
<i>Erophylla bombifrons</i>	Phyllostomidae		yes																				
<i>Erophylla sezekorni</i>	Phyllostomidae		yes																				
<i>Glossophaga longirostris</i>	Phyllostomidae																						
<i>Glossophaga soricina</i>	Phyllostomidae																						
<i>Macrotus waterhousii</i>	Phyllostomidae		yes						extinct														
<i>Monophyllus plethodon</i>	Phyllostomidae		yes						extant	extant												extant	
<i>Monophyllus redmani</i>	Phyllostomidae		yes																				
<i>Phyllonycteris aphylla</i>	Phyllostomidae		yes																				
<i>Phyllonycteris major</i>	Phyllostomidae	yes	yes																				
<i>Phyllonycteris poeyi</i>	Phyllostomidae		yes																				
<i>Phyllops falcatus</i>	Phyllostomidae		yes																				
<i>Phyllops silvai</i>	Phyllostomidae	yes	yes																				
<i>Phyllops vetus</i>	Phyllostomidae	yes	yes																				
<i>Stenoderma rufum</i>	Phyllostomidae		yes	extant		extant		extant															

Supplementary Table 1

Species	Family	Extinct in West Indies	Endemic to West Indies	St. Thomas	Tortola	Vieques	Virgin Gorda	St. Croix	Anguilla	St. Barthelemy	St. Eustatius	St. Martin	Tintamarre	Antigua	Barbuda	Barbados	Dominica	Bequia	Carriacou	Mustique	Union	Guadeloupe	La Desirade

Supplementary Figure S4. Species-area curves and observed vs. predicted richness for the Bahamas at the LGM and present after excluding *Artibeus jamaicensis* and accounting for coral accretion since the LGM. Shaded areas indicate the 95% confidence interval around the mean of the curves. Left: SARs fitted to observed current and estimated LGM values. Right: predicted vs. observed species richness. The curve of slope = 1 indicates where the LGM SAR perfectly predicts current species richness. The LGM SAR underestimates current species richness in the area below the curve, and overestimates current richness in the area above the curve.

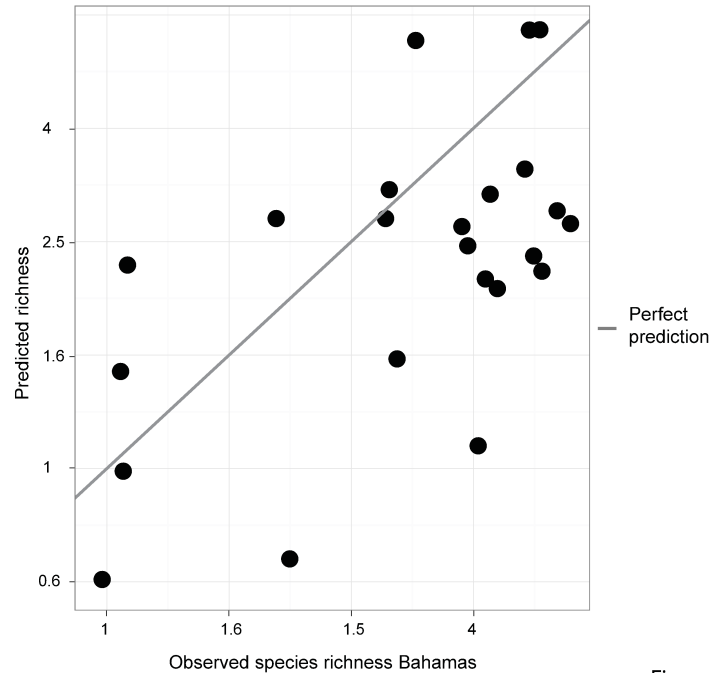
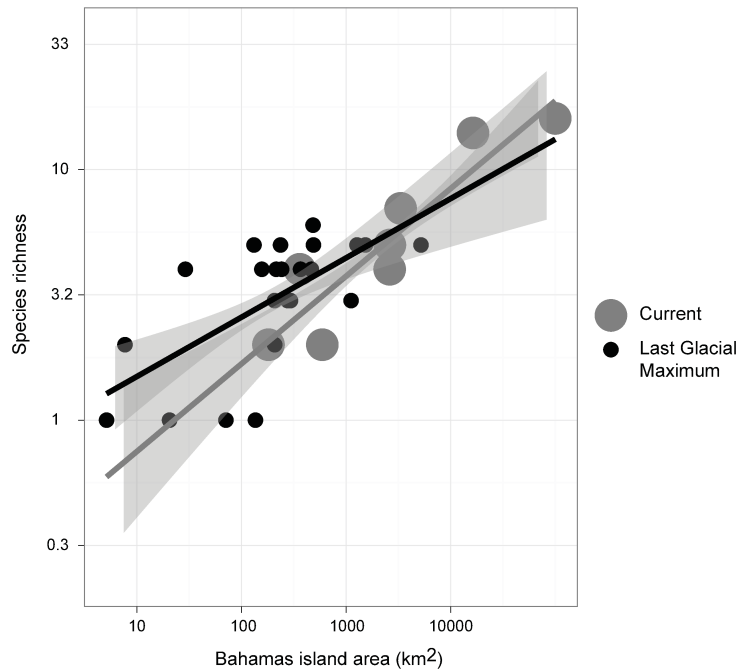


Figure S4