1	Patch-scale biodiversity retention in fragmented landscapes:
2	reconciling the habitat amount hypothesis with the island
3	biogeography theory
4	
5	Running title: Biodiversity retention in habitat patches
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7	Anderson Saldanha Bueno and Carlos A. Peres
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9 10 11 12 13	A.S.B. (corresponding author) School of Environmental Sciences, University of East Anglia, Norwich, Norfolk, NR4 7TJ, United Kingdom, and Instituto Federal de Educação, Ciência e Tecnologia Farroupilha, Júlio de Castilhos, RS, CP 38, CEP 98130-970, Brazil. Email: buenoas@gmail.com. ORCID: 0000-0001-7416-6626
14 15	C.A.P. School of Environmental Sciences, University of East Anglia, Norwich, Norfolk, NR4 7TJ, United Kingdom. Email: c.peres@uea.ac.uk. ORCID: 0000-0002-1588-8765

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## 28 ABSTRACT

- 29 Aim: To test whether the species richness of understorey insectivorous birds on forest islands
- 30 induced by a major hydroelectric dam is best explained by either the island biogeography
- theory (IBT) or the habitat amount hypothesis (HAH). Given the low dispersal ability of the
- 32 focal species group and the hostile water matrix, we predict that the species richness will be
- predominantly driven by an island effect as posited by the IBT, rather than a sample area
- 34 effect as posited by the HAH.
- Location: Forest islands within the Balbina Hydroelectric Reservoir, central Brazilian
  Amazonia.
- 37 **Taxon:** Birds.
- 38 Methods: We mist-netted birds at 33 forest islands (0.63–1,699 ha), totalling 874 individuals
- of 59 species. The size of the local landscape used to calculate the habitat amount was
- 40 determined by a multi-scale analysis in which buffers around mist-net lines ranged from 50 to
- 41 2,000 m. We applied four tests to examine whether the species richness on forest islands is
- 42 predominantly driven by either an island effect (island size) or a sample area effect (habitat
- 43 amount).
- Results: From the four tests applied, one was consistent with an island effect, two were
  regarded as inappropriate to test the HAH, and one could not be adequately addressed due to
  island size being highly correlated with habitat amount in the local landscape (200-m buffer).
- Main conclusions: Some of the proposed ways of testing the HAH may lead to misleading conclusions. The relative importance of island size in determining the species richness of understorey insectivorous birds on forest islands is higher than that of surrounding habitat amount, thereby providing stronger support for IBT. We propose a conceptual framework, based on the degree of matrix permeability and species dispersal ability, to determine to what extent a patch- or landscape-centric worldview in landscape ecology provides the most appropriate framework to assess the effects of habitat fragmentation on biodiversity.
- 54
- 55 Key-words: Amazonia, habitat amount hypothesis, habitat fragmentation, habitat loss,
- 56 insularization, island biogeography theory, SLOSS, species richness, species-area
- 57 relationship, tropical forest
- 58

#### 59 INTRODUCTION

60 MacArthur & Wilson's (1967) island biogeography theory (hereafter, IBT) has been 61 widely applied as a paradigmatic conceptual framework in habitat fragmentation ecology, implying that habitat patches are analogous to oceanic islands surrounded by a hostile matrix 62 (Haila, 2002; Laurance, 2008). However, such analogy has been repeatedly challenged since 63 64 IBT does not account for many factors operating in fragmented landscapes (Laurance, 2008; Wiens, 2008), which were later incorporated into a landscape ecology framework (Haila, 65 66 2002). For example, species move among suitable habitat patches as a function of varying 67 degrees of terrestrial matrix permeability (Powell et al., 2013), indicating that habitat patches exert weaker boundaries to local populations and their derivative assemblages compared to 68 oceanic islands. If habitat patches fail to behave as discrete spatial units, the universally 69 celebrated species-area relationship (hereafter, SAR) – which is widely observed in 70 fragmented landscapes (Matthews et al., 2016) – may be governed at spatial scales larger than 71 that of island effects driven by habitat patch size. 72

With this in mind, Fahrig (2013) proposed the habitat amount hypothesis (hereafter, 73 HAH), which posits that (1) habitat patches are not discrete spatial units, and (2) the habitat 74 surrounding any given patch is the main source of immigrants. The underlying mechanism of 75 SARs in fragmented landscapes is therefore predicted to be the sample area effect, rather than 76 the island effect. Accordingly, sample sites within larger habitat patches harbour more 77 species because they are also associated with a greater amount of surrounding habitat. 78 79 Meanwhile, sample sites associated with the same amount of landscape-scale habitat should 80 harbour the same number of species, regardless of patch size (Fig. 7 in Fahrig, 2013). Such notion implies that conservation efforts should primarily focus on increasing the overall 81 habitat amount (i.e. the proportion of habitat in the landscape) without necessarily 82 considering its spatial arrangement (i.e. size and isolation of individual habitat patches) 83 84 (Seibold et al., 2017).

85 The generalisation of the HAH was initially criticised since its application was considered to be restricted to small-scale landscapes containing large habitat amounts 86 (Hanski, 2015), although the HAH was yet to be tested (Fahrig, 2015). Recent empirical 87 88 studies carried out in a variety of natural (e.g. forest fragments, fluvial islands, calcareous 89 grasslands) and experimental fragmented landscapes (e.g. dead-wood microhabitats, moss 90 fragments), across a wide range of taxonomic groups (e.g. small and arboreal mammals, birds, vascular plants, saproxylic beetles, and micro-arthropods), have either supported (Melo 91 et al., 2017; Rabelo et al., 2017; Seibold et al., 2017) or refuted (Evju & Sverdrup-Thygeson, 92 2016; Haddad et al., 2016; Torrenta & Villard, 2017) the HAH. Therefore, further empirical 93 94 studies are needed to appraise the degree to which the HAH can be generalised to different 95 landscape scenarios and taxonomic groups (Rabelo et al., 2017).

The IBT and HAH were originally developed within a context of oceanic islands 96 (MacArthur & Wilson, 1967) and habitat patches within terrestrial landscapes (Fahrig, 2013), 97 respectively. These two landscape scenarios may be seen as extremes along a continuum. In a 98 global synthesis, Matthews et al. (2016) showed that z-values of SARs are higher in true 99 islands than in habitat patches. They also reported gradients in z-values ranging from inland 100 101 water-body to oceanic islands, and from forest to mountaintop habitat patches. Hence, the magnitude of island effects is context-dependent regarding the type of matrix surrounding 102 habitat patches (Prugh et al., 2008). Patterns of species richness in intermediate landscape 103 scenarios, such as inland water-body islands and mountaintops, could therefore be explained 104 105 by either IBT or HAH.

The HAH was erected under the assumption that species perceive the wider 106 macrohabitat mosaic as functionally connected (Fahrig, 2013). Matrix permeability, as 107 measured by the structural similarity between habitat patches and any surrounding matrix 108 109 (Prevedello & Vieira, 2010), along with inherent differences in species dispersal ability (Lees & Peres, 2009), would then determine whether species use their habitat primarily at the 110 111 patch- or landscape-scale. Accordingly, we hypothesise that patterns of species richness in fragmented landscapes can be better explained under either the HAH if species exhibit high 112 levels of dispersal ability across a permeable matrix, or the IBT if species exhibit low 113 114 dispersal ability across a hostile matrix.

115 Here, we examined whether the HAH can be extended to anthropogenic archipelagic landscapes using the number of understorey insectivorous bird species on forest islands 116 117 induced by a large hydroelectric dam in central Brazilian Amazonia. We focused on understorey insectivorous birds because they are particularly vulnerable to forest loss and 118 fragmentation (Powell et al., 2015), and exhibit low dispersal ability through non-forest 119 matrix habitats (Sekercioglu et al., 2002; Laurance et al., 2004). We show that the number of 120 121 understorey insectivorous bird species on forest islands is best explained by an island effect, 122 which is consistent with the IBT. Moreover, we propose a conceptual framework, based on the degree of matrix permeability and species dispersal ability, to determine which point 123 along the continuum between a patch- and landscape-centric worldview in fragmentation 124 ecology – represented here by either IBT or HAH – provides the most appropriate guiding 125 126 framework for biodiversity studies in fragmented landscapes.

127

## 128 MATERIALS AND METHODS

## 129 Study area

This study was carried out within the Balbina Hydroelectric Reservoir (BHR) in
central Brazilian Amazonia (1°40' S, 59°40' W; Fig. 1). The BHR spans *ca*. 300,000 ha and

- 132 was formed by the damming of the Uatumã River in 1987 (Fearnside, 2016), creating over
- 133 3,500 land-bridge islands of variable size (range = 0.2-4,878 ha), which are surrounded by a
- 134 vast water reservoir often containing dead tree snags rising above the water level (Benchimol
- 135 & Peres, 2015a). To offset the environmental impacts of the dam, the left bank of the former
- 136 Uatumã River, including all islands, became strictly protected by the 940,358-ha Uatumã
- 137 Biological Reserve (IUCN category Ia), the largest of its category in Brazil (Fig. 1).
- 138 The vegetation is comprised primarily of submontane dense ombrophilous (*terra*-
- *firme*) forest, although *igapó* forest subjected to seasonal flooding formerly occurred along
- 140 the margins of the Uatumã River before damming. Forest structure varies among islands due
- 141 to both island size and associated edge-mediated disturbance: smaller islands are species-poor
- and dominated by pioneer tree species, whereas larger islands are species-rich and contain a
- 143 higher dominance of large-seeded canopy tree species (Benchimol & Peres, 2015a).
- 144 According to the Köppen classification, the climate is equatorial fully humid (Af), with mean
- annual precipitation and temperature of 2,464 mm and 26.5 °C, respectively (Alvares et al.,
- **146** 2013).
- 147

# 148 Sampling design

- We selected 33 forest islands within the BHR, ranging in size from 0.63 to 1,698.84 ha. Sixteen islands were on the left bank, whereas 17 islands were on the right bank (Fig. 1). The combined study meta-landscape encompassed 177,720 ha where sample sites were spaced apart by an average distance of 27.9 km (SD = 15.0 km; range = 2.0–68.4 km).
- We surveyed birds using mist nets  $(12 \times 2.5 \text{ m}, Ecotone \ 1016/12)$  from July to 153 December in two consecutive years (2015 and 2016). We placed 16 mist nets end-to-end in 154 the understorey along a continuous near-linear net-line (ca. 200 m) whenever possible, but 155 used a cross-shaped net-line design on islands smaller than 4 ha, thereby ensuring the same 156 sampling effort across all 33 surveyed islands. Herein, each mist-net line corresponds to one 157 sample site. Mist nets were operated from 06:00 to 15:00 h for two days at each site each 158 159 year, resulting in a total sampling effort of 19,008 net-hours (16 mist nets  $\times$  9 hours  $\times$  2 days  $\times$  2 years  $\times$  33 sites). To avoid double-counting, we ringed birds with coded aluminium rings 160 and excluded recaptures. Rings were provided by the Brazilian National Center for Bird 161 162 Conservation and Research (CEMAVE) under research permits SISBIO 49068 and CEMAVE 3984. 163
- 164

# 165 **Response variable and species group**

We used the number of species of understorey insectivorous birds as the response 166 variable, and limited our analysis to forest species because the habitat type (i.e. forest) must 167 be appropriately defined for the focal species group (Fahrig, 2013). We defined forest species 168 as those classified as having 'medium' or 'high' levels of forest dependency (sensu BirdLife 169 International, 2018), and insectivorous species as those classified under the 'invertebrate' 170 171 dietary category (sensu Wilman et al., 2014). The only forest insectivorous species omitted from the analysis was the Amazonian Pygmy-Owl (Glaucidium hardyi) because surveys were 172 diurnal, and this species is nocturnal (Wilman et al., 2014). Since understorey mist nets 173 primarily capture understorey birds and occasionally those that walk on the ground or forage 174 at forest strata higher than 2.5 m (Karr, 1981), we considered all species captured as 175 understorey birds to avoid misinterpretation. 176

177

### 178 **Predictor variables**

We extracted data on *island size* and *habitat amount* for all 33 sample sites using a 179 180 classified image (Collection 2, 2015, Amazon) derived from 30-m resolution LANDSAT imagery downloaded from the Brazilian Annual Land Use and Land Cover Mapping Project 181 182 (available at http://mapbiomas.org). To do so, we used the QGIS software (QGIS Development Team, 2016) and the LecoS plugin (Jung, 2016). Island size corresponds to the 183 total forest area (in hectares) within an island, and habitat amount corresponds to the 184 percentage of forest cover within a given surrounding landscape at varying scales. In 185 extracting the predictor variables, only 'dense forest' (pixel value 3) was defined as forest, 186 187 because other pixel values effectively represent either heavily degraded forests or non-forest land cover types. 188

189

## 190 Data analysis

## 191 Scale of effect

Species-landscape relationships are strongly affected by the scale at which landscape 192 attributes are measured (Jackson & Fahrig, 2015). We therefore employed a multi-scale 193 194 analysis to determine the 'scale of effect' – the landscape scale at which the relationship between the number of species and habitat amount peaks (Jackson & Fahrig, 2015). We 195 defined the scale of effect as the 'local landscape' for understorey insectivorous birds at the 196 Balbina forest archipelago. Our multi-scale analysis examined 40 different buffer sizes 197 around sample sites (i.e. mist-net lines), ranging from 50 to 2,000 m at 50-m intervals. The 198 199 smallest landscape scale (50 m) corresponds to the average between the reluctance of 200 Amazonian understorey birds to cross forest clearings as narrow as 30 m (Laurance et al., 2004) and an assemblage-wide avian gap-crossing ability of up to 70 m (Lees & Peres, 2009). 201

The largest landscape scale (2,000 m) includes those frequently used in avian fragmentation studies (Jackson & Fahrig, 2015; Morante-Filho et al., 2015; Aurélio-Silva et al., 2016). For this analysis, we included all 33 surveyed islands and log-transformed the response and predictor variables ( $\log_{10} x + 1$ ).

- 206
- 207 IBT vs. HAH

The number of species in fragmented landscapes can be explained by either patch size 208 209 (e.g. Torrenta & Villard, 2017) or habitat amount (e.g. Melo et al., 2017), which represent two worldviews in assessing the total area of suitable habitat. This means that the iconic SAR 210 (Rosenzweig, 1995) holds true regardless of its spatial drivers (patch size or habitat amount), 211 but that the underlying mechanism may be either the island effect driven by patch size as 212 213 predicted by the IBT, or the sample area effect driven by habitat amount as predicted by the HAH (Fahrig, 2013). We applied four tests to determine whether the IBT or the HAH is the 214 most appropriate theoretical framework to explain the number of understorey insectivorous 215 216 bird species on forest islands within the BHR.

217

#### 218 Test 1: Multiple linear regression

We used multiple linear regression analysis to examine the independent effects of island size and habitat amount in the local landscape on species richness. This method allows one to estimate how much of the variation in the response variable (i.e. number of species) can be attributed solely to a predictor variable (e.g. island size), once the effects of another predictor (e.g. habitat amount) are controlled for (Legendre & Legendre, 1998).

An effect of island size, rather than one of habitat amount, would provide support for IBT, whereas the reverse would provide support for HAH (Fig. 2). The response and predictor variables were log-transformed ( $\log_{10} x + 1$ ) prior to analysis. The predictor variables were also standardised (mean = 0, SD = 1) to allow comparison of regression slopes. Finally, we examined the strength of correlation values between island size and habitat amount across the entire spectrum of 40 landscape scales (50–2,000 m) to assess the suitability of the multiple linear regression test.

231

232 Test 2: Z-values

We used the logarithmic form of the SAR (type IV curve *sensu* Scheiner, 2003) to fit simple linear regression models (Rosenzweig, 1995) for islands surrounded by either low habitat amounts (up to *ca*. 50% of the landscape; Morante-Filho et al., 2015) or high habitatamounts, according to the following equation:

237  $\log_{10}(S+1) = z \times \log_{10}(A+1) + \log_{10}(c),$ 

where S = number of species, z = regression slope, A = island size, c = regression intercept.

239 To assess whether the *z*-values derived from either SARs were significantly different (p < p

240 0.05), we performed an ANCOVA model with habitat amount (low or high) as an

241 independent categorical variable. To support the IBT, the SAR for islands with low habitat

amounts should have a higher *z*-value than those with high habitat amounts (Fig. 2).

243 Conversely, *z*-values should be statistically equivalent to support the HAH (Fig. 2).

244

245 Test 3: Species accumulation curves

We compared the cumulative number of species on all 33 surveyed islands ordered according to both increasing (small-to-large) and decreasing (large-to-small) island sizes (Quinn & Harrison, 1988), which may lead to three possible outcomes. First, the small-tolarge accumulation curve lies below the large-to-small, supporting IBT (Fig. 2). Second, the curves overlap, supporting HAH (Fig. 2). Third, the small-to-large accumulation curve lies above the large-to-small, supporting neither IBT nor HAH.

252

# 253 Test 4: Extrapolation of SAR

We fit a SAR model (see Test 2: *Z*-values) to all 33 surveyed islands. We further extrapolated the number of species to a hypothetical island containing the area (+ 1) of all 33 surveyed islands combined (7,841.4 ha), and compared the overall number of species observed across surveyed islands with the extrapolated value (Yaacobi et al., 2007). Compared with the extrapolated value, a lower observed number of species would support IBT (Fig. 2); a statistically equivalent number, HAH (Fig. 2); a higher number, neither IBT nor HAH.

261

# 262 **RESULTS**

We captured a total of 874 individual understorey insectivorous birds representing 59 species, 49 genera and 19 families across all 33 sample sites (see Table S1 in Supporting Information). The number of individuals per island ranged from 0 to 84 (26.5  $\pm$  23.4), and the number of species from 0 to 27 (8.8  $\pm$  7.5; see Table S2).

#### 268 Scale of effect

The correlational peak between the number of species and habitat amount (i.e. the 269 scale of effect) occurred at 200-m buffers around sample sites (r = 0.873; see Fig. S1), 270 thereby corresponding to the local landscape for understorey insectivorous birds at the 271 Balbina forest archipelago. The fact that this local landscape is intermediate between the 272 smallest and the largest landscape scales examined here indicates that our multi-scale analysis 273 included the true scale of effect (Jackson & Fahrig, 2015). Therefore, habitat amount is 274 275 defined as the percentage of forest cover only within 200-m buffer local landscapes for all 276 subsequent analyses.

277

## 278 Test 1: Multiple linear regression

A multiple linear regression model including island size and habitat amount showed 279 280 that both predictor variables were strongly and positively related to the number of species ( $R^2$ = 0.80, p < 0.001). Partial regressions also showed that habitat amount had a slightly better fit 281 and higher regression slope than island size (see Fig. S2), which in itself would lend more 282 support for HAH than IBT. However, island size and habitat amount were positively 283 correlated across the entire range of 40 landscape scales examined and peaked exactly at the 284 285 scale of effect (200-m buffer; see Fig. S3). Due to the high collinearity between predictors (r = 0.857), regression coefficients could change depending on the random component in the 286 response variable (Legendre & Legendre, 1998), thereby precluding us from raising any 287 conclusions derived from Test 1. 288

289

290 Test 2: Z-values

The species-area relationship for islands surrounded by low habitat amounts (< 55%) had a statistically higher *z*-value (0.747) than islands surrounded by high habitat amounts (> 70%; 0.311), as shown by an ANCOVA test (p = 0.009; Fig. 3). This outcome supports an island effect, rather than solely a sample area effect (Fahrig, 2013), thereby lending stronger support for IBT.

296

297 Test 3: Species accumulation curves

Species accumulation curves did not overlap whether sampling sites were ordered according to increasing (small-to-large) or decreasing (large-to-small) island sizes. The small-to-large curve lay above the large-to-small curve (Fig. 4), which contradicts both IBT and HAH.

## 303 Test 4: Extrapolation of SAR

304	We observed a larger number of species $(59 + 1)$ across all 33 surveyed islands than
305	that extrapolated (55.2) to a hypothetical island containing the area (+ 1) of all surveyed
306	islands combined (7,841.4 ha). However, the difference between the observed and
307	extrapolated number of species was not significant (Fig. 5), which lends support for HAH.

308

## 309 DISCUSSION

310 Compared to the HAH, there was more evidence giving support to the IBT in explaining the number of understorey insectivorous bird species within Amazonian forest 311 islands in one of the largest hydroelectric reservoirs on Earth. Considering the four tests 312 applied, Test 2 (z-values) was consistent with an island effect as posited by the IBT, Tests 3 313 314 (SACs) and 4 (extrapolation of SAR) were regarded as inappropriate to test the HAH (see 315 below), and Test 1 (multiple linear regression) could not be adequately addressed due to a prohibitively high correlation between island size and habitat amount at the local landscape. 316 Since both an island effect (e.g. Evju & Sverdrup-Thygeson, 2016) and a sample area effect 317 (e.g. Rabelo et al., 2017) may explain patterns of species richness in fragmented landscapes, 318 319 the key question becomes which of these two theoretical frameworks provides the best fit to 320 different scenarios in 'real-world' fragmented landscapes. This question has critical implications to biodiversity conservation strategies since empirical evidence primarily 321 supporting IBT would imply a management focus on the spatial arrangement of remaining 322 habitat patches, whereas support for HAH would imply a management strategy focused on 323 retaining the maximum overall amount of habitat regardless of its configuration (Seibold et 324 al., 2017). 325

The independent effects of predictor variables may be disentangled using statistical 326 methods such as multiple regression analysis. However, as the degree of collinearity between 327 predictor variables increases, the accuracy in determining their independent effects decreases, 328 particularly above a high threshold (r > 0.7) from which parameter estimates begin to be 329 severely distorted in regression-type analyses (Dormann et al., 2013). In our set of sample 330 sites, the highest correlation between island size and habitat amount occurred exactly at the 331 scale of effect (i.e. 200-m buffer; r = 0.857), which precluded us from directly testing the 332 predictions of the HAH. Ideally, patch size and habitat amount should be either orthogonally 333 independent or negatively correlated (Fig. 7 in Fahrig, 2013). However, the pervasive 334 positive correlation between habitat patch size and habitat amount in landscapes worldwide is 335 336 the rule rather than the exception (Fahrig, 2003), and this correlation becomes even stronger and more ubiquitous for smaller local landscapes (Rabelo et al., 2017). For instance, island 337

size and habitat amount were more likely to be independent in our study system at largerscales, well beyond a demographically realistic local landscape for our focal species group.

The scale of effect is indeed unlikely to be known before sampling design is 340 established, thereby a multi-scale analysis is necessary to determine the local landscape 341 (Fahrig, 2013). This implies that sample sites selected a priori to control for the positive 342 correlation between patch size and habitat amount may fail to achieve this goal if the size of 343 the local landscape is different than initially thought. To illustrate this, consider a set of 344 345 sample sites where the size of the focal habitat patches increases while the amount of habitat 346 remains constant (Fig. 6). If the size of the local landscape derived from a multi-scale analysis is found to be half of that defined *a priori*, patch size and habitat amount will be 347 positively correlated (Fig. 6). Therefore, directly testing the HAH under its main assumptions 348 is expected to be less feasible if the scale of effect is small or not known a priori. Despite 349 these shortcomings, there are alternative ways of testing the HAH (Fahrig, 2013). 350

First, if patch size *per se* does not affect the number of species, as predicted by the 351 HAH, z-values derived from species-area relationships are expected to be the same in 352 landscapes with either low or high habitat amounts (Fig. 2). We showed that the z-value for 353 354 islands at landscapes isolated by low habitat amounts (< 55%) is statistically higher than that at landscapes connected by high habitat amounts (> 70%; Fig. 3), which contradicts a 355 prediction of the HAH. Such a difference could be attributed to the selected cut-off (55%) 356 that distinguishes low from high habitat amounts. To test whether the difference in z-values is 357 358 sensitive to this threshold, we reran the *z*-value test using different cut-offs for habitat amount 359 (54%, 50%, 48.5%, 45%, 43.1%, 30%; see Supporting Information). The differences in zvalues held true except when the species-area relationship for islands surrounded by low 360 habitat amounts was not significant. Since all islands were smaller than 10 ha for the lower 361 cut-offs ( $\leq 45\%$ ), the lack of a relationship may be explained by the small island effect (i.e. 362 363 for small islands, the variation in island size does not affect the number of species; Lomolino 364 & Weiser, 2001). In sum, forest islands surrounded by low habitat amounts had a steeper decline in species richness as a function of island size reduction (i.e. higher z-value) than 365 those surrounded by high habitat amounts whenever the SAR models were significant. 366

367 The difference between z-values is attributed to an island effect, which is consistent with the IBT (Fahrig, 2013). Accordingly, larger and less isolated islands are more species-368 369 rich than smaller and more isolated islands because the former experience lower extinction (area effect) and higher colonisation rates (distance effect) (MacArthur & Wilson, 1967). 370 Thus, if islands within their local landscapes were functionally connected as assumed by the 371 372 HAH, the number of immigrants reaching focal islands would mainly depend on the amount 373 of habitat surrounding those islands (sample area effect), thereby compensating species declines through rescue effects (Fahrig, 2013; see Seibold et al., 2017). 374

Second, the species accumulation curves (SACs) from either small to large patches or 375 376 from large to small patches (Fig. 2) should roughly coincide to support HAH since this is caused by a sample area effect, meaning that the long-celebrated dichotomy between a Single 377 Large Or Several Small patches (SLOSS) should harbour a similar number of species. 378 Alternatively, a faster accumulation in the number of species from large to small patches, 379 380 compared to that from small to large patches, would be attributed to an island effect (IBT). The cumulative number of understorey insectivorous bird species at the Balbina forest 381 archipelago rose faster from small to large patches than from large to small ones, which 382 supports neither IBT nor HAH. 383

The fact that several small patches (islands) apparently harboured more species than a 384 single large patch (Fig. 4) is consistent with the literature (Fahrig, 2017). However, we did 385 not find support for several possible explanations for this pattern at the Balbina forest 386 archipelago. First, we strictly focused on forest species implying that the pattern was not 387 confounded by the inclusion of disturbance-adapted species, which would increase the overall 388 number of species across small patch sites (Lovei et al., 2006). Second, habitat heterogeneity, 389 390 regarding vegetation structure in Amazonian terra firme forests, is associated with elevation (Castilho et al., 2006), below-ground vertical distance to the water table (Schietti et al., 2014) 391 and horizontal distance to perennial streams (Drucker et al., 2008). Thus, several small 392 393 patches could harbour more species than a single large patch if they covered wider topographic and hydrologic gradients, resulting in higher levels of habitat heterogeneity 394 395 (Báldi, 2008). However, our islands consist of upland habitat remnants resulting from hilltop 396 terrains of the once continuous forest. As such, they span similar elevations and streams were missing from all but two very large islands (Beco do Catitu and Mascote). As a result, 397 surveyed islands shared relatively low levels of intra-patch habitat heterogeneity regarding 398 closed-canopy forest structure. Third, the Balbina islands are isolated by a hostile water 399 400 matrix which likely hinders the dynamic of colonisation and extinction (Palmeirim et al., 2017), particularly for species that are unable to either cross wide gaps or use dead tree snags 401 as stepping stones. Indeed, the disappearance of understorey insectivorous birds from forest 402 fragments has been largely attributed to dispersal limitation (Sekercioglu et al., 2002), which 403 404 along with a severely hostile water matrix explain patterns of bird species occupancy on 405 forest islands (Moore et al., 2008). The relatively small local landscape threshold (200-m buffer) for understorey insectivorous birds at the Balbina forest archipelago provides 406 407 additional evidence of such dispersal limitation (Jackson & Fahrig, 2012).

The most likely explanation for the observed SACs (Fig. 4) relies on a bias associated with this method. In a SLOSS-type study, Gavish et al. (2012) compared four methods to examine the effects of habitat loss and fragmentation on the species richness of spiders. They concluded that only SACs (Quinn & Harrison, 1988) should be avoided as this method was biased towards detecting more species in several small habitat patches than in a single large patch. This occurs because the method is sensitive to sampling intensity (i.e. proportion of
patch area that is sampled), which could lead to an apparent higher number of species in
small but more intensively sampled patches (Gavish et al., 2012). Since the proportion of the

416 island area we sampled in smaller islands was immensely higher than in larger islands, the

417 result of the SACs is likely to be misleading. Moreover, the pattern of SACs was inconsistent

418 with the HAH even in an experiment designed to decouple the independent effects of patch

- 419 size and habitat amount on saproxylic beetles whose revealed strong support for HAH
- 420 (Seibold et al., 2017).

421 Third, the extrapolation of the SAR model (Yaacobi et al., 2007) suggests that several small islands did *not* harbour more understorey insectivorous bird species than a single large 422 island containing the same aggregate area of several small islands (Fig. 5). This result is 423 presumably consistent with the HAH (Fahrig, 2013; MacDonald et al., 2018). However, had 424 this method been suitable to test the HAH, oceanic archipelagos should harbour fewer species 425 426 than that predicted by the extrapolation of SAR models derived from their constituent islands. Indeed, observed and extrapolated values of species richness for most oceanic archipelagos 427 428 are statistically the same (75% to 95% of 40 case studies; Santos et al., 2010). Collectively, this means that neither SACs nor an extrapolation of SAR models seem to be reliable 429 methods to test the HAH. 430

Testing the HAH is by no means a trivial task for two main reasons. First, as a general 431 rule, habitat patch size and habitat amount tend to be positively correlated (Fahrig 2003). 432 433 However, these two predictors should be either largely orthogonal or negatively correlated to properly test predictions derived from the HAH (Fig. 7 in Fahrig, 2013). Depending on the 434 landscape, this constraint may however be overcome if the scale of effect (sensu Jackson & 435 Fahrig, 2015) is known prior to the establishment of the experimental design. Second, species 436 assemblages are comprised of species with varying degrees of dispersal ability, although 437 438 within some groups, such as understorey insectivorous birds, such a trait is broadly similar across species (Laurance et al., 2004). Thus, the scale of effect for a given species 439 440 assemblage will result from a combination of species with either lower or higher dispersal ability (Lees & Peres, 2009). Therefore, we believe the most robust way forward in testing 441 the HAH would be to focus on individual species (Hanski, 2015) whose dispersal ability 442 443 through the matrix (i.e. landscape vagility) is known a priori and derived from in situ studies (e.g. Awade & Metzger 2008). 444

445

# 446 Moving beyond: a conceptual framework to assess the role of patch size and habitat 447 amount in explaining species responses to habitat fragmentation

We can reasonably expect that local assemblage structure is primarily governed by 449 patch-level characteristics in a hypothetical situation in which species seldom if ever exit the 450 patch, due to low dispersal ability, low matrix permeability, or both (Moore et al., 2008). 451 Conversely, landscape-level characteristics should matter most in a hypothetical situation in 452 which species often move among patches within the local landscape, due to high dispersal 453 ability, high matrix permeability, or both (Walter et al., 2017). Hence, the degree to which 454 either a patch- or landscape-centric worldview is most pertinent in fragmentation ecology 455 studies should be determined by the species vagility within the local landscape, which is 456 largely a combination of matrix permeability (a landscape attribute) and dispersal ability (a 457 species trait) (Fig. 7). Accordingly, increasing support for IBT should be expected for a 458 species assemblage with low dispersal ability in patches surrounded by an impermeable 459 matrix (Fig. 7c; this study; Palmeirim et al., 2017). Conversely, increasing support for HAH 460 would be expected for a species assemblage in which high dispersal ability is prevalent and 461 habitat patches are surrounded by a permeable matrix (Fig. 7b; Melo et al., 2017). Under 462 intermediate scenarios (Fig. 7a and 7d), the most appropriate theoretical framework – IBT or 463 HAH – would depend on the relative contributions of matrix permeability and species 464 dispersal ability. For instance, support for HAH would be expected if species successfully 465 move among patches even if they are surrounded by an impermeable matrix (Fig. 7a; Storck-466 Tonon & Peres, 2017), whereas support for IBT would be expected if species fail to move 467 468 among patches even if they are surrounded by a relatively permeable matrix (Fig. 7d; Munguía-Rosas & Montiel, 2014). 469

470 A recent empirical study testing the HAH (MacDonald et al., 2018) provides further support for our conceptual framework. Accordingly, the inclusion of highly mobile species in 471 the species pool led to stronger support for HAH in explaining the number of butterfly 472 species on islands within a natural archipelagic landscape (Fig. 7a), whereas excluding highly 473 474 mobile species led to stronger support for IBT (Fig. 7c). Our conceptual framework also accounts for dynamic matrix habitats that change over time. As such, for a given forest 475 landscape dominated by a regenerating vegetation matrix that accumulates aboveground 476 phytomass, a patch-centric approach should be gradually replaced by a landscape-centric 477 478 approach as the matrix becomes more permeable, ultimately enhancing species vagility of even the most sedentary species (Powell et al., 2013). The Biological Dynamics of Forest 479 Fragments Project (BDFFP) in central Brazilian Amazonia is an iconic example of a dynamic 480 tropical landscape, in which a cattle pasture matrix surrounding primary forest fragments has 481 been fully replaced by an ageing secondary forest over the past ca. 35 years (Stouffer et al., 482 2011). As the structural contrast between forest fragments and their adjacent matrix 483 484 decreases, forest species can resume movements between forest fragments (Stouffer et al., 2011), exploit newly available matrix resources (Blake & Loiselle, 2001), and incorporate 485 matrix habitats into their territories (Stouffer et al., 2006). In such situation, a dichotomous 486

classification of the landscape into either habitat or non-habitat is at best misleading (Stouffer
et al., 2006), and a landscape-centric approach would be most appropriate.

At the Balbina forest archipelago, the structural contrast between habitat patches 489 (forest islands) and the matrix (open-water) could not be greater, and is aggravated by the fact 490 that matrix recovery, by definition, cannot occur within hydroelectric reservoirs with stable 491 water levels. Such harsh landscape scenario restricts populations of species with low 492 dispersal ability to fewer islands compared to species that can traverse the matrix. Indeed, the 493 494 inherent swimming capacity – a measure of dispersal ability on open-water – of large 495 vertebrate species at the Balbina forest archipelago is positively related to species island occupancy (Benchimol & Peres, 2015b). Based on both patch- and landscape-scale 496 predictors, that study also found island size to be the single best predictor of island occupancy 497 for most species. Likewise, island size was a powerful predictor of species richness of 498 terrestrial and arboreal vertebrates ( $r^2 = 0.910$ , Benchimol & Peres, 2015c), birds ( $r^2 = 0.808$ , 499 Aurélio-Silva et al., 2016), lizards ( $r^2 = 0.870$ , Palmeirim et al., 2017) and frogs ( $r^2 = 0.891$ , 500 Lima et al., 2015) within Amazonian forest archipelagos. Given this bulk of evidence 501 502 showing a strong island size effect on species richness, a patch-centric approach (IBT) is likely to be the most appropriate in true archipelagic landscapes. Nevertheless, species with 503 high dispersal ability (e.g. orchid bees, Storck-Tonon & Peres, 2017; butterflies, MacDonald 504 et al., 2018) may still be able to cross hostile expanses of water, which would justify a 505

- 506 landscape-centric approach (HAH).
- 507

## 508 CONCLUSIONS

509

We tested the habitat amount hypothesis (HAH) under one extreme of the continuum of 510 matrix permeability and species dispersal ability (Fig. 7c) and found stronger support for the 511 512 island biogeography theory (IBT). Meanwhile, we hypothesise that stronger support for HAH is expected under the opposite extreme of this continuum (Fig. 7b), and to either IBT or HAH 513 514 under intermediate scenarios (Fig. 7a and 7d). This notion implies that most species responses to habitat fragmentation lie somewhere along these extremes. Hence, IBT and 515 HAH should not be seen as a mutually exclusive dichotomy, but instead a continuum in 516 517 explaining patterns of species retention in habitat patches. The conceptual framework we propose (Fig. 7) also considers fragmented landscapes with dynamic (e.g. vegetation 518 519 regrowth following land abandonment) or managed matrices (e.g. restored habitats following human intervention). In such landscapes, patch-centric patterns of occupancy (IBT) should 520 gradually transition into those dominated by entire landscapes (HAH) given the role of matrix 521 type in mediating species-area relationships (Freeman et al., 2018). Conversely, matrix 522

- 523 habitat degradation would revert the emphasis back to prime habitat patches. Although
- ameliorating the harshness of water matrices is virtually impossible, other hostile matrix
- habitats, such as bauxite mining (Kennedy & Marra, 2010), can be managed to enhance
- 526 functional connectivity among habitat patches (Fig. 7 in Villard & Metzger, 2014). Finally,
- 527 we conclude that the most appropriate worldview in fragmentation ecology (IBT or HAH) is
- 528 not only context-dependent but also dynamic. Therefore, the best conservation strategy –
- 529 focusing on either the spatial arrangement of remaining habitat patches or the overall habitat
- 530 amount in the landscape is neither static nor can be generalised to a wide spectrum of
- 531 landscape scenarios and taxonomic groups.

## 533 **FIGURE LEGENDS**

- Figure 1. (a) Location of the study area in central Brazilian Amazonia, indicated by a solid
- rectangle containing (b) the Balbina Hydroelectric Reservoir (BHR) landscape, showing the
- 536 boundaries of the Uatumã Biological Reserve, a strictly-protected area safeguarding most of
- this landscape; (c) larger inset map showing the spatial distribution of the 33 surveyed
- islands; and (d) the 200-m buffer area (red polygon) around a mist-net line (white line)
- representing the local landscapes derived for the understorey insectivorous birds examined
- 540 here. Photo credit: Eduardo M. Venticinque.
- 541
- 542 Figure 2. Possible conceptual relationships of the four empirical tests applied to determine
- 543 whether either the island biogeography theory (IBT; graphs on the left) or the habitat amount
- 544 hypothesis (HAH; graphs on the right) is the most appropriate theoretical framework to
- 545 explain the number of understorey insectivorous bird species on forest islands within the
- 546 Balbina Hydroelectric Reservoir in central Brazilian Amazonia.

547

- 548 Figure 3. Divergent species-area relationships for understorey insectivorous birds surveyed at
- 549 33 islands surrounded by either low ( $r^2 = 0.598$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, n = 15) or high ( $r^2 = 0.547$ , p < 0.001, r = 0.547, p < 0.001, r = 0.00
- 550 0.001, n = 18) habitat amounts. Circle sizes scale to the landscape-scale habitat amount, with
- smaller and larger circles representing islands surrounded by either low or high habitat
- amounts, respectively. Note the different *z*-values for these two landscape scenarios (p =
- 553 0.009) and the base 10 logarithmic scales along both axes.

554

- Figure 4. Species accumulation curves of understorey insectivorous birds for islands ordered
  according to either increasing (light grey circles, dashed line) or decreasing (dark grey
  circles, solid line) island size.
- 558
- 559 Figure 5. Species-area relationship for understorey insectivorous birds surveyed at 33 islands.
- 560 The white circle shows the extrapolated number of species (55.2) to a hypothetical island
- containing the area (+ 1) of all 33 surveyed islands combined (7,841.4 ha), whereas the black
- 562 circle shows the total number of species observed in this study (59 + 1). Dashed lines show
- the 95% confidence intervals of the predicted line. Note the base 10 logarithmic scales alongboth axes.

- 566 Figure 6. Sampling design established to control for the positive correlation between patch
- size and habitat amount. The solid black circle correspondents to the local landscape defined
- 568 *a priori* (i.e. before the scale of effect is known). The dashed black circle corresponds to the
- local landscape derived from a multi-scale analysis (i.e. post data analysis). The difference
- 570 between the two landscapes scales (solid and dashed black circles) implies that even a well-
- 571 designed study may fail to control for the collinearity between predictors. Figure modified
- 572 from Fahrig (2013).
- 573
- 574 Figure 7. Conceptual framework based on both the degree of matrix permeability and species
- 575 dispersal ability in determining whether the island biogeography theory (IBT) or the habitat
- amount hypothesis (HAH) is the most appropriate guiding theoretical framework for
- 577 biodiversity studies in fragmented landscapes.







582

583 Figure 2



585 Figure 3





591 Figure 5



594 Figure 6





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  2412.
- 768

## 769 **BIOSKETCHES**

- 770 Anderson Saldanha Bueno is a PhD student at the School of Environmental Sciences,
- 771 University of East Anglia, UK. Since 2008, he has been working on avian community
- ecology in Brazilian Amazonia, focusing on species distributions along environmental
- gradients. His doctoral research seeks to understand patterns of species assembly and
- extinction risk in experimental island landscapes.
- 775 **Carlos A. Peres** is a Brazilian-born Professor of Tropical Conservation Biology at the
- University of East Anglia with 35 years of field and modelling experience on a wide
- spectrum of research topics on Neotropical forest wildlife ecology and conservation. Much of
- his research program focuses on the disturbance ecology of Amazonian forests.
- 779
- Author contributions: A.S.B. and C.A.P. conceived the ideas; A.S.B. collected the field data;
- A.S.B. analysed the data; and A.S.B. wrote the manuscript with substantial inputs from
- 782 C.A.P.