1Small mammal responses to Amazonian forest islands are 2modulated by their forest dependence

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16We assessed the extinction dynamics of small mammals in man-made Amazonian forest 17 islands, providing unprecedented information that can be used in areas affected/to be 18 affected, by hydroelectric dams.

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20Author Contributions: AFP, MVV and CAP conceived and designed the experiments. AFP 21performed the experiments, and MB provided additional data. AFP analysed the data and 22wrote the manuscript under the supervision and advice of MVV and CAP; MB also 23provided editorial advice.

24Abstract

25Hydroelectric dams have induced widespread loss, fragmentation and degradation of 26terrestrial habitats in lowland tropical forests. Yet their ecological impacts have been 27widely neglected, particularly in developing countries, which are currently earmarked for 28exponential hydropower development. Here we assess small mammal assemblage 29responses to Amazonian forest habitat insularization induced by the 28-year old Balbina 30Hydroelectric Dam. We sampled small mammals on 25 forest islands (0.83 - 1,466 ha) and 31 four continuous forest sites in the mainland to assess the overall community structure and 32species-specific responses to forest insularization. We classified all species according to 33their degree of forest-dependency using a multi-scale approach, considering landscape, 34patch and local habitat characteristics. Based on 65,520 trap-nights, we recorded 884 35individuals of at least 22 small mammal species. Species richness was best predicted by 36island area and isolation, with small islands (<15 ha) harbouring an impoverished nested 37 subset of species (mean \pm SD: 2.6 \pm 1.3 species), whereas large islands (>200 ha; 10.8 \pm 381.3 species) and continuous forest sites (∞ ha; 12.5 \pm 2.5 species) exhibited similarly high 39species richness. Forest-dependent species showed higher local extinction rates and were 40often either absent or persisted at low abundances on small islands, where non-forest-41dependent species became hyper-abundant. Species capacity to use non-forest habitat 42matrices appears to dictate small mammal success in small isolated islands. We suggest 43that ecosystem functioning may be highly disrupted on small islands, which account for 4462.7% of all 3,546 islands in the Balbina Reservoir.

45**Key words:** habitat fragmentation, hydroelectric dams, island biogeography, land-bridge 46islands, tropical forests.

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48Introduction

49Hydroelectric dams have become major drivers of habitat loss and fragmentation 50worldwide (Lees et al. 2016; Jones et al. 2016). Still, their ecological and social impacts 51have been widely neglected, particularly in hyper-diverse tropical developing countries 52(Castello et al. 2013; Lees et al. 2016), which are currently the primary targets for 53hydropower development (Zarf et al. 2015). By flooding low-elevation areas, river 54damming often converts previous ridgetops into land-bridge islands, creating a complex 55insular landscape within hydroelectric reservoirs. A total of 145 dams in operation or under 56construction have flooded or will flood ~1.5 Mha of pristine forests in the Amazon basin 57alone, and current government plans include the construction of 263 additional dams 58(ECOA 2016).

Biological communities isolated on land-bridge islands are likely to be affected by 60several multi-scale drivers – including landscape, patch and habitat quality features of any 61given site – which may be more evident in habitat fragments than in true islands (Arroyo-62Rodriguez et al. 2013). At the landscape scale, the degree of isolation from other islands 63and the mainland affects species colonization rates (MacArthur and Wilson 1967; Watling 64and Donnelly 2006). At the patch level, the remaining habitat area affects both population 65sizes and (micro)habitat diversity (Hutchinson 1957; MacArthur and Wilson 1967), and the 66severity of edge effects that can penetrate deeply into small islands (Benchimol and Peres 672015a). Edge effects are further linked to local habitat quality, which determines the 68spectrum of ecological niches available and, consequently, local species diversity (August 691983).

Species persisting in insular habitat patches are further prone to be affected by 71intrinsic eco-morphological traits (Cosson et al. 1999; Lynam and Billick 1999). In 72addition to dispersal ability, traits associated with success in disturbed habitat remnants 73play important roles in structuring animal communities isolated within land-bridge islands

74(Lynam and Billich 1999). Thus, classifying species according to their ecological traits can 75help us to understand how spatial and habitat characteristics affect species persistence 76(Ewers and Didham 2006; Santos-Filho et al. 2016), and how species composition changes 77across disturbance gradients (Pardini et al. 2009). Forest-dependent species unable to 78transverse unsuitable habitat between forest patches are therefore considered more 79extinction-prone in fragmented landscapes (Henle et al. 2004), whereas more structurally 80complex islands are expected to boost the persistence of forest-specialist species (Devictor 81et al. 2008). This pattern of selective extinctions may further result in a nested structure 82along the gradient of fragmentation, with species persisting at progressively smaller islands 83comprising subsets of assemblages retained at larger sites (Wright et al. 1998).

84 In the Neotropics, small non-volant mammals (marsupials and rodents) occupy a 85central position in forest food webs, acting either as seed predators and dispersers (Mangan 86and Adler 2000; Terborgh et al. 2001), pollinators (Vieira et al. 1991), arthropod predators 87(Carvalho et al. 2005), and as a resource for higher trophic levels (Wright et al. 1994). 88Although detailed ecological information on most species is scant, small mammals play 89critical roles in ecosystem functioning including forest regeneration (Terborgh et al. 2001; 90Galetti et al. 2015). Several studies have assessed the effects of habitat fragmentation on 91small mammals, including tropical reservoir islands (Adler and Seamon 1996; Granjon et 92al. 2002; Lambert et al. 2003; Wang et al. 2010a, Gibson et al. 2013). Indeed, islands 93isolated in the aftermath of hydroelectric dams offer several advantages over non-insular 94forest patches abutting a terrestrial matrix, including the equitability of isolation time, an 95equally hostile uniform open-water matrix, and convergent histories of anthropogenic 96disturbance (Diamond 2001; Wu et al. 2003; Benchimol and Peres 2015a). Studies in 97hydroelectric reservoirs can therefore rule out confounding effects of matrix types, 98enabling assessments of fragmentation effects per se (Cosson et al. 1999). Nevertheless, no 99small mammal study to date has been carried out in hyper-diverse Amazonian reservoirs, 100where currently ambitious governmental plans are expected to greatly expand hydropower 101infrastructure (Lees et al. 2016). Understanding how biodiversity is affected by 102hydropower projects is then of critical importance for management actions in lowland 103Amazonia.

104 Here, we assess small mammal assemblage responses to habitat insularization 105induced by a major hydroelectric dam in Central Amazonia 28 years after isolation. The 106Balbina Hydroelectric Dam created a reservoir of 443,772-hectares 107(FUNCATE/INPE/ANEEL, 2000), comprising 3,546 forest variable-sized islands. We 108conducted quantitative surveys on 25 forest islands and four widely-separated mainland 109sites in neighbouring continuous forests to determine the main predictors of small mammal 110persistence within forest islands, and investigate whether small mammal assemblages 111exhibited a nested structure along the gradient of habitat insularization. We used a multi-112scale approach, initially considering potential area effects, and the combined effects of 113patch, landscape and habitat-quality metrics. We further classified each species in terms of 114their degree of forest-dependency to investigate the influence of this ecological trait in 115explaining patterns of local species extinction across islands, also testing for potential 116nestedness structure. We hypothesized that small mammal assemblages on smaller and 117more isolated islands that had been more degraded were represented by a simplified subset 118of mainly non-forest-dependent species. Conversely, nearly complete assemblages 119comprised of both forest and non-forest-dependent species are expected to persist on 120larger, less isolated islands containing high-quality forest habitat. In particular, non-forest-121dependent species are expected to present a lower degree of nestedness than forest-122dependent species.

124Methods

125Study area

126This study was carried out in the archipelagic landscape of the Balbina Hydroelectric 127Reservoir and its immediate surroundings, in Central Brazilian Amazonia (1°48'S, 12859°29'W; Fig. 1). The Balbina dam was built in 1986 on the Uatumã River, a left-bank 129tributary of the Amazon River. Given the typically flat topography of the area, this dam 130flooded a vast area of 312,900 ha of primary forest, within the 443,772-ha hydroelectric 131reservoir (FUNCATE/INPE/ANEEL 2000). In the aftermath of damming, the former 132hilltops of the pre-inundation forest area remained above-water as 3,546 land-bridge 133 islands widely distributed across the reservoir lake. Many dead relics of emergent trees are 134still standing within the open-water matrix, as the submerged primary forest had not been 135clear-cut. Most islands consist of dense closed-canopy terra firme forest, but many small 136islands were strongly affected by edge-related windfalls and ephemeral wildfires, which 137occurred during a late-1997 to early-1998 El Niño drought (Benchimol and Peres 2015a). 138The area within and around the former left bank of the Uatumã river has been legally 139protected since 1990 by the 942,786-ha Uatumã Biological Reserve, the largest reserve in 140its category in Brazil. This contributes to low levels of post-damming human disturbance 141across the reservoir. The mean annual temperature in this region is 28°C and the mean 142annual rainfall is 2,376 mm (IBAMA 1997). Small mammal communities were sampled on 14325 islands and four continuous forest sites (hereafter, CF; Fig. 1). Islands were selected 144according to their size, degree of isolation and spatial distribution, resulting in a wide range 145of island configurations. Surveyed islands ranged in area from 0.55 to 14,660 ha (mean \pm 146SD: 199 ± 344 ha; Table S1), and isolation distances from each focal island to the nearest 147mainland continuous forest ranged from 44 to 11,872 m (4,351 \pm 3,386 m).

149Small mammal sampling

150Small mammals were sampled using transects during sessions of 16 consecutive nights 151over two field seasons, from April to November 2014 and April to November 2015. Each 152transect consisted of a set of nine stations of live traps (hereafter, LTs), followed by an 153array of three pitfall units. Each LT station was placed 20-m apart from others and included 154two Sherman traps (23 x 9 x 8 cm, H. B. Sherman Traps, Inc., Tallahassee, Florida) and 155one wire mesh trap (30 x 17.5 x 15 cm, Metal Miranda, Curitiba, Paraná). At each LT 156station, one trap was set on the ground, one in the understory (~1.5 m high), and one in the 157(sub)canopy (>10 m high). Traps of different types were placed alternatively on the ground 158and in the understory across consecutive stations, but only Sherman traps were placed in 159the canopy due to logistic limitations. The forest canopy stratum was sampled using an 160adaptation of the method described by Lambert et al. (2006). LTs were baited with a mix of 161bananas, peanut powder, sardines and oatmeal. Pitfall traps (100L) were also spaced at 20-162m intervals and connected by a plastic drift fence 50-cm high and 10-cm underground, with 16310 m of fence extending beyond the two terminal pitfalls.

The number of transect placed at different sites varied according to their area. This 165allowed us to obtain a higher number of individuals at larger forest sites, where overall trap 166density, and consequently the probability of an individual passing near a trap, were lower 167(Table S1). Due to spatial restrictions in small islands, alternative smaller transects were 168established therein. Thus, all islands smaller than 2 ha and those between 2 and 10 ha were 169sampled by transects containing only three LT stations followed by an array of one pitfall, 170and by six LT stations followed by an array of two pitfalls, respectively. Larger islands 171were sampled by as many as four transects, according to their size classes: 10 to 50 ha, 50 172to 200 ha, 200 to 500 ha and >500 ha, respectively; CF sites were sampled by either six 173(CF₁ and CF₃), 10 (CF₂) and 12 transects (CF₄). Due to logistic restrictions, CF₂ and CF₄

174were only sampled during either the first (2014) or second (2015) field season,
175respectively, which reflects the higher number of transects placed therein (for further
176details on sampling effort per site, see Table S1).

177 We conducted a total 65,520 trap-nights across 79 transects. All traps were 178 inspected daily and whenever live captures could not be identified in the field, a maximum 179 of five voucher specimens per species per site were collected during the first season, and 180 deposited at the Mammal Collection of the Instituto Nacional de Pesquisas da Amazônia 181(INPA), in Manaus, Brazil. All other individuals recorded were weighted and tagged (Fish 182and Small Animal Tag, size 1; National Band and Tag Co., Newport, Kentucky), so that 183 any subsequent recaptures could be distinguished. Additionally, tissue samples were 184collected from all individuals recorded and deposited at INPA. However, we were not 185always able to identify at the species-level records of *Proechimys* spp. (*P. cuvieri* and *P.* 186guyanensis) and Oecomys spp. 1 (O. roberti and O. bicolor) at all sites. Because these 187congeners are ecologically very similar (Jones et al. 2009), we further refer to those taxa as 188'ecospecies'. To streamline, we hereafter use 'species' to refer to both species and 189ecospecies. We followed ASM guidelines (Sikes et al. 2016), and the study was approved 190by the appropriate Brazilian institutional animal care and use committee (SISBIO license 191No. 39187-4).

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193*Patch*, *landscape* and *habitat* quality variables

194Patch and landscape metrics were calculated using ArcMap 10.1 (ESRI 2012), based on 195high-resolution multi-spectral RapidEye imagery (5-m resolution with 5-band color 196imagery) of the entire study landscape. At the patch scale, we measured island area and 197shape, and at landscape-scale, we obtained for each surveyed island a proximity (PROX) 198metric to other islands and CF sites, and their nearest distances to CF (Table 1; for further

199details on imagery processing and landscape metrics, see Benchimol and Peres 2015a, 2002015b). We used Generalized Linear Models (GLMs) to analyse the relationship between 201the species richness in the islands and the proximity index considering in separate models 202at multiple scales of effects in terms of increasing buffer radius (Jackson and Fahrig 2012) 203–250 m, 500 m and 1 000 m. We then ordered the models according to its Akaike 204Information Criteria (AIC; Burnham and Anderson 2002). For further analyses, we 205retained the PROX index obtained using the 500 m-radius (AIC = 129.2), rather than 1 000 206m (129.2) or 250 m (130.64).

Small mammal species responses to habitat fragmentation are closely linked to 208habitat structure, such as overstory and understory vegetation density (Delciellos et al. 2092015). We thus obtained habitat variables by measuring forest composition and vegetation 210structure, which further reflect the habitat quality for small mammals. Vegetation variables 211were obtained from floristic surveys within 0.25-ha (250m × 10m) forest plots established 212on each focal island and CF site, where all trees \$10cm diameter at breast height (DBH) 213were measured and identified at species-level. The number of plots surveyed per site varied 214to the area of each site, ranging from one to four plots (see Table 1 for a description of 215floristic variables; details in Benchimol and Peres 2015c). Additionally, we conducted a 216semi-supervised classification to obtain four land cover classes (closed-canopy forest, 217open-canopy forest, bare ground, and water) using ArcMap 10.1, and obtained the 218percentage of closed-canopy forest within the island from the RapidEye imagery (Table 1).

220Forest-dependency index

221Degree of forest habitat-dependency (FD) was defined as the species-specific ratio 222between the capture rate in the open habitat matrix and in continuous primary forest. 223Forest-dependency estimates were based on small mammal data obtained during a 5-year

224study by J. R. Malcolm (1991) at the Biological Dynamics of Forest Fragments Project 225(BDFFP), a fragmented landscape ~100 km from the Balbina reservoir, which shares the 226same small mammal fauna. Forest and non-forest-dependent species were those for which 227a FD < 1 and FD > 1, respectively. We obtained the community-averaged FD values by 228summing the FD values of all individuals recorded at each survey site, and dividing this by 229the number of individuals therein. Community-averaged FD values ranged between 0.061 230–0.688, higher values corresponding to an increased prevalence of non-forest-dependent 231species. Species FD were previously transformed to range between 0 (highest forest-232dependency) and 1 (lowest forest-dependency). Information on capture rates at the BDFFP 233landscape within either primary forest or the matrix was unavailable only for the squirrel 234Guerlinguetus aestuans, which occurs chiefly in primary forests (Patton et al. 2007). Based 235on ecological similarities between closely related taxa (Jones et al. 2009), we assigned FD 236values to congener species for which information was unavailable (Oecomys, Neacomys 237and Marmosops). Capture rates and FD values are summarized for all species in Table S2.

239Data analysis

240We excluded from analysis two Echimyidae — $Makalata\ didelphoides$ and Echimys 241chrysurus — that were recorded only once during the study. These species feed on seeds 242and leaves, in addition to some fruit (Patton et al. 2000), and consequently are rarely 243attracted to the bait used here. The adequacy of small mammals sampling was evaluated 244using the sample coverage estimator (sensu Chao and Jost 2012), which estimates the 245proportion of the total number of individuals in an assemblage that belong to the species 246represented in the sample. Our small mammal trapping was satisfactorily accurate in 247relation to our sampling effort, averaging 0.92 ± 0.07 per site in the most complete 248assemblage, except for two small islands ('Xibé' and 'Abu'), in which sample coverage

249was zero (Table S1). Despite the lack of sampling representativeness in those two islands, 250we still retained them in the analysis because trap density therein was much higher (~4.57 251traps/ha) than in islands larger than 100 ha (0.08 - 0.54 traps/ha) and CF sites (~0.002) 252traps/ha; Table S1). Nevertheless, to account for any eventual undetected species and avoid 253 any potential bias in species richness values, we estimated a bootstrapped number of 254species. To do so, we applied a bootstrap method that determines confidence intervals 255around Hill numbers, facilitating the comparison of multiple assemblages of extrapolated 256samples (Chao et al. 2014). For each site, the bootstrapped number of species was obtained 257 from the extrapolation of individual-based Hill numbers assuming twice the number of 258individuals recorded at that site, using the R codes provided by Chao et al. (2014). We 259further verified if the variable number of traps deployed per site caused any bias on 260 observed numbers of species. To do so, we correlated the number of traps deployed per site 261with sample coverage estimations and the number of individuals recorded. Spatial 262autocorrelation was examined by correlating matrices of binary species composition and 263geographic distances among sites, applying a Mantel test using the 'vegan' R package 264(Oksanen et al. 2007).

Considering all 25 islands and their bootstrapped species richness, we performed 266species-area relationships (SARs) comparing eight possible SAR models: power, 267exponential, negative exponential, Monod, rational function, logistic, Lomolino and 268cumulative Weibull (for each model formula, see Table S3), using the 'mmSAR' R package 269(Guilhaumon et al. 2010). In addition, given the importance in identifying critical 270thresholds beyond which species responses change abruptly, we also included a piecewise 271regression model to explain small mammal SARs (Toms and Lesperance 2003), using the 272'segmented' R package (Muggeo 2017). Data fitting for each model was compared

273according to Akaike Information Criterion values corrected for small sample sizes (AICc: 274Burnham and Anderson 2002).

We further estimated the degree of nestedness in small mammal assemblages for 276each sampling site using NODF, a metric based on overlap and decreasing fill of presence-277absence matrix data (Almeida-Neto et al. 2008). Using this method, probability levels can 278be assessed using *Z* scores, i.e., by comparing the observed nestedness value with the mean 279of a series of values obtained by reshuffling the original matrix to produce a number of 280random matrices according to a certain null model (Strona et al. 2014). These analyses 281were performed using the 'Nestedness for Dummies' Program (Strona et al. 2014), 282considering the 'proportional column and row totals' algorithm to construct 999 simulated 283random matrices. Interaction matrices of small mammal species and sites ordered by 284species richness were further obtained using the '*Vegan*' R package. Prior to analysis, we 285added the data from all four mainland continuous forest sites. Nestedness analyses were 286further repeated separately for each species grouping (i.e., forest- and non-forest-287dependent), using the same procedure

Species composition was also analysed using Principal Coordinate Analysis 289(PCoA) based on a quantitative Bray-Curtis similarity matrix of species composition. 290Species abundances were previously standardized for each site due to differences in 291sampling effort per site. Aggregate biomass (log₁₀ x) was defined as the sum of the body 292mass of all individuals recorded at any given site per unit of trapping effort. For that, we 293considered the body mass of each live capture weighted during the field work.

To evaluate the combined effects of patch, landscape and habitat quality metrics on 295small mammal assemblages (bootstrapped species richness, species composition, 296standardized abundance, and aggregate biomass), we performed Generalized Additive 297Models (GAMs). GAMs do not require a linear relationship between the response and the

298explanatory variables, and were performed using a Gaussian error structure. We controlled 299 for high levels of variable inter-dependence by performing a Pearson correlation matrix, 300retaining weakly correlated variables (r < 0.70, P > 0.05). 'Island shape', 'percentage of 301old-growth trees' and 'fire severity' were correlated and therefore excluded from further 302analysis. We did not include CF sites in this analysis because that additionally overinflated 303the correlation between 'island area' and 'proximity' (r = 0.86, P < 0.0001), further 304invalidating the inclusion of one these variables in the model. We also calculated the 305 Variation Inflation Factors (VIFs), any of the remaining variables was moderately 306redundant, presenting VIF < 5 (Dormann et al. 2013). A candidate model set was further 307constructed, using all additive combinations of the eight explanatory variables retained, 308and models were ranked based on their AICc, using the 'MuMIn' R package (Bartoń 2014). 309To account for model uncertainty in multi-model inference, a model-averaging approach 310was performed using only the most plausible models (i.e., $0 < \Delta AICc > 2$, $\Delta AIC = AIC_i -$ 311AIC_{min} in which $i = i^{th}$ model). The relative importance (RI) of each variable contained in 312that model set was obtained by the sum of the Akaike weights of the models in which that 313 variable had been included (Rhodes et al. 2009). Explanatory variables were previously 314standardized (x = 0, σ = 1) to place coefficient estimates into the same scale. The same 315modelling procedures were repeated considering the number and abundance of forest and 316non-forest-dependent species separately. When analysing graphically the isolated effects of 317island area and proximity on species richness and abundance of forest and non-forest-318dependent species, we improved data fitting by performing simple GLMs, both including 319and excluding the quadratic term of each explanatory variable. AIC values were compared 320between the models including and excluding the quadratic term of the explanatory variable 321(Burnham and Anderson 2002). We then added the quadratic term of the explanatory 322variable when examining the relationship between the number of forest-dependent species

323and island area, and between the number of non-forest-dependent species and island 324proximity (Table S4). All data analyses were performed in R (R Development Core Team 3252015).

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327Results

328A total of 884 small mammal individuals was recorded across the 29 sampled sites, 329amounting to an overall mean capture success of 1.35% per trap-night (excluding 330recaptures). A total of 22 species was recorded considering all sites, representing 12 331rodents and 10 marsupials (see Table S5). Excluding the two singleton records of E. 332*chrysurus* and *M. didelphoides*, the number of species recorded per site ranged from 1 to 33316 (mean \pm SD: 5.76 \pm 3.80 species) for all islands, and from 9 to 16 for CF sites (12.5 \pm 3342.5 species). Nearly all species recorded on islands were also present in at least one CF 335site, except for the small terrestrial rodent *Neacomys paracou*, which was recorded at only 336one island (N = 2 individuals). The most abundant species were the arboreal marsupial 337Marmosa demerarae (N = 195) and the terrestrial rodent Hylaeamys megacephalus (N = 195) 338187); while the arboreal echimyid rodent *Isothrix pagurus* (N = 6) and sciurid 339 Guerlinguetus aestuans (N = 4) were among the least detected species across all sites (see 340Table S3). On the basis of those species, our index of forest-dependency resulted in 11 341species classed as forest-dependent and 9 species as non-forest-dependent (Table S2). 342Furthermore, the number of traps deployed at each site was neither correlated with sample 343coverage estimations (r = -0.012, P = 0.951) nor with the overall number of individuals 344recorded (r = 0.245, P = 0.200), supporting the notion that the number of traps deployed at 345any given site did not induce any bias in the number of species recorded therein. Also, 346small mammal species composition was largely unaffected by the geographic distance 347between sites (r = 0.037, P = 0.269, N = 406 pairwise comparisons).

349Species-area relationships and nestedness

350The power, rational function, logistic and cumulative Weibull SAR models performed 351similarly well in explaining the bootstrapped species richness (S) of small mammals across 352all 25 sampled islands, explaining 68.7 - 71.0% of the variation in S (Table S3). 353Henceforth, to facilitate comparisons with most other studies, we focused our results on the 354power model. As such, the observed SAR clearly showed fewer species over a decreasing 355gradient of island size (z-value = 0.289, c-value = 2.089, R^2 = 0.687; Table S3). Islands 356smaller than 15 ha retained highly impoverished small mammal assemblages (mean \pm SD: 3572.6 \pm 1.3 species), whereas islands larger than 200 ha on average harboured 10.8 \pm 1.3 358species, which was comparable to CF sites in the mainland (12.5 \pm 2.5; Fig. 2).

Across the decreasing gradient of island size, small mammal community 360disassembly further appears to be mediated primarily by the selective extirpation of forest-361dependent species, in that small islands mostly retained non-forest-dependent species, as 362denoted from the high community-averaged FD values. In contrast, community-averaged 363FD values remained relatively low across CF sites (range = 0.08 - 0.11; Fig. 2). Moreover, 364overall small mammal assemblages presented a significantly nested structure (matrix size: 365520, fill = 0.313; $NODF_{row} = 62.493$, Z = 9.731, P < 0.001). Similar nestedness results 366were obtained when considering only either forest-dependent species (matrix size: 220, fill 367 = 0.341; $NODF_{row} = 56.055$, Z = 4.086, P < 0.001), or non-forest-dependent (matrix size: 368234, fill = 0.376; $NODF_{row} = 60.682$, Z = 4.645, P < 0.001; Fig. 3).

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370Combined effects of patch, landscape and habitat quality metrics

371Multiple-scale landscape, patch and habitat quality variables clearly affected the overall 372persistence of small mammal species across all 25 islands surveyed, with species

373composition best predicted by both island area ($\beta_{AREA} = 0.229$, P = 0.023, RI = 0.84) and 374tree species richness ($\beta_{S.TREES} = 0.148$, P = 0.033, RI = 0.67; Fig. 4a, Table S6). In fact, 375despite the lack of significant nestedness among the survey sites, the narrow set of non-376forest-dependent species persisting at all islands smaller than 2 ha was virtually the same. 377Most of the other islands, ranging from 2 to 800 ha, retained highly idiosyncratic species 378compositions, as indicated by the poorly predicted PCoA1 values obtained for those 379islands ($R^2 = 0.417$; Fig. S1).

The bootstrapped number of species at each island was best predicted by both 381 island area ($\beta_{AREA} = 4.227$, P = 0.001, RI = 0.75) and proximity ($\beta_{PROX} = 3.312$, P = 0.002, 382RI = 1.00; Fig. 4b, Table S5). When species were distinguished based on their degree of 383 forest-dependency, the number of forest-dependent species responded similarly, being 384 positively affected by both island area ($\beta_{AREA} = 3.107$, P = 0.001, RI = 1.00) and proximity 385($\beta_{PROX} = 1.289$, P = 0.039, RI = 1.00; Fig. 4c). Richness of non-forest-dependent species, 386 however, was positively affected only by proximity to other land-masses ($\beta_{PROX} = 1.227$, P = 0.011, RI = 0.88; Fig. 4d, Table S7). Again, the turnover in the number of forest and non-388 forest-dependent species can be observed across the size gradient from insular and 389 continuous forest sites: forest-dependent species were often missing from small islands but 390 were more abundant than non-forest-dependent species at CF sites (Fig. 5a). The same was 391 not clearly observed across the gradient of proximity to other land-masses (Fig. 5c).

Although none of the variables considered here performed well in predicting 393 overall species abundance (Fig. 4e, Table S6), the abundance of forest-dependent species 394 was positively affected by island area ($\beta_{AREA} = 6.597$, P = 0.005, RI = 0.84) and negatively 395 affected by the aggregate basal area of trees bearing fleshy fruits ($\beta_{FRUIT} = -2.799$, P = 3960.027, RI = 0.16; Fig. 4d). The abundance of non-forest-dependent species was negatively 397 affected by both island area ($\beta_{AREA} = 6.597$, P = 0.005, RI = 0.49) and proximity ($\beta_{PROX} = 3960.027$).

3986.597, P = 0.005, RI = 0.35; Fig. 4g, Table S7). Non-forest-dependent species accounted 399for a disproportionate fraction of the overall abundance at islands smaller than 15 ha 400(median (min – max) = 100% (50 –100%); Fig. 5b), while the abundance of the two 401functional groups in terms of forest habitat-dependency was comparable between weakly 402isolated large islands and CF sites (Fig. 5b, d). Indeed, except for the least detected species 403(*N. guianensis*, *N. paracou* and *Monodelphis arlindoi*), non-forest-dependent species were 404clearly widespread across all island sizes and CF sites. Forest-dependent species instead 405are completely absent from very small islands (< 2 ha) and 45.5% only occurred in islands 406larger than 200 ha (Fig. S2). Finally, the aggregate biomass of small mammals across all 25 407islands could not be predicted by any of the variables considered here (Table S6).

408

409Discussion

410As new capital investment fuels hydropower frontier expansion across wilderness regions 411in tropical countries (Castello et al. 2013; Zarf et al. 2015), decision-makers should 412carefully weigh the landscape-wide ecological impacts of this infrastructure against their 413overall socioeconomic benefits (Fearnside and Pueyo 2012; Lees et al. 2016). In lowland 414Amazonia, damming creates disproportionately large reservoirs and highly fragmented 415archipelagos of land-bridge forest islands (Junk and Mello 1990). Under this landscape 416context, we document widespread local extinctions of small mammal species across the 417archipelago, which is comprised of myriad of small islands. Small mammal assemblages 418further exhibited a nested structure, so that smaller and more isolated Balbina islands 419exhibited depauperate small mammal assemblages, in which forest-dependent species were 420typically either missing or persisted at very low abundances.

421

422Island biogeography revisited

423Island size and degree of isolation were the strongest predictors of the number of small 424mammal species at Balbina forest islands. Islands larger than 200 ha and mainland forest 425sites contained up to 12 and 15 species, respectively, while those smaller than 15 ha 426supported a maximum of four species, or fewer than one-third of the number of species 427expected to persist in large islands and mainland forest sites. Although large islands 428harboured nearly full complements of species, they account for only 1.8% of the 3,546 429islands in Balbina reservoir, whereas those smaller than 15 ha correspond to 62.7% of all 430islands (see histogram in Fig. 1). Our findings are consistent with the overall negative 431response of terrestrial species and communities to isolation in man-made reservoir islands 432reported worldwide (Jones et al. 2016). Indeed, medium and large-bodied terrestrial 433vertebrates were greatly affected by island size within the Balbina reservoir, with 95% of 434all islands retaining fewer than 60% of all 35 native vertebrate species (Benchimol and 435Peres 2015a). In other tropical reservoirs containing forest islands smaller than 350 ha, 436small mammals experienced local extinctions at similar (Guri Reservoir, Venezuela: 437Lambert et al. 2003) or higher rates than those observed here, often culminating in the 438persistence of only a single dominant species (Gatun Lake, Panama: Adler and Seamon 4391996; Saint-Eugène reservoir, French Guiana: Grajon et al. 2002; Chiew Larn Reservoir, 440Thailand: Gibson et al. 2013). Yet those studies do not provide a comprehensive scenario, 441failing to include large islands (but see Wang et al. 2010a for a reservoir in southern 442China). Given the observed ability of certain terrestrial mammal species to swim between 443insular forest patches (e.g. Grajon et al. 2002), large islands play a critical biotic role in the 444archipelagic landscape, by harbouring resident populations and operating as a source of 445emigrants for some species, depending on island isolation (Adler and Seamon 1996). 446 Second to island area, patterns of small mammal species richness at Balbina were 447best predicted by isolation. This is in part consistent with seminal ideas from Island

448Biogeography Theory (IBT), in which islands experiencing higher colonization rates
449should harbour more species (MacArthur and Wilson 1967). However, higher colonization
450rates were primarily a function of the functional connectivity of an island in the landscape
451(as inferred by the Proximity index) rather than the distance to the mainland, as predicted
452by IBT. Studies in terrestrial landscapes have found similar patterns of fragmentation
453effects in terrestrial landscapes for both small mammals (e.g. Goodman and
454Rakotondravony 2000; Pardini et al. 2005), and other taxa (see Predevello and Vieira
4552010). Our findings therefore unequivocally support the notion that small, isolated
456reservoir islands have more limited conservation value than non-island habitat fragments in
457entirely terrestrial landscapes, which agrees with paleotropical small mammal data from
458reservoir islands in southern Thailand (Gibson et al. 2013).

459

460Determinants of community composition

461Communities isolated in land-bridge islands may exhibit nested structures over time, as has 462already been detected for small mammal, birds and lizard assemblages within the ~50 year 463old Thousand Island Lake, China (Wang et al. 2010a). In Balbina, the nested structure 464observed in small mammal assemblages are probably related to the differential species 465susceptibility to extinction (Lynam and Billick 1999). Indeed, the subset of small mammal 466species persisting at Balbina islands was closely related to both patch and local habitat 467characteristics, as suggested by the importance of island area and tree species richness in 468predicting species composition. Small islands likely retain a very limited spectrum of 469habitat resources (August 1983), so they are consistently occupied by the same subset of 470species, which were mainly non-forest-dependent. As island area increases, more 471ecological niches become available (Pardini et al. 2005), which facilitates higher species 472packing, including forest-dependent species. Similarly, habitat structure was one of the

473main determinants of small mammal assemblages in Brazilian Atlantic Forest fragments 474(Pardini et al. 2005; Delciellos et al. 2015). Nevertheless, although forest-dependent 475species exhibited an expected nested structure, the same was observed for non-forest-476dependent species. This might be due to the inclusion of certain species rarely captured 477throughout the sampling, that were particularly evident at larger islands and continuous 478forest sites (e.g., *Monodelphis arlindoi*, *N. paracou*, and *N. guianae*).

479 Moreover, despite the nested structure of the Balbina small mammal assemblages, 480larger islands and continuous forest sites still presented an idiosyncratic species 481composition (Figs. S1, S2). This result contrasts with other faunal groups at the Balbina 482landscape, in which species composition converged along the gradient of island area 483(Benchimol and Peres 2015b; Aurélio-Silva et al. 2016). The idiosyncrasy in species 484composition observed in this study could result from the reduced sample coverage in larger 485islands and continuous forest sites. Regardless of the overall satisfactory sample coverage 486of estimators obtained, the difference between observed and bootstrapped species richness 487 was higher in continuous forest sites (Fig. S3). Even considering the higher sampling effort 488at larger forest sites, the density of traps was 21 - 207 times lower therein (0.20 and 0.02) 489traps/ha at islands > 200 ha and CF sites, respectively) compared to small islands (4.14 490traps/ha). Therefore, the probability of an individual passing near a trap was much higher 491in small islands due to higher trap saturation, and also due to vertical forest compression, 492as larger forest areas were both more saturated and multi-layered (AFP, CAP, MB, unpubl. 493data). In addition, large islands' transects may cover a limited part of the ecological 494distribution of species, which are often aggregated due to patchy trophic resources 495(Charles-Dominique et al. 1981). Our results may therefore underestimate the number of 496species in larger islands and continuous forest sites. This further amplifies the strength of 497species-area effects, and suggesting that SAR patterns presented here are conservative.

499Vulnerability of forest-dependent species

500In entirely terrestrial fragmented landscapes, habitat-generalist species responses are most 501likely due to their smaller spatial requirements, ability to access and take advantage of 502matrix resources, and trophic plasticity (Pardini et al. 2005; Umetsu and Pardini 2007; 503Santos-Filho et al. 2012). Interestingly, those general patterns also hold true in a real island 504landscape such as Balbina, even though matrix resources are entirely unavailable. 505Likewise, forest specialist small mammal species greatly declined in forest patches 506surrounded by small amounts of forest cover in the Brazilian Atlantic Forest, whereas those 507non-forest-dependent species were unaffected by habitat loss (Estavillo et al. 2013). Using 508the same species classification, Santos-Filho et al. (2016) noted the same trend in species 509richness of small mammals in southern Amazonian forest fragments. Unlike forest-510dependent species, habitat generalists were able to persist in tiny islands with highly 511reduced trophic and structural resource availability. Rodent species persisting in islands 512within China's Three Gorges Dam also exhibited dietary shifts compared to their baseline 513diet in the mainland (Wang et al. 2010b).

In contrast to species richness, the abundance of forest-dependent species was not affected 515by island isolation. The abundance of these species may therefore be mainly determined by 516within-island processes rather than movement between islands. For forest-dependent species, 517such processes may be idiosyncratic in relation to each island, and unrelated to landscape 518variables. In contrast, non-forest-dependent species were hyper-abundant in small isolated 519islands, so that overall species abundance was compensated for at those sites. Because only non-520forest-dependent species increased in abundance in small islands, higher species abundances 521there are unlikely a consequence of "fence effects" in which increased island isolation limits 522dispersal (Adler et al. 1986). Previous neotropical small mammal studies also reported relatively

523higher abundances in small fragments compared to continuous forest (Malcolm 1991; Passamani 524and Fernandez 2011; but see Santos-Filho et al. 2012), including both true islands (Glanz et al. 5251990; Lambert et al. 2006), and forest patch isolates embedded within terrestrial matrices 526(Laurance 1994; Lynam and Billick 1999; Pardini et al. 2005; Vieira et al. 2009). Apparently, 527 species that could persist in small isolated islands were additionally well positioned to increase 528their abundances therein. This over-inflated abundance could be explained by the absence of 529predators (Glanz et al. 1990; Adler and Seamon 1996; Terborgh et al. 1997; Lambert et al. 2006) 530or density compensation in the absence of other small mammal species (Fonseca and Robinson 5311990). Benchimol and Peres (2015b) reported the local extinction of most medium and large-532bodied vertebrates in small Balbina islands, including mammalian predators of small mammals, 533and this is also the case of diurnal and nocturnal raptors (A.S. Bueno, unpubl. data). Moreover, 534despite the low trophic resource availability at small disturbed islands, non-forest-dependent 535species may increase in abundance by additionally exploring novel resources (Pardini et al. 2005; 536Wang et al. 2010b), which may become available once other species have become locally 537extirpated (Fonseca and Robinson 1990). The negative relationship between the abundance of 538 forest-dependent species and the aggregate basal area of trees bearing fleshy fruits was 539unexpected. This could be due to the higher occupancy or abundance of large-bodied terrestrial 540mammals, such as white-lipped peccary *Tayassu pecari* (Benchimol and Peres 2015b), which 541 could negatively affect the abundance of forest-dependent species. Further studies should 542investigate interspecific interactions between large and small mammals.

543

544Conservation implications

545In line with previous findings in non-insular forest fragments (Watling and Donnelly 2006; 546Vieira et al. 2009), fragmentation effects on small mammal assemblages across a true 547archipelagic landscape were best predicted by island area and isolation. As explained

548above, however, this illustrates a slight departure from the central tenets of Island 549Biogeography Theory, and highlights the importance of taking into account the spatial 550configuration of habitat remnants in landscape-scale conservation plans. This should be 551extended to environmental impact assessments (EIAs) prior to licensing of hydropower 552development, or other infrastructure projects resulting in fragmented forest landscapes 553(Fahrig 2003). In addition, species classification according to degree of forest-dependency 554could efficiently distinguish species exhibiting different patterns of persistence across the 555archipelagic landscape. Species ability to traverse the matrix should therefore also be 556considered, as previously noted (Pardini et al. 2010; Santos-Filho et al. 2016). Moreover, 557as the aquatic matrix acts as a strong environmental filter, ecosystem functioning could 558undergo severe changes at small, isolated islands occupied by the same set of 559hyperabundant species, including the loss of ecosystem processes such as forest 560regeneration and arthropod control (Terborgh et al. 1997, 2001). As the vast majority of 561Balbina islands are small, their biodiversity and ecosystem functioning trajectories can 562already be seriously compromised. In this way, creation of myriad small islands within vast 563hydroelectric archipelagos is not conducive to maintaining high levels of biodiversity and 564forest functionality in the long term (Jones et al. 2016), which should be considered prior 565to licensing future hydropower projects. If any given mega-dam becomes completely 566unavoidable, hydraulic engineers should prioritize the creation of large islands by 567reassessing the relationship between maximum operational water level of a reservoir and 568the spatial configuration of its archipelago.

We finally emphasize that our study illustrates a 'best-case' scenario in terms of 570ecological impacts caused by habitat insularization, given that the Balbina flooded area and 571its surroundings have been strictly protected by the Uatumã Biological Reserve since 1990. 572Other hydroelectric reservoirs in the Brazilian Amazon lacking protection were rapidly

573occupied and deforested by smallholders (e.g. Tucuruí Hydroelectric Reservoir; Fearnside 5742001). As such, we highlight the importance of implementing protected areas as a 575mitigation measure, preventing subsequent hunting, fire disturbance and unplanned 576settlements therein. To make matters worse, insular ecosystems are likely to pay an 577extinction debt by gradually losing species even many years in the aftermath of island 578creation (Jones et al. 2016). Careful planning of hydropower development is therefore 579critical to avoid mass species extinctions and losses in ecosystem services in the world's 580mega-diverse tropical forests.

581

582Acknowledgements

583We thank all 15 volunteers/field assistants who assisted in the field; the Reserva Biológica 584do Uatumã and its staff for logistical support; M.N.F. da Silva for help with species 585identification and D. Storck-Tonon for assistance with landscape metrics. This study was 586funded by the Amazon Region Protected Areas Program (ARPA), Amazonas Distribuidora 587de Energia S.A., and Associação Comunidade Waimiri Atroari; Rufford Foundation (grant 588number 13675-1); Idea Wild; and a NERC grant (NE/J01401X/1) awarded to CAP. AFP 589and MB were funded by Coordenação de Aperfeiçoamento de Pessoal de Nível Superior 590(CAPES) scholarship, and MVV was funded by Fundação de Amparo à Pesquisa do 591Estado do Rio de Janeiro (FAPERJ) and Conselho Nacional de Desenvolvimento 592Científico e Tecnológico (CNPq).

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794Figure legends

Figure 1. Surveyed sites within the Balbina Hydroelectric Reservoir of Central Brazilian 797Amazonia: 25 land-bridge islands (in red and highlighted by a 1,000-m buffer contour) and 798four continuous forest sites (CF₁, CF₂, CF₃ and CF₄; indicated by red rectangles). Inset 799photographs illustrate an aerial view of the Balbina archipelagic landscape (photo credit: E. 800M. Venticinque) including the aquatic matrix. Histogram represents the size distribution of 801all 3,546 islands; red dots in the histogram represent the size distribution of surveyed 802islands.

804Figure 2. Species-area relationship for small mammal assemblages recorded across 25 805land-bridge islands and species richness at four continuous forest (CF) sites surveyed at the 806Balbina Reservoir landscape, considering the bootstrapped species richness. Data points 807are color-coded according to the community-averaged forest habitat-dependency (FD) 808values (log₁₀ x; see text). Shaded area represents the 95% confidence region. Boxplots 809indicate the median, 1st and 3rd quartiles, and minimum and maximum values of species 810richness in CF sites; solid dots indicate observed values.

812Figure 3. Nested matrices of small mammal species and sites: 25 land-bridge islands and 813an aggregation of the four continuous forest sites ('*Tfall*'). Sites are ordered top to bottom 814by species richness. Matrices, from left to right, represent the overall small mammal 815assemblage (grey cells), forest-dependent species (green cells), and non-forest-dependent 816species (orange cells). For each matrix, we indicate NODF_{row} and P-values. Trend lines 817illustrate the hypothetical maximum nestedness for each data set. Sites are described in 818Table S1, and the full species nomenclature is listed in Table S3.

Figure 4. Estimates of averaged models and their 95% confident intervals for predictors of 820(a) species composition (denoted by the PCoA axis 1), (b) bootstrapped species richness, 821number of (c) forest-dependent and (d) non-forest-dependent species, (e) overall species 822abundance, and abundance of (f) forest-dependent and (g) non-forest-dependent species. 823Predictors included: proximity (Prox), distance (Dist) to the mainland, island size (Area), 824tree species richness (S.trees), tree density (D.trees), proportion of closed-canopy forest 825(CC), aggregate basal area of fleshy-fruiting trees (Fruit) and density of lianas (Lianas). 826Statistically significant negative and positive coefficients are shown as red and blue 827colours, respectively.

Figure 5. Relationships between island area (top panels; (a) and (b)) and island proximity 830(bottom panels; (c) and (d)) and the richness (left panels) and abundance (right panels) of 831forest-dependent (green circles) and non-forest-dependent species (red circles) at the 832Balbina reservoir landscape. Species abundances are standardized by sampling effort. 833Shaded areas represent the 95% confidence regions. Boxplots for CF sites indicate the 834median, 1^{st} and 3^{rd} quartiles, and minimum and maximum values of species richness and 835abundance. The number of forest-dependent species increases with island area (P < 0.001) 836and proximity (P = 0.011), whereas the number of non-forest-dependent species increases 837only with island area (P = 0.005), while the abundance of non-forest-dependent species P = 0.005 (P = 0.005), while the abundance of non-forest-dependent species P = 0.005 (P = 0.005).

840Table

Table 1. Description of patch, landscape, and habitat quality variables quantified to 843examine small mammal assemblage properties within the Balbina landscape. The overall 844range, mean and standard deviation are provided for each variable.

Name (code name)	Variable description	Range (mean ± SD)
Landscape		
Proximity (PROX)	The sum of all island areas divided by the squared sum of edge-	2.75 – 4.54 x 10 ⁹
	to-edge distances from each focal island to all islands within a	$(1.89 \times 10^8 \pm 9.07 \times$
	500 m-buffer ($log_{10} x + 1$; McGarigal et al. 2012).	108)
Distance (DIST)	Euclidean distance from each island to the nearest	44 – 11,872 m
	neighbouring mainland forest site.	$(4,351 \pm 3,318)$
Patch		
Island size (AREA)	Total island area of each focal island $(\log_{10} x)$.	0.83 – 1466.00 ha
		(199.0 ± 344.1)
Island shape	Total perimeter length of each focal island divided by the total	0.004 - 0.106
(SHAPE)	island area.	(0.018 ± 0.022)
Habitat quality		
Trees richness	Number of tree species calculated from floristic surveys in	14 – 66
(S.TREES)	0.25-ha forest plots within each focal island or mainland site.	(54.6 ± 11.5)
	Turner density coloniated from file sixtic comments 0.25 l. f.	
Trees density	Trees density calculated from floristic surveys in 0.25-ha forest plots within each focal island or mainland site.	84 – 176

(D.TREES)		(123.4 ± 22.8)
Closed-canopy	Percentage of closed-canopy forest within the focal island.	37.5 – 99.35
forest (CC)		(76.2 ± 15.5)
	Fire severity within each focal island or mainland site, scored	
Fire severity	as an ordinal scale based on the extent of each forest site	1 - 3
(FIRE)	affected by surface fires and the number of charred trees and	(1.96 ± 0.60)
	height of char marks on each tree.	
Old-growth trees	Percentage of old-growth trees calculated from floristic surveys	10.71 – 82.34
(OGT)	in 0.25-ha forest plots within each focal island or mainland site.	(64.1 ± 17.0)
Basal area of	Basal area of trees bearing fleshy fruits, calculated from	12.1 –33.0
fleshy-fruiting trees (FRUIT)	floristic surveys of all live trees ≥10 cm DBH in 0.25-ha forest plots within each focal island or mainland site.	(20.6 ± 5.0)
Lianas (LIANAS)	Mean number of lianas (> 2.5 cm DBH) calculated from floristic surveys in 0.25-ha forest plots within each focal island or mainland site.	$0 - 40.5$ (21.5 ± 10.3)
	or maintaine site.	

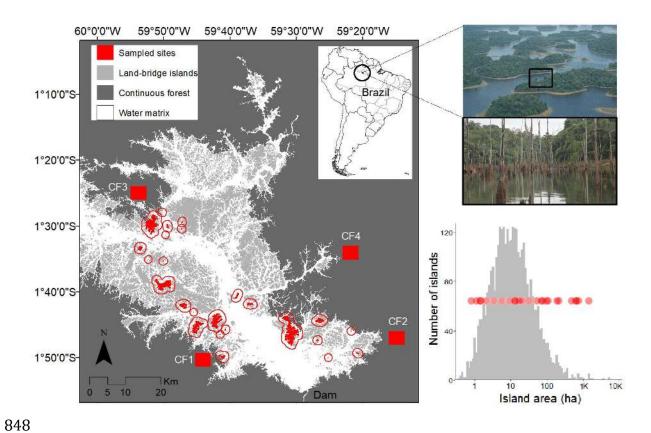


Figure 1

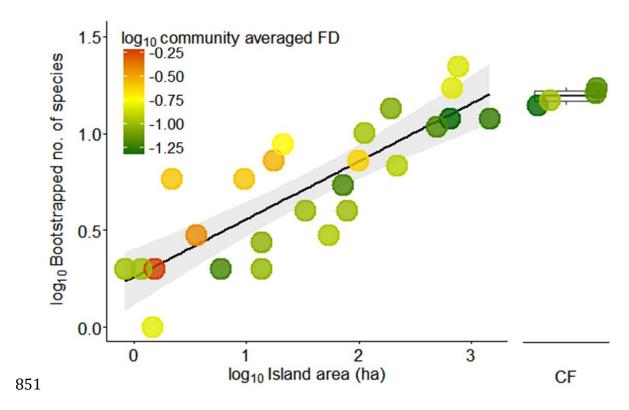


Figure 2

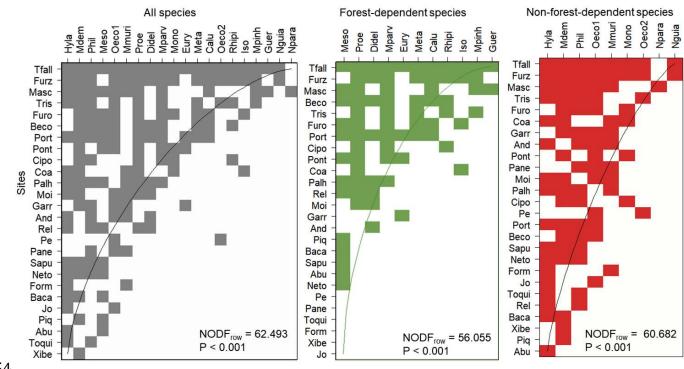
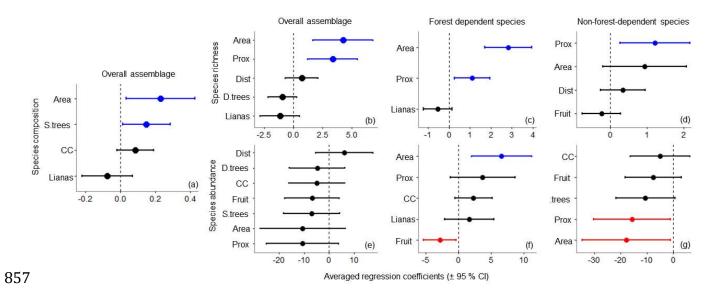
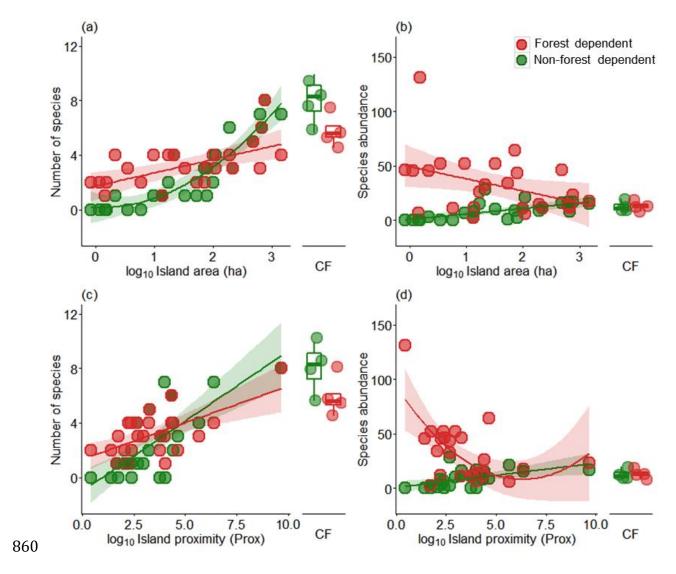


Figure 3



858Figure 4



861Figure 5