

1 LRH: Mayhew *et al.*

2 RRH: BIRD COMMUNITIES IN SECONDARY FOREST

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6 **Connectivity with primary forest determines the value of secondary tropical forests for**
7 **bird conservation**

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24 **ABSTRACT**

25

26 Species extinctions caused by the destruction and degradation of tropical primary forest may be
27 at least partially mitigated by the expansion of regenerating secondary forest. However, the
28 conservation value of secondary forest remains controversial, and potentially underestimated,
29 since most previous studies have focused on young, single-aged, or isolated stands. Here we use
30 point count surveys to compare tropical forest bird communities in 20–120-yr-old secondary
31 forest with primary forest stands in central Panama, with varying connectivity between
32 secondary forest sites and extensive primary forest. We found that species richness and other
33 metrics of ecological diversity, as well as the combined population density of all birds, reached a
34 peak in younger (20-yr-old) secondary forests, and appeared to decline in older secondary forest
35 stands. This counter-intuitive result can be explained by the greater connectivity between
36 younger secondary forests and extensive primary forests at our study site, compared with older
37 secondary forests that are either (1) more isolated, or (2) connected to primary forests that are
38 themselves small and isolated. Our results suggest that connectivity with extensive primary
39 forest is a more important determinant of avian species richness and community structure than
40 forest age, and highlight the vital contribution secondary forests can make in conserving tropical
41 bird diversity, so long as extensive primary habitats are adjacent and spatially connected.

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43

44 **ABSTRACT IN SPANISH**

45

46 La extinción de especies causada por la destrucción y la degradación del bosque tropical
47 primario puede ser mitigada, en cierta parte, por la expansión del bosque secundario. Sin
48 embargo, la importancia del bosque secundario para la conservación es un tema controversial y
49 potencialmente subestimado ya que la mayoría de los estudios previos se han centrado en
50 bosques muy jóvenes de una sola edad, o aislados de bosques con diferentes edades. En este
51 estudio utilizamos puntos de conteo para comparar las comunidades de aves en bosques
52 secundarios de 20 a 120 años de edad y en bosques primarios en Panamá, y con conectividad
53 variable entre los bosques secundarios y los bosques primarios. Encontramos que la riqueza de
54 especies, y otras métricas de la diversidad ecológica, alcanzaron su máximo en los bosques
55 secundarios más jóvenes (20 años de edad) y parecieron disminuir en los bosques secundarios
56 más antiguos. Este resultado aparentemente contradictorio se puede explicar por la mayor
57 conectividad entre los bosques secundarios más jóvenes y los bosques primarios de gran
58 extensión, y porque los bosques secundarios más antiguos están (1) más aislados o (2) están
59 conectados a bosques primarios aislados y de menor extensión. Nuestros resultados indican que
60 la riqueza de especies y la estructura de la comunidad de aves dependen más de la conectividad
61 entre bosques primarios extensos que de la edad del bosque. Así mismo, nuestros resultados
62 resaltan la importancia de los bosques secundarios para la conservación de las aves tropicales,
63 especialmente estos son contiguos y están conectados a bosques primarios extensos.

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- 66 **Key words:** Bird communities; community structure; conservation; landscape management;
- 67 land-use change; tropical extinction crisis; secondary forest; Panama

68 BETWEEN 2010 AND 2015 THERE WAS AN ANNUAL LOSS OF APPROXIMATELY 7.6 MILLION HA of
69 forest globally, with most of this deforestation occurring in the tropics (FAO 2015). Most of the
70 world's biodiversity is found in the tropics (Dirzo & Raven 2003), and the continued loss and
71 degradation of tropical forests is likely to cause mass species extinctions (Dent & Wright 2009,
72 Wright & Muller-Landau 2006). It has been proposed that primary forest (PF) losses may be
73 offset by the planting and natural regeneration of secondary forests (SF) on previously deforested
74 land (Wright 2005). However, the long-term conservation value of SF depends on whether these
75 habitats can maintain similar species composition and ecosystem functions as PF (Chazdon *et al.*
76 2009, Dent & Wright 2009).

77 Many studies that assess the conservation value of tropical SF have focused on birds, one
78 of the best studied faunal groups in the tropics (e.g. Barlow *et al.* 2007b, Lees & Peres 2006,
79 Robinson 1999, Stotz *et al.* 1996, Willis 1974). Birds provide important ecosystem services, such
80 as pollination and seed dispersal (Sekercioglu 2006, Whelan *et al.* 2008), and their diverse
81 habitat and dietary requirements often lead to species-specific responses to habitat disturbance
82 (Hughes *et al.* 2002, Petit & Petit 2003). Birds are also convenient indicators for ecological
83 assessments because they are relatively easy to identify and survey.

84 Studies comparing avian species richness and community structure in SF and PF report
85 inconsistent results. Some studies have found equivalent or higher species richness in SF
86 compared to PF (Andrade & Rubio-Torgler 1994, Blake & Loiselle 2001, Borges 2007, O'Dea
87 & Whittaker 2007, Schulze & Waltert 2004), while other studies report reduced species richness
88 in SF (Barlow *et al.* 2007b, Bowman *et al.* 1990, Gibson *et al.* 2011, Terborgh & Weske 1969,
89 Tvardíková 2010). These conflicting results may stem from three key factors: the age of SF
90 studied, the landscape context, and the responses of different avian groups to habitat change.

91 In terms of SF age, most studies examining avian diversity in tropical SF have only
92 included relatively young, and single-aged stands (less than 35 yr; e.g. Barlow *et al.* 2007b,
93 Blake & Loiselle 2001, Borges 2007, Terborgh & Weske 1969). With increasing forest age, SF
94 typically develops greater structural complexity, resembling PF over time (Guariguata &
95 Ostertag 2001, Pena-Claros 2003). More complex forest structure offers an increased breadth of
96 ecological niches for forest birds (DeWalt *et al.* 2003, Zahawi *et al.* 2015). Thus, the structural
97 complexity that develops over SF succession should provide habitats for increasingly diverse and
98 complex bird communities (MacArthur & MacArthur 1961, Casas *et al.* 2016). Studies focusing
99 on young SF may therefore underestimate the longer-term value of SF for bird conservation.
100 While estimates of avian species richness in young SF are often inflated by non-forest species,
101 species composition in young tropical SF tends to differ from PF (Barlow *et al.* 2007a, Borges
102 2007, Tvardíková 2010). With increasing time since abandonment both forest structure and bird
103 communities become more similar to those of PF (Andrade & Rubio-Torgler 1994, Borges 2007,
104 Raman 1998). Thus estimates of conservation value need to consider the recovery of forest
105 species composition and abundance rather than richness alone.

106 The landscape context of SF, defined by connectivity to PF source populations and
107 isolation within the countryside matrix, plays a critical role in determining avian community
108 reassembly (Chazdon *et al.* 2009, Dent & Wright 2009, Wolfe *et al.* 2015). Many tropical forest
109 birds are highly dispersal limited with poor gap-crossing abilities, and may not be able to
110 colonise SF unless it is contiguous with PF (Van Houtan *et al.* 2007, Lees & Peres 2009, Moore
111 *et al.* 2008, Tobias *et al.* 2013). In addition, bird species occurring in isolated SF embedded
112 within a non-forest matrix may be more sensitive to random population fluctuations and local
113 extinction. Connectivity to PF is an important factor in SF recovery, and the species composition

114 of bird communities in isolated SF may never fully converge with that of PF (Jones *et al.* 2016,
115 Wolfe *et al.* 2015).

116 Finally, the response of bird species to forest succession may be mediated by their degree
117 of ecological specialisation. It has been suggested that generalist, migratory or forest-edge
118 species proliferate in SF as their wider niche breadth makes them better adapted to the conditions
119 found in younger forest (Barlow *et al.* 2007b, Stotz *et al.* 1996). By contrast, forest specialists
120 are likely to require foraging and nesting resources only found in more mature forest (Barlow *et*
121 *al.* 2007b, DeWalt *et al.* 2003). Forest isolation has also been shown to adversely affect forest-
122 dependent, understorey insectivore species more severely than other functional groups (Bradfer-
123 Lawrence *et al.* 2018, Barlow *et al.* 2006, Ferraz *et al.* 2007, Stouffer *et al.* 2006). Thus, the
124 conservation value of SF for birds is affected by species-specific responses mediated by both site
125 and landscape factors, including habitat age, and level of isolation and connectivity to PF.

126 Here, we examine the species richness and composition of bird communities in central
127 Panama across the longest SF chronosequence studied to date, spanning forest ages from 20 to
128 120 yr, as well as PF controls (see Fig. 1). Across this age gradient, we sampled forests that were
129 either isolated from or connected to extensive PF. This landscape presents an opportunity to
130 examine how bird communities change across both successional and isolation gradients, and to
131 investigate the relative importance of forest age versus isolation in determining the conservation
132 value of SF. Including forest age and landscape context introduces a new level of complexity to
133 classical forest fragmentation studies, which arguably reflects the reality of most human-
134 modified tropical forest landscapes.

135 In this context, we assessed the relative role of secondary forest age versus connectivity
136 with primary forest in determining bird diversity – estimated as (1) bird species richness and

137 other diversity metrics, (2) bird population density, and (3) the similarity of avian community
138 composition to PF. In all cases, we examined the extent to which variation in bird communities is
139 mediated by landscape context, such as isolation by water barriers.

140

141 **METHODS**

142

143 **STUDY SITES.** – We conducted field surveys in the Panama Canal Watershed, where vegetation is
144 classified as tropical moist forest (Holdridge & Budowski 1956). The climate is seasonal with a
145 distinct dry season, typically from mid-December until early May, and annual rainfall of 1900 –
146 3600mm (Windsor, 1990; Turner et al. 2015). We selected study sites in the Barro Colorado
147 Nature Monument, Soberania National Park and the adjacent Agua Salud Project (Fig. 1). The
148 Barro Colorado Nature Monument (5,600 ha; 9°9' N, 79°51' W) is comprised of five peninsulas
149 and Barro Colorado Island (BCI), all situated in Lake Gatun, which was formed in 1914 by the
150 flooding of the Panama Canal. The Barro Colorado Nature Monument is a mosaic of PF mixed
151 with SF stands of different ages that were used for cattle pasture or fruit production between the
152 1880s and the establishment of the park in 1979 (Leigh *et al.* 1982). Soberania National Park
153 (22,000 ha; 9°9' N, 79°44' W) was established in 1980 and is a mix of PF and very old SF (Van
154 Bael *et al.* 2013). The Agua Salud Project research site (664 ha; 9°13' N, 79°47' W) was once
155 predominantly cattle pasture or small-scale shifting cultivation but, after establishment in 2008,
156 the landscape is now predominantly SF of relatively young age (Van Breugel *et al.* 2013). The
157 difference in annual rainfall between our northernmost and southernmost sites (separated by a
158 latitudinal distance of 9.8 km) is 159 mm pa (Rompre *et al.* 2007). As this variation is minor

159 compared to a difference of 2100 mm pa across the full rainfall gradient in Central Panama
160 (Rompre *et al.* 2007), we treated the study area as a single climatic band (see Fig. 1).
161
162 SITE SELECTION. – We selected secondary forest sites along a chronosequence of approximately
163 20, 40, 60, 90 and 120 yr since abandonment, with two replicates per forest age. We selected
164 sites within existing research areas having data on long-term vegetation dynamics, tree
165 communities and accurate age estimates compiled from historical records, aerial photographs and
166 interviews with residents; for details see Denslow & Guzman (2000) and van Breugel *et al.*
167 (2013). The youngest SF in the Barro Colorado Nature Monument is 40-yr-old, while SF in
168 Agua Salud is 10–34 yr old (mean = 19 yr old). For ease of presentation, we refer to Agua Salud
169 sites as 20-yr-old. We selected four PF sites: two in a relatively small patch (c. 800 ha) of
170 isolated PF on Barro Colorado Island (henceforth referred to as isolated PF) and two in an
171 extensive area of mainland PF in Soberania National Park (c. 22,000 ha; henceforth referred to as
172 extensive PF). The PF are at least 500 yr old and there is no indication that they have ever been
173 logged or cultivated (Piperno 1990). There is no ongoing disturbance (such as logging or
174 hunting) in Barro Colorado Nature Monument, whereas in Agua Salud there may be some forest
175 clearance and disturbance in the wider landscape. Across the Barro Colorado Nature Monument
176 chronosequence, average canopy height and structural complexity increases with SF age
177 (DeWalt *et al.* 2003, Mascaro *et al.* 2012). Further details of vegetation structure and
178 composition are available in Dent *et al.* (2013), DeWalt *et al.* (2003), and Mascaro *et al.* (2012).

179 Habitat patch size is an important determinant of species' persistence in fragmented
180 landscapes (Bender *et al.* 1998). However, the importance of patch size relates to the
181 composition of the surrounding matrix. The SF sites in our study are embedded within a mixed-

182 age forest matrix, which buffers the effects of fragment size and limits our ability to accurately
183 calculate areas of single-aged fragments. The three forest areas in which study sites are
184 embedded include Barro Colorado Island (1,560 ha), Gigante peninsulas (2,600 ha), and
185 Soberania National Park and surrounding contiguous forest (22,000 ha; see Fig. 1 for details).
186 The SF and PF sites in this study experience different connectivity. The 20-yr-old Agua Salud SF
187 sites form part of a large forest network connected to extensive PF in Soberania National Park,
188 while both island and peninsula SF sites are smaller, isolated areas of forest surrounded by water.
189 Island SF sites (90–120 yr old) are connected only to isolated PF, and are separated from
190 extensive mainland PF by water. Secondary forest on the Gigante Peninsula is more extensive,
191 and contains older patches (>200 yr old) interspersed with patches of 40–60 yr old SF, but is
192 separated from extensive PF by either water or agriculture. We sampled PF sites on both island
193 and mainland settings to examine the effects of different types and extents of forest isolation, and
194 to provide a baseline for studying the effects of SF age on bird communities. Due to the
195 restrictions of available PF and SF in the study landscape, it was not possible to replicate within
196 categories (e.g., isolation type x isolation extent x forest type), and even where replicates were
197 possible, the sampling design is weakened because some sites are embedded within the same
198 geographical feature (e.g. BCI) and therefore to some extent non-independent. We take these
199 factors into account in our analyses (see below) and emphasise that the study landscape has
200 distinct advantages—not least the comparison across different levels of isolation, and the
201 availability of background data on the history of forest regeneration—which provide a unique
202 opportunity to understand secondary forests in a spatial and temporal context.
203

204 BIRD SURVEY METHODS. - At each of the 14 sites, we established nine point counts with each
205 point separated by a minimum of 100 m from other points, and by at least 50 m from forest of a
206 different age (Van Bael *et al.* 2013, Robinson *et al.* 2000). Two trained observers surveyed one
207 site per morning, with the first count beginning ten mins before sunrise and the last completed by
208 10:30 h. All nine stations at a site were sampled once during a survey visit, with a minimum of
209 three days between surveys; no surveys were conducted on days with heavy rain or strong wind
210 because these limit bird activity and detectability.

211 Point counts were 10 mins in duration, and all birds seen or heard within a 50 m-radius
212 were identified, following previous studies (De Bonilla *et al.* 2012, Martin & Blackburn 2014,
213 O’Dea & Whittaker 2007, Raman & Sukumar 2002). Limiting counts to a 50 m radius can help
214 to reduce the differences in detectability of birds among habitat types due to vegetation structure,
215 and minimises biases and errors in species identification and distance estimates (Petit *et al.*
216 1995). For each bird seen or heard, observers used a laser rangefinder to estimate the Euclidean
217 distance from the centre of the point count to the bird (Buckland *et al.* 2008). Distance estimates
218 to birds detected only by ear are likely to be less consistent than estimates based on visual
219 detections, but in most cases the location of calling birds can be judged reasonably accurately.
220 Birds flying above the canopy were excluded from the survey. Along with the point count data,
221 we kept a list of additional species encountered as we walked between the point count stations
222 during a survey. We conducted surveys over three years: July to October 2014 (wet season),
223 January 2015 and January to March 2016 (dry season). Each site was surveyed a total of ten
224 times over the three years; five times in the wet season and five times in the dry, giving a total of
225 1,260 point counts.

226 Observers had considerable ornithological field experience, including in tropical forest
227 habitats. Two observers were Panamanian, with many years' experience identifying local
228 avifauna. All observers received training before data collection began, including detection tests
229 to check for any bias in identification ability and for consistency in estimations of distance.
230 Recordings of calls and songs were used intensively to improve identification skills and check
231 identifications based on vocalisations.

232

233 DATA ANALYSIS. - Prior to analysis, we removed unidentified birds from the dataset (1.8% of
234 total number of detections). We conducted all analyses both on the remaining bird species
235 (henceforth, all birds), and on a dataset restricted to birds with a higher dependency on forest
236 habitats (henceforth, forest specialists). We defined forest specialists as species characteristic of
237 the interior of undisturbed forest, breeding almost invariably within forests, occurring less often
238 away from forest interior and rarely seen in non-forest habitats, even though they may persist in
239 secondary forest and forest patches if their particular ecological requirements are met (BirdLife
240 International 2018, Buchanan *et al.* 2011). We note that classification of forest dependency in
241 birds is potentially subjective, partly because species vary in their habitat selection
242 geographically. We used the most recent classification of forest specialism (BirdLife
243 International 2018) because it is global in focus, comprehensive and widely accessible. We
244 found results to be very similar when we used alternative, geographically restricted
245 classifications of forest dependency, including published descriptions by Ridgely & Gwynne
246 (1989), habitat codes of Stotz *et al.* (1996), and habitat scores of Tobias *et al.* (2016).

247 We calculated rarefaction curves to compare rates of species accumulation among forest
248 age classes for all birds and forest specialists. When scaled by the number of samples, curves

249 reached or approached the asymptote for all forest ages and species sets, suggesting survey effort
250 was adequate (Fig. S1 and S2). However, curves did not reach asymptotes for some forest ages
251 and species sets when scaled by individuals, suggesting some sites were under-sampled (Fig. S1
252 and S2). To identify species that were missing from the extensive PF dataset, we compared our
253 dataset to the species list reported in a previous survey of the same extensive PF forest (Robinson
254 *et al.* 2000). This long-term study used intensive survey methods to describe the species
255 composition of the extensive PF site and so provides a complete picture of the species present at
256 this locality.

257 We calculated species richness and the percentage of PF species present in SF by
258 combining both the point count data and the additional species encounters. All other analyses
259 used data from point counts only. We conducted analyses using R (Version 3.4.1, R Core Team
260 2017).

261
262 SPECIES RICHNESS, DIVERSITY AND DOMINANCE. - We compared species richness, Shannon-
263 Weiner diversity indices and dominance across forest ages using data from all surveys combined.
264 We calculated dominance as the percentage of individual birds represented by the five most
265 common species in each site.

266
267 BIRD POPULATION DENSITY. - We used the R package ‘Distance’ (Laake *et al.* 2015) to estimate
268 bird community population density among forest ages pooled over the 1,260 point counts,
269 following methods described in Buckland *et al.* (2015). We pooled visual and audial detections,
270 and stratified analyses by forest age to allow for differences in detectability among habitats.
271 Using the function ‘ds’ (‘Distance’ R package; Laake *et al.* (2015), we fitted 36 detection

272 functions with various combinations of covariates (year, season, detection method and observer)
273 per forest age and used AIC model selection to choose the best-fit models (Burnham *et al.* 2011).
274 The detection functions provided an estimation of bird population density (number of individuals
275 per hectare) in each of the forest ages. The function ‘ds’ requires a minimum of 80 observations
276 within a category to give reliable estimates per species, thus we did not calculate detection
277 functions for individual species, since only 4–9 bird species in each forest age category had more
278 than 80 detections. Our results should be interpreted with caution since pooling community
279 detectability data assumes that each species is equally detectable across each of the 14 sites.

280

281 SPECIES COMPOSITION AND SIMILARITY TO PRIMARY FOREST. – We calculated the percentage of
282 bird species detected in PF that were also detected in SF separately for isolated PF and extensive
283 PF sites by pooling data for each forest age category.

284 We used the Morisita-Horn abundance-based similarity index (S_{MH}) to compare species
285 composition between pairs of assemblages. The S_{MH} is robust to uneven and insufficient
286 sampling and thus suited to determine if reassembly of PF communities occurs in SF in terms of
287 relative abundance (Chao *et al.* 2006). We examined whether species composition of SF
288 converged with either isolated PF or extensive PF over time by comparing the similarity in
289 composition (S_{MH}) of each SF forest site to each of the PF sites. We examined similarity to
290 isolated and extensive PF sites separately because isolation-related extirpations have altered the
291 island bird communities (Robinson 1999). Similarity values were produced using the function
292 ‘vegdist’ (‘vegan’ R package; Oksanen *et al.*, 2016).

293 To determine if forest age or geographic location explained patterns in species
294 composition across sites, we performed Mantel tests on three matrices of pair-wise distances

295 among sites: Euclidean geographic distance, difference in forest age, and dissimilarity in species
296 composition ($1 - S_{MH}$). We assigned PF sites a nominal age of 500 yr to include these sites in the
297 distance matrix for forest age. Mantel tests were performed using the function ‘mantel’ (‘vegan’
298 R package; Oksanen et al., 2016).

299 We explored qualitative similarities in species composition among sites with non-metric
300 multidimensional scaling (NMDS; Anderson *et al.* 2011). This approach uses rank order, rather
301 than absolute abundances of species, to represent the original position of communities in
302 multidimensional space as accurately as possible using a reduced number of dimensions. We
303 used similarity matrices generated from both the S_{MH} abundance-based and Jaccard incidence-
304 based similarity values (S_J). We included the S_J similarity values to investigate whether PF
305 species were present in SF, even if patterns of relative abundance were different from those in
306 PF. Ordinations were performed using the function ‘metaMDS’ (‘vegan’ R package; Oksanen et
307 al., 2016).

308 To assess the significance of observed differences in species composition as related to SF
309 age, isolation level (isolated or connected), forest type (SF or PF) and distance to extensive PF,
310 we conducted a series of permutational MANOVAs, an analysis of variance using distance
311 matrices. This analysis uses pseudo-F values to compare among-group to within-group similarity
312 and assesses significance by permutation. We also investigated the effect of season (wet or dry)
313 on species composition by conducting a permutational MANOVA at survey level. Permutational
314 MANOVAs were produced using the function ‘adonis’ (‘vegan’ R package; Oksanen et al.,
315 2016).

316 We calculated the mean number of migratory bird detections in different forest age
317 categories based on count data with no distance corrections. This gives a relative abundance of

318 migratory birds in habitats for those species with similar detection probabilities. We also used
319 the point count data to list the five most abundant species per forest age, and classified these
320 species using diet and habitat information from Ridgely & Gwynne (1989) and Wilman *et al.*
321 (2014).

322

323 **RESULTS**

324

325 Our surveys recorded a total of 183 bird species from 42 families, of which 55 species from 24
326 families were forest specialists (Table S1). We detected 13,894 individual birds in fixed radius
327 point counts, of which 5,256 were forest specialists (BirdLife International 2018).

328

329 PATTERNS OF SPECIES RICHNESS, DIVERSITY, AND DOMINANCE. - No clear relationship was found
330 between species richness and forest age (Table 1). The youngest SF (20-yr-old) had higher
331 species richness than all other sites, and a species richness of forest specialists similar to
332 extensive PF (Table 1). The oldest SF (120-yr-old) had the lowest species richness for all birds
333 and forest specialists. These counter-intuitive patterns of species richness appear to be influenced
334 by differences in connectivity among sites, with higher species richness found in sites that were
335 connected to extensive PF (Fig. 2). Compared with extensive mainland PF sites, the isolated PF
336 sites had lower species richness for both datasets. Species diversity (Shannon-Weiner index)
337 showed similar patterns across sites, while dominance values were highest in isolated sites and
338 lowest in connected sites (Table 1).

339

340 BIRD POPULATION DENSITY. - There was no clear pattern in bird community population density
341 estimates across the different forest ages or levels of isolation. For all birds, the 20-yr-old SF had
342 the greatest density of birds, estimated at 29 individual birds/ha (95% CI: 26, 31; Fig. 3). This
343 compares with the lowest density estimate of 17 individual birds/ha (95% CI: 16, 19) in the 120-
344 yr-old SF. Qualitatively similar patterns were found for forest specialists.

345

346 SIMILARITY TO PRIMARY FOREST. – Focusing on all birds, we found no clear relationship between
347 SF age and the percentage of PF bird species detected in SF sites (as estimated by our surveys),
348 but there was a relationship between isolation and percentage of PF species present in SF.
349 Percentage of PF species present was consistently highest in connected sites, and lower in
350 isolated sites. When comparing SF ages, we found that the highest percentage of PF species
351 occurred in the 20-yr-old connected SF (86% when compared with extensive PF sites as
352 estimated by our surveys; Fig. 4). Surprisingly, the 120-yr-old isolated SF had the lowest
353 percentage of PF species present, with only 72 percent in common with isolated PF and 57% in
354 common with extensive PF. This is likely due to shifts in species richness driven by isolation
355 effects in the island PF, where we detected just 62% of the species that we found in extensive
356 mainland PF.

357 For all birds, compositional similarity to isolated (but not extensive) PF increased with
358 forest age (Fig. 5). The highest similarity in species composition between PF and SF was
359 recorded on BCI where the community composition of the oldest isolated SF (120-yr-old; $n = 2$
360 sites) was very similar to isolated PF ($n = 2$ sites; similarity index [S_{MH}]: 0.87 ± 0.03). By
361 contrast, the lowest similarity was between the 20-yr-old SF and the isolated PF (0.58 ± 0.03);
362 these sites span the widest range in both isolation level (mainland vs. island) and age (20-yr vs.

363 PF). Similarity was higher between extensive PF sites and 20-yr-old SF (0.69 ± 0.03) than
364 between extensive PF sites and the older, isolated 120-yr-old SF (0.61 ± 0.04). Mantel tests
365 indicated that geographic distance among sites ($R^2 = 0.74$, $P = < 0.001$) explained a greater
366 proportion of variation in species composition than forest age ($R^2 = 0.30$, $P = < 0.05$). Similar
367 patterns were found for forest specialist species, except forest age was not a significant predictor
368 of species composition (geographic distance: $R^2 = 0.66$, $P = < 0.01$; forest age: $R^2 = 0.21$, $P =$
369 0.06).

370 The NMDS of abundance-based species composition (S_{MH}) across all bird species
371 showed a separation of sites in relation to both forest age and isolation level (Fig. 6). Sites
372 displayed a clear split along Axis 1 that related to site location (connected or isolated), while the
373 age of forest sites tended to increase along Axis 2. The NMDS comparisons for forest specialists
374 showed very similar patterns as those seen for all birds, as did the NMDS results for both
375 datasets using S_J , although the effect of forest age became less apparent when restricting analyses
376 to species presence/absence data (Fig. 6).

377 The permutational MANOVA using S_{MH} indicated that forest isolation level explained a
378 greater portion of the variation in community composition of all bird species than forest age or
379 forest type (SF vs PF; Table 2). Distance to extensive PF was not a significant predictor of
380 community composition. The permutational MANOVA using S_J for all birds showed very
381 similar results (Table 2). Season had a significant effect on community composition, but it did
382 not change the patterns observed for forest isolation, forest age or forest type, although distance
383 to extensive PF became significant (Table 2). Community composition of forest dependent
384 species (using S_{MH} or S_J) was largely dictated by forest connectivity. The same patterns were
385 found for forest specialists as the all-bird dataset when season was taken in to account (Table 2).

386

387 COMPOSITIONAL CHANGES. - The composition of the five most abundant bird species differed
388 across forest ages (Table S2). Only one species, Black-crowned Antshrike (*Thamnophilus*
389 *atrinucha*), was consistently abundant across all sites. Southern Bentbill (*Oncostoma olivaceum*)
390 was among the top five most abundant species in the youngest forest sites (20, 40 and 60-yr-old
391 SF), while Red-lored Amazon (*Amazona autumnalis*) appeared in the top five for both the
392 isolated and extensive PF, as well as the 90-yr-old SF. The five most abundant species in the
393 isolated and extensive PF, and the 90-yr-old SF exhibited a greater diversity of feeding guilds
394 and foraging strata than those found in the younger SF sites (Table S2).

395 We detected 15 species in isolated PF that we did not see in extensive PF, including
396 species such as Crested Guan (*Penelope purpurascens*) that are susceptible to hunting and
397 therefore extirpated from most mainland localities. By contrast, 44 species were detected in
398 extensive PF that were not seen in isolated PF. These were predominantly understory,
399 insectivorous species, many of which have become extinct on BCI since its isolation (Robinson
400 1999, Willis 1974).

401 The number of migratory birds detected per point count was highest in the 20-yr-old SF
402 (0.533 ± 0.091) and declined with increasing SF age to only 0.078 ± 0.032 migrants detected per
403 point count in the 120-yr-old SF (Fig. S3). The number of migrants detected in PF sites was
404 about half the number detected in the 20-yr-old SF (island PF: 0.27 ± 0.07 ; mainland PF: $0.29 \pm$
405 0.06).

406

407

408 **DISCUSSION**

409

410 Our survey data, sampled across tropical SF of varying ages and isolation levels revealed that
411 variation in avian species richness was best explained by connectivity to extensive PF, rather
412 than forest age. We found the highest species richness in the youngest SF sites, which were
413 adjacent to extensive PF. Similarly, connectivity, rather than the forest age, was the strongest
414 predictor of community composition. Finally, the highest bird population density was also found
415 in younger SF, although broader patterns in density did not appear to be driven by either forest
416 age or connectivity.

417

418 SPECIES RICHNESS, DIVERSITY AND DOMINANCE. - High species richness and abundance of birds
419 in SF or successional areas has been documented in many studies (Blake & Loiselle 2001, Johns
420 1991, Karr 1976, Petit & Petit 2003), supporting the hypothesis that intermediate levels of
421 disturbance may lead to high species richness (Connell 1978). In general, species richness in
422 younger SF is boosted by an influx of non-forest, open habitat and generalist species, although it
423 also may contain an important component of forest species (Barlow *et al.* 2007b, Dunn &
424 Romdal 2005). Most studies report that species richness and community structure of tropical
425 secondary forests progressively approaches that of PF over time, and tends to track the
426 increasing structural complexity of secondary forests (Raman 1998, Dent & Wright 2009). In our
427 study, however, species richness and abundance did not increase with forest age, and were
428 instead highest in the youngest SF even when non-forest bird species were removed. We also
429 found that species richness and abundance both increased with greater connectivity to extensive
430 PF, suggesting that high species richness in the 20-yr-old SF is driven by proximity to extensive
431 PF in adjacent Soberania National Park.

432 Proximity to undisturbed habitats has been shown to increase the diversity of bird
433 communities in degraded sites (Johns 1991, Terborgh & Weske 1969, Waltert *et al.* 2004). In La
434 Selva, Costa Rica, PF was the primary habitat and source population for many of the bird species
435 found in SF (Blake & Loiselle 2001). This pattern is supported by our findings, in which a
436 greater number of forest specialists were found in well-connected 20-yr-old SF, than in isolated
437 PF. In a landscape of mixed ages of SF and varying connectivity among forest patches, our
438 findings suggest that the key factor determining avian diversity is connectivity to extensive PF,
439 rather than forest age. Extinction and colonisation dynamics shape the avifaunas of forests within
440 this landscape, with potentially lower colonization rates in isolated forest sites, and greater
441 colonisation rates in regenerating forests that are sufficiently well connected to PF (Robinson
442 1999, Bradfer-Lawrence *et al.* 2018).

443
444 BIRD POPULATION DENSITY. - The density of birds in SF and PF varied across sites, with the
445 highest density estimates in the youngest SF, matching patterns previously reported for the
446 Neotropics (Blake & Loiselle 2001, Johns 1991, Karr 1976, Petit & Petit 2003). Earlier studies in
447 Soberania National Park have reported densities 2–3 times higher than our PF estimates (Van
448 Bael *et al.* 2013, Robinson *et al.* 2000). The disparity in figures may result from differing
449 methodologies, particularly the spot mapping and smaller point count radius used by previous
450 studies. By contrast, previous population density estimates for young SF (5–6-yr-old) from Agua
451 Salud were about 45 percent lower than estimated population densities from our surveys (20-yr-
452 old), but comparable with our estimates from older SF (Van Bael *et al.* 2013). Higher population
453 density in younger forest may in part reflect increased detectability of some species, particularly
454 those associated with the forest canopy, which is harder to survey in PF (Robinson *et al.* 2018).

455 Our results may also reflect the increased resource availability of both fruit and insects often
456 found in younger SF (Blake & Loiselle 1991, Levey 1988, Martin 1985), which may encourage
457 birds from PF to use adjacent SF for foraging.

458

459 SIMILARITY TO PRIMARY FOREST. - Most studies comparing the similarity of avian species
460 composition between SF and PF report increasing similarity to PF with SF age (Borges 2007,
461 Dent & Wright 2009, Raman 1998). All our SF sites had high levels of compositional similarity
462 to PF, and upper figures were within the range of similarity found in extensive PF. In line with
463 our hypothesis, SF community composition became increasingly similar to that of isolated PF
464 across the chronosequence. However, SF community composition did not converge on that of
465 extensive mainland PF sites with increasing SF age. Similarly, there was no relationship between
466 SF age and the percentage of PF species detected: the highest percentage of PF species was
467 found in the youngest SF that, critically, was also the least isolated and most well-connected to
468 extensive PF.

469 Based on our findings, isolation plays a greater role than forest age in determining the
470 reassembly of bird communities in SF. Despite the persistence of high-stature PF forest on BCI,
471 many species have disappeared from the local community since it was isolated by the inundation
472 of Lake Gatun (Robinson 1999, Willis 1974). While habitat size effects and isolation by water
473 have influenced patterns local extinction, they do not appear to drive our results since peninsula
474 sites have similar bird communities to the island PF, with relatively low species richness. By
475 contrast, bird communities in extensive mainland PF sites include forest specialists that have
476 been lost from both BCI and peninsula sites, and are unlikely to recolonise SF unless it is
477 contiguous with PF that harbours these species. In summary, SF avian communities are at least

478 partially dependent on contiguous PF source populations. If connected PF populations have low
479 species richness then SF will likely never develop the bird communities associated with
480 extensive PF forest stands (Ferraz *et al.* 2007, Jones *et al.* 2016, Stouffer *et al.* 2006). However,
481 if SF sites are adjacent to extensive PF, forest specialists may recolonise relatively rapidly. For
482 example, understorey insectivores increased in abundance just 10 yr after SF was abandoned
483 adjacent to PF in Amazonia (Andrade and Rubio-Torgler, 1994). Our findings highlight that
484 connectivity is critical for reassembly of avian communities in regenerating tropical forests
485 (Barlow *et al.* 2006, Lees & Peres 2009).

486

487 COMPOSITIONAL CHANGES. - Despite the key role of connectivity in determining avian
488 composition, forest age may still influence bird community reassembly, as demonstrated by the
489 increasing similarity of communities in older isolated SF to that of isolated PF. However, six
490 forest species present in isolated PF on BCI were missing from the adjacent 120-yr-old SF,
491 including the forest specialists Long-billed Gnatwren (*Ramphocaenus melanurus*), Rufous
492 Mourner (*Rhytipterna holerythra*), Scaly-throated Leaf-tosser (*Sclerurus guatemalensis*),
493 Semiplumbeous Hawk (*Leucopternis semiplumbeus*), Spot-crowned Antwren (*Dysithamnus*
494 *puncticeps*), and Wood Thrush (*Hylocichla mustelina*). Conversely there were no forest
495 specialists detected in the 120-yr-old SF that were not also present in the isolated PF. Although
496 several studies report a high representation of PF species present in SF (> 70% of PF species),
497 SF communities often lack rare species, or those with highly specialised dietary or habitat
498 requirements (Chazdon *et al.* 2009, Dent & Wright 2009).

499 The loss of forest species from isolated sites across this landscape is striking, and is
500 especially evident when comparing isolated PF with extensive PF. BCI is a relatively large forest

501 fragment (1560 ha), but it has been isolated for > 100 yr and during this time numerous avian
502 extinctions have been documented (Chapman 1938, Eisenmann 1952, Karr 1990, 1982,
503 Robinson 1999, Willis & Eisenmann 1979); 65 species have been lost from the island, including
504 30 forest species and 35 edge species (Robinson 1999). Many of the forest species missing from
505 the PF sites on BCI are understory insectivores such as Dusky Antbird (*Cercomacroides*
506 *tyrannina*), Ocellated Antbird (*Phaenostictus mcleannani*) and Black-faced Antthrush
507 (*Formicarius analis*). In addition, we only detected two of the ten species identified by Robinson
508 (1999) as forest birds that are close to extirpation on BCI: Black-tailed Trogon (*Trogon*
509 *melanurus*) and Rufous Piha (*Lipaugus unirufus*). The isolation of BCI within a large waterbody
510 makes recolonization by many forest species unlikely as they are poorly adapted to sustained
511 flight, and unwilling or incapable of dispersing across open water (Moore *et al.* 2008, Tobias *et*
512 *al.* 2013).

513 Species richness and relative abundance of migratory birds was highest in younger SF,
514 with numbers decreasing with increasing SF forest age. Similarly, Van Bael *et al.* (2013) found
515 more migrant species in SF (5–6-yr-old) than PF sites in central Panama (0.5 and 0.2 birds/point
516 count for SF and PF respectively). Migrant birds may occupy degraded and open habitats
517 because they are excluded from optimal habitats by resident species, or because they are better
518 able to adapt to the resources offered by SF (Greenberg *et al.* 1994, Willis 1980, Wunderle &
519 Latta 1996). Our results add to a growing body of evidence confirming that secondary and
520 degraded tropical forests are important habitats for migrant bird species (Van Bael *et al.* 2007,
521 Greenberg *et al.* 1994, 1997, Wunderle & Latta 1996).

522 It is possible that other aspects of community structure, such as functional and
523 phylogenetic composition, may be affected by forest age and connectivity (Bregman *et al.* 2016,

524 Pigot *et al.* 2016). For example, if SF provides a simplified range of structural and dietary
525 resources for roosting and foraging birds then young forests may not be able to support as many
526 closely related or functionally similar species, resulting in functional and phylogenetic over-
527 dispersion (Bregman *et al.* 2016). In addition, isolation may increase functional and phylogenetic
528 clustering as certain groups are selected against due to their inability to cross gaps between forest
529 fragments (Bregman *et al.* 2016). Further studies are needed to clarify how forest successional
530 status and connectivity across the wider landscape interact to shape bird community
531 composition.

532

533 CAVEATS. – Our study design is limited by the historical and geographical features of the study
534 landscape, making it impossible to establish a fully replicated study within the different levels of
535 isolation and forest age (Denslow & Guzman 2000). Thus, some of the patterns detected may be
536 specific to the local context of water barriers associated with the Panama Canal. Nonetheless,
537 while further studies are now required to assess how far our conclusions can be generalised to
538 other tropical landscapes, we argue that the water barriers and detailed history of forest
539 regeneration in central Panama provide a valuable setting for testing the relative effects of
540 isolation and forest age on the conservation value of SF.

541 Effectively surveying birds across differing habitats remains a methodological challenge
542 for studies such as ours (Buckland *et al.* 2008). We standardised survey effort at all sites at the
543 risk of generating spurious differences in bird communities due to variation in detectability
544 (Bregman *et al.* 2016, Robinson *et al.* 2018). For example, it is possible that estimates of species
545 richness and population density in younger SF are inflated because (1) individual birds tend to be
546 more detectable at forest edges and in lower-stature forests (Barlow *et al.* 2007b, Buckland *et al.*

547 2008, Ruiz-Gutiérrez *et al.* 2010) and (2) PF bird species may have been missed by our surveys
548 because they are relatively quiet, inactive, inconspicuous or occur at low population densities
549 (Terborgh *et al.* 1990, Robinson *et al.* 2018). To explore how detectability may have affected our
550 results, we compared our species counts to a dataset from a previous study surveying the same
551 extensive PF site with more intensive survey methods (including mist-netting), over a far longer
552 survey period, and across an expanded set of species (Robinson *et al.* 2000). Overall, Robinson
553 *et al.* (2000) reported an additional 132 species in the community. However, the majority (65%)
554 of these were either unavoidably or deliberately excluded from our study because they fall into
555 one of three categories: 1) nocturnal, vagrant or migratory species which are not core
556 components of the diurnal communities we studied (45%), 2) aerial species (e.g. swifts and
557 raptors) (15%), and 3) birds associated with aquatic landscape features (5%). Because of our
558 study aims, we did not survey species in the second and third categories, and indeed discounted
559 all individual birds detected on waterbodies or in flight passing over the forest. Excluding these
560 species, 46 species from Robinson *et al.* (2000) were not observed, most of which are rare or
561 difficult to detect. While the absence of these 46 species from our censuses may underestimate
562 the importance of PF for conservation in our analysis, we note that this is a relatively minor
563 component of overall biodiversity and represents a small number of individual birds. Moreover,
564 increased surveying and mist-netting of our SF sites would no doubt also increase species
565 richness in SF by an unknown amount, thus we do not believe our main conclusions would be
566 altered by further surveying.

567 A separate issue relates to the breeding status or viability of populations of forest birds in
568 younger SF. It has been hypothesised that populations of many PF bird species in SF may be
569 non-viable, and therefore less important to conservation, because they are largely made up of (1)

570 transient individuals or (2) temporary territories with infrequent breeding and low breeding
571 success (Tobias *et al.* 2013). We cannot rule out this possibility based on our results, and more
572 research is needed to clarify population demography and viability in SF. However, the relatively
573 high population density of primary forest bird species in younger SF suggests that, at a
574 minimum, SF can greatly increase the population carrying capacity of adjacent PF, thus
575 increasing its importance for conservation.

576

577 CONCLUSIONS. - Our results suggest that the link between tropical bird communities and
578 successional trajectories differs between isolated and non-isolated SF sites; in isolated sites, bird
579 communities converge with isolated PF over time, whereas highly-connected SF sites converge
580 rapidly with extensive PF. In addition, we find evidence that SF, even when relatively young,
581 can support dense populations of PF species, so long as forests are connected to extensive PF.
582 Crucially, even if these populations are transitory, SF may theoretically increase the population
583 carrying capacity of PF, reducing the risk of local extinction. The importance of habitat
584 connectivity highlighted by our results is specifically relevant to tropical forests as species at
585 higher latitudes are often better adapted for survival in SFs or dispersal between patches of PF
586 (Stratford & Robinson 2005). These findings emphasise the importance of reforesting and
587 maintaining existing SF at the borders of extensive tropical forest, and highlight the need for
588 improved protection of SF in protected area buffer zones throughout the tropics.

589

590

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592

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603

604

605 **Data Availability:** Data available from the Dryad Digital Repository: <https://doi.org/10.5061/dryad.77m6100>.

607

608

609 **LITERATURE CITED**

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845

846 **TABLES**

847

848 TABLE 1. Approximate forest age, level of isolation, species richness, Shannon-Weiner
 849 diversity index and dominance statistics for bird communities of ten secondary forest and four
 850 primary forest sites in central Panama, for both forest specialists and all bird species.

Site	Age (yr)	Level of Isolation	All Birds			Forest Specialists ^a		
			Species richness	Shannon–Wiener index	Dominance (%)	Species richness	Shannon–Wiener index	Dominance (%)
1	20	Connected	117	3.97	29.86	40	3.07	46.19
2	20	Connected	113	4.03	27.65	37	3.01	49.22
3	40	Isolated	84	3.63	36.75	26	2.72	54.55
4	40	Isolated	90	3.73	38.01	25	2.75	53.67
5	60	Isolated	76	3.61	37.16	25	2.83	50.00
6	60	Isolated	89	3.73	34.35	30	2.93	45.21
7	90	Connected	95	4.04	22.32	34	3.19	40.05
8	90	Isolated	83	3.77	32.12	26	2.77	54.25
9	120	Isolated	63	3.27	48.37	22	2.45	66.57
10	120	Isolated	62	3.37	46.42	23	2.49	63.84
11	Primary	Isolated	74	3.63	36.89	27	2.78	54.77
12	Primary	Isolated	75	3.64	35.94	28	2.79	54.17

13	Primary	Extensive	99	4.11	21.02	39	3.41	28.74
14	Primary	Extensive	96	3.95	25.13	38	3.20	39.87

851

852 ^a Forest Specialists: species that are scored as having high forest-dependence (BirdLife

853 International 2018).

854

855 TABLE 2. Permutational MANOVA results assessing species composition using community
856 similarity matrices generated with both Morisita-Horn abundance-based similarity index (S_{MH})
857 and Jaccard incidence-based similarity index (S_J). We tested observed differences between forest
858 age, isolation level (isolated or connected), forest type (SF or PF) and geographic distance to
859 extensive mainland PF. We also investigated the effect of season (wet or dry) on species
860 composition by conducting a perMANOVA at survey level.

		All Birds				Forest Specialists ^a			
		R^2	F	df	P	R^2	F	df	P
S_{MH}	Forest isolation	0.15	9.12	1	< 0.01	0.18	6.28	1	< 0.05
	Forest age	0.13	7.80	1	< 0.01	0.02	0.84	1	ns
	Forest type (SF vs PF)	0.08	5.10	1	< 0.01	0.01	0.51	1	ns
	Distance to extensive PF	0.02	1.35	1	ns	0.04	1.31	1	ns
S_J	Forest isolation	0.10	2.19	1	< 0.05	0.11	2.18	1	< 0.05
	Forest age	0.11	2.42	1	< 0.05	0.07	1.37	1	ns
	Forest type (SF vs PF)	0.09	1.93	1	< 0.05	0.06	1.14	1	ns
	Distance to extensive PF	0.06	1.29	1	ns	0.06	1.17	1	ns
S_{MH}^b	Season (wet v. dry)	0.05	11.69	1	< 0.01	0.04	8.84	1	< 0.01
	Forest isolation	0.03	7.45	1	< 0.01	0.05	9.38	1	< 0.01
	Forest age	0.05	10.29	1	< 0.01	0.04	7.15	1	< 0.01
	Forest type (SF vs PF)	0.03	6.30	1	< 0.01	0.02	5.35	1	< 0.01
	Distance to extensive PF	0.03	5.69	1	< 0.01	0.02	3.56	1	< 0.05

861 ^a Forest Specialists: species that are scored as having high forest-dependence (BirdLife
862 International 2018).

863 ^bData analysed at survey level, with season included.

864 **FIGURE LEGENDS**

865

866 **FIGURE 1.** Map of the 14 study sites in central Panama. Sites are colour-coded by forest age (PF
867 = primary forest; SF = secondary forest). This area of central Panama is composed of a mosaic of
868 contiguous forest stands of different age interspersed among a matrix of water and agricultural
869 land. The main areas in which forest stands are embedded among other patches of forest (and
870 therefore difficult to discretely identify) are the Barro Colorado National Monument (BCNM),
871 including Barro Colorado Island (1,560 ha) and Gigante peninsulas (2,600 ha), Soberania
872 National Park (SNP) and surrounding contiguous forest (22,000 ha), including the Agua Salud
873 Project (ASP).

874

875 **FIGURE 2.** Bird species richness by forest age, forest type, and degree of isolation from
876 extensive PF. “Forest Specialists” are species that are scored as having high forest-dependence
877 (BirdLife International 2018).

878

879 **FIGURE 3.** Population density estimates and 95% confidence intervals for all bird species
880 (number of birds per hectare) using distance corrections. Species data has been pooled for the
881 two sites in each of seven forest age categories.

882

883 **FIGURE 4.** Percentage of bird species detected in primary forest (PF) that were also detected in
884 secondary forest (SF) in five SF age categories for isolated PF sites and extensive PF sites.

885 Species data has been pooled for the two sites in each of seven forest age categories.

886

887 FIGURE 5. Similarity (Morisita-Horn) between secondary forest sites (SF) and both isolated and
888 extensive primary forest (PF). Each bar represents the mean similarity index (± 1 SE) between the
889 two SF sites in each age category (20, 40, 60, 90 and 120 yr old) and the PF sites. Calculated
890 using the all bird dataset.

891

892 FIGURE 6. Non-metric multidimensional scaling (NMDS) plots of bird communities in two
893 forest sites in each of five secondary forest (SF) age categories (20, 40, 60, 90, 120-yr-old), and
894 isolated primary forest (PF) and extensive PF. NMDS were generated using the Morisita-Horn
895 index (all birds stress = 0.07; forest specialists stress = 0.08) and Jaccard index (all birds stress =
896 0.07; forest specialists stress = 0.06). Isolation levels are represented by different symbols.

897 “Forest Specialists” are species that are scored as having high forest-dependence (BirdLife
898 International 2018).

899

900 FIGURE S1: Species accumulation curves scaled by (A) sample and (B) by number of individual
901 birds detected for bird communities in each of five secondary forest (SF) age categories: 20, 40,
902 60, 90, 120-yr-old, and isolated primary forest (PF) and extensive PF for all birds.

903

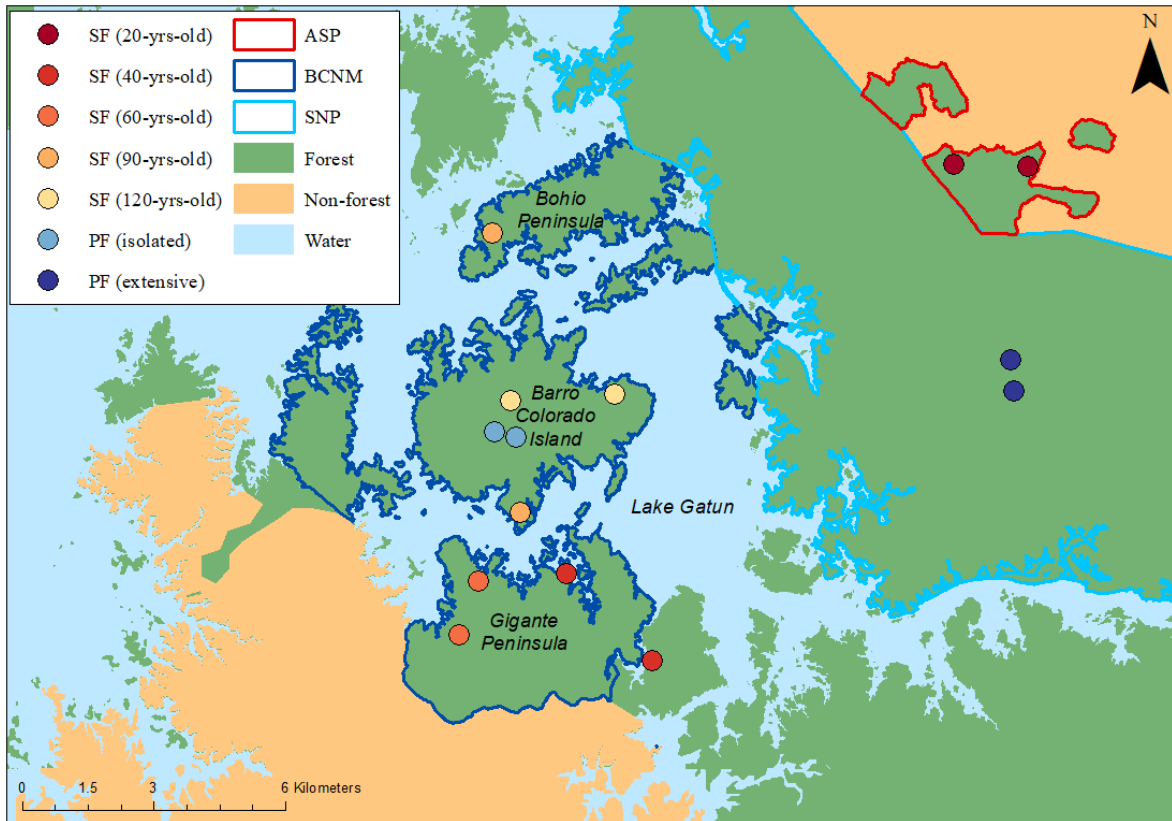
904 FIGURE S2: Species accumulation curves scaled by (A) sample and (B) by number of individual
905 birds detected for bird communities in each of five secondary forest (SF) age categories: 20, 40,
906 60, 90, 120-yr-old, and isolated primary forest (PF) and extensive PF for forest specialists.

907 “Forest Specialists” are species that are scored as having high forest-dependence (BirdLife
908 International 2018).

909

910 FIGURE S3. Mean number of migratory birds detected per point count (± 1 SE), from 126
911 locations.

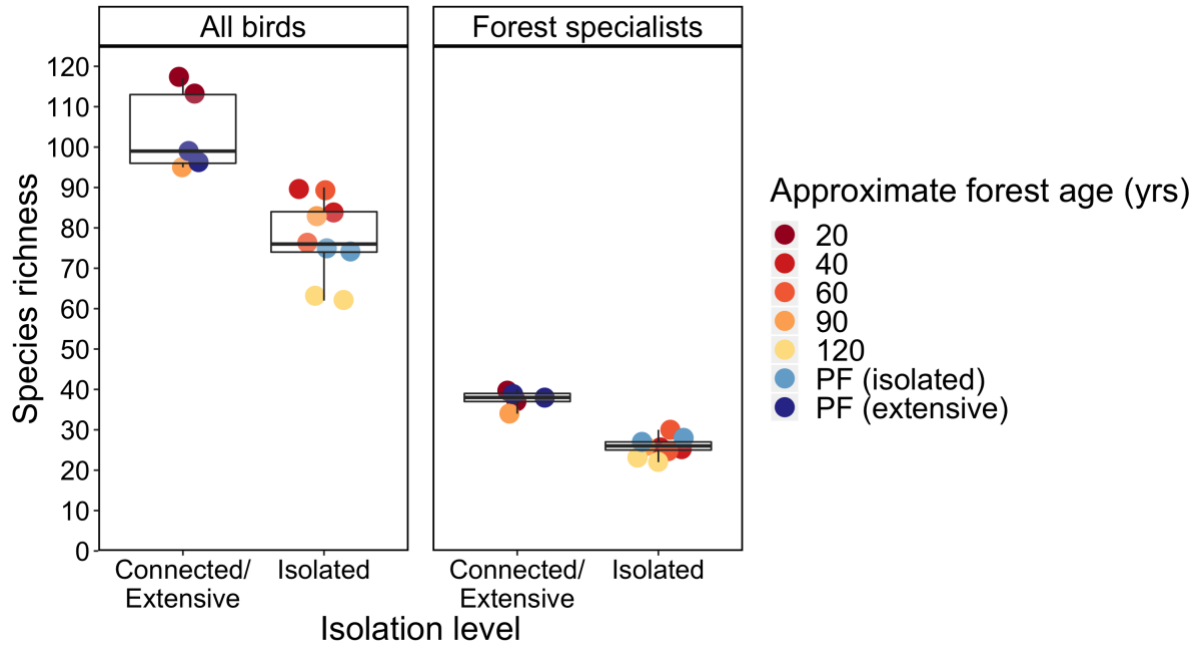
912 **FIGURES**



913

914 **FIGURE 1.**

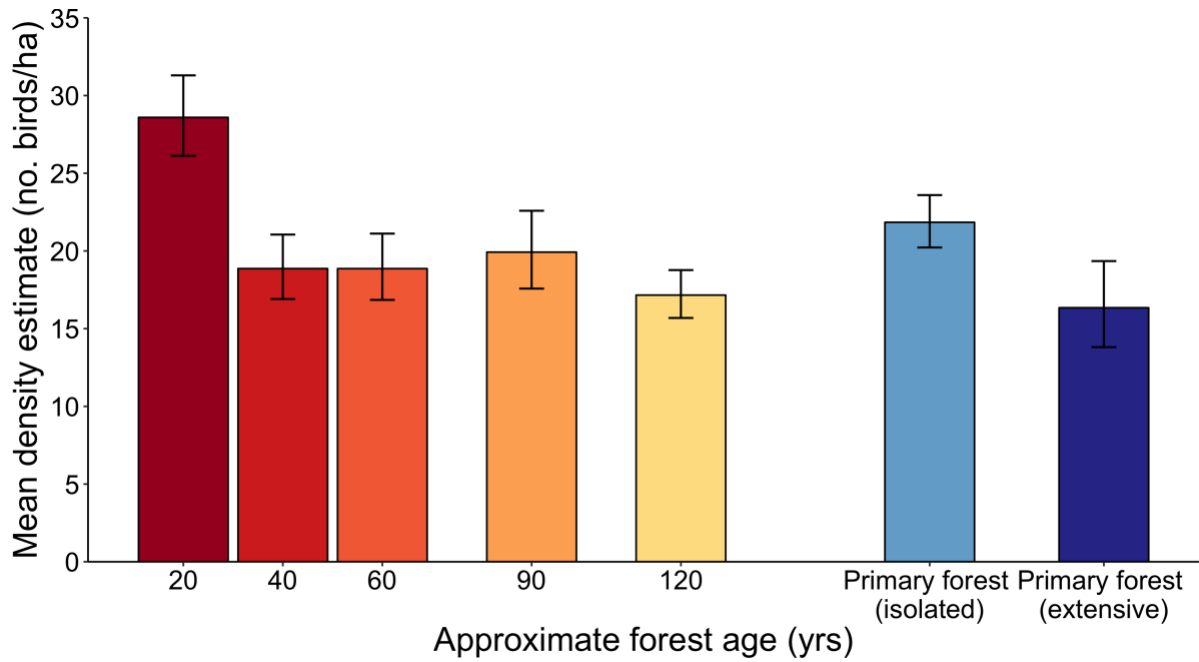
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917 FIGURE 2.

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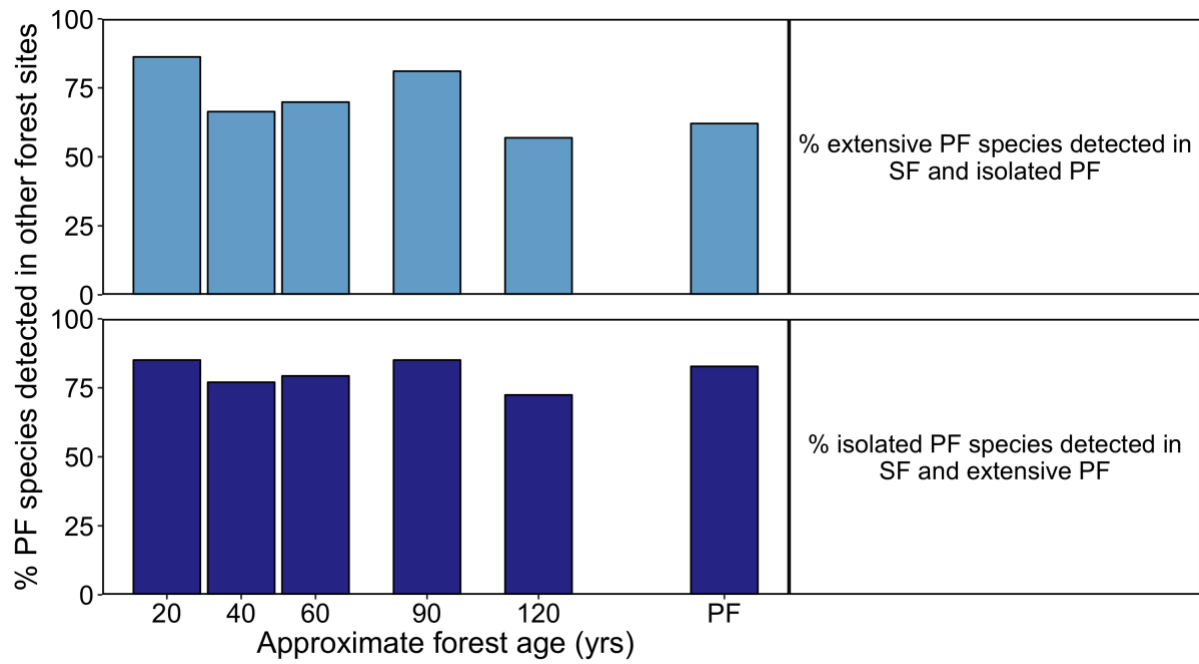


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920 FIGURE 3.

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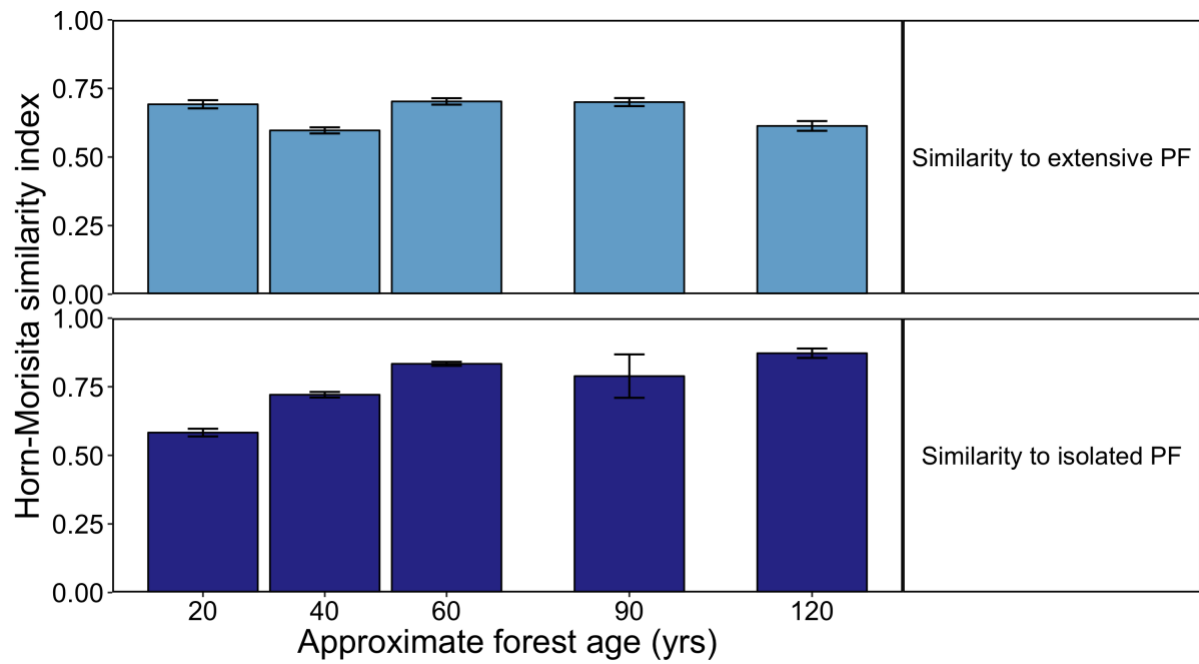
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924 FIGURE 4.

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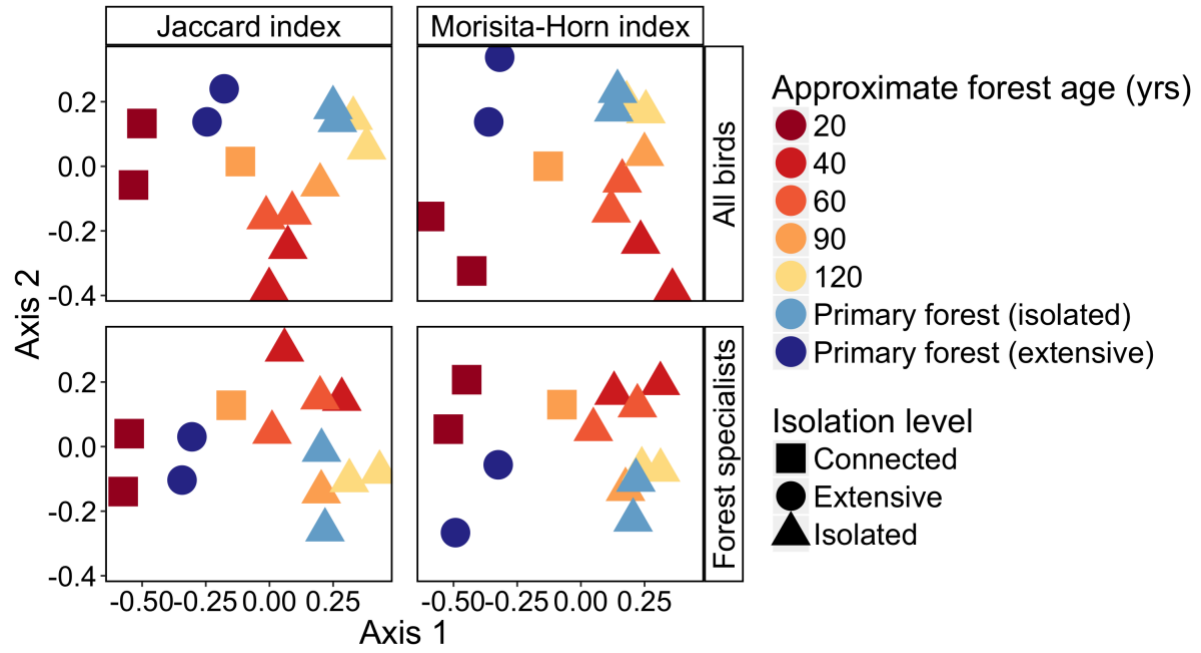


926

927 FIGURE 5.

928

929



930

931 FIGURE 6.

932 **SUPPLEMENTARY INFORMATION**

933

934 **TABLE S1: Bird species list from our surveys in central Panama, with ecological classifications**

935 used in analyses and species presence/absence in sites.

Scientific Name	Migrant Species	Forest Specialist ^b	Site ^c													
			1	2	3	4	5	6	7	8	9	10	11	12	13	14
<i>Tinamus major</i>	0	1	1	0	1	1	1	1	1	1	1	1	1	1	1	1
<i>Crypturellus soui</i>	0	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0
<i>Penelope purpurascens</i>	0	1	0	0	0	0	0	0	0	1	1	1	1	1	0	0
<i>Ortalis cinereiceps</i>	0	0	0	1	1	1	0	1	0	0	0	0	0	0	0	0
<i>Odontophorus gujanensis</i> ^a	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Patagioenas speciosa</i>	0	0	1	1	1	0	0	1	0	1	0	0	1	0	1	0
<i>Patagioenas cayennensis</i>	0	0	1	0	1	1	1	1	1	1	0	0	1	1	1	1
<i>Patagioenas nigrirostris</i>	0	0	1	1	1	1	0	0	1	0	0	1	1	1	1	1
<i>Geotrygon montana</i>	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0
<i>Leptotila verreauxi</i>	0	0	0	0	1	1	1	1	0	0	0	0	0	0	0	0
<i>Leptotila cassinii</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Columbina talpacoti</i>	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0
<i>Claravis pretiosa</i>	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Nyctibius griseus</i> ^a	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Chordeiles minor</i> ^a	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0

<i>Nyctidromus albicollis</i>	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1	0
<i>Antrostomus rufus</i>	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0
<i>Florisuga mellivora</i>	0	0	1	1	0	0	0	0	0	1	1	0	0	0	1	0
<i>Phaethornis striigularis</i>	0	0	1	1	1	1	1	1	1	1	1	0	0	0	1	1
<i>Phaethornis longirostris</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Heliothryx barroti</i>	0	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Thalurania colombica</i>	0	0	1	1	1	0	0	0	0	0	0	0	0	0	1	0
<i>Amazilia tzacatl</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Amazilia amabilis</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Lepidopyga coeruleogularis</i>	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Juliamyia julie</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Dromococcyx phasianellus</i>	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0
<i>Piaya cayana</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	0
<i>Aramides cajaneus</i>	0	0	0	0	0	1	1	0	1	0	0	0	0	0	0	0
<i>Pulsatrix perspicillata</i>	0	0	0	0	0	1	0	0	0	0	0	1	0	0	0	0
<i>Leptodon cayanensis</i>	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0
<i>Elanoides forficatus</i>	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
<i>Harpagus bidentatus</i>	0	0	0	0	0	0	0	0	0	1	1	0	0	0	0	1
<i>Buteogallus anthracinus</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Buteogallus urubitinga</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Pseudastur albicollis</i>	0	1	0	0	0	0	1	0	1	0	0	0	0	0	0	0
<i>Leucopternis semiplumbeus</i>	0	1	0	0	1	0	0	0	0	0	0	0	1	1	0	0
<i>Buteo nitidus</i>	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0

<i>Buteo platypterus</i>	1	0	0	0	0	0	0	1	0	1	0	0	0	0	0	0
<i>Buteo albonotatus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
<i>Trogon massena</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Trogon melanurus</i>	0	0	1	0	1	1	1	1	1	1	0	0	0	0	1	0
<i>Trogon chionurus</i>	0	0	1	1	0	0	0	0	1	1	0	1	1	0	1	0
<i>Trogon caligatus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Trogon rufus</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Momotus subrufescens</i>	0	0	0	1	1	1	1	1	0	0	0	0	0	0	0	0
<i>Baryphthengus martii</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Electron platyrhynchum</i>	0	1	1	1	0	1	0	1	1	0	0	0	0	0	1	1
<i>Notharchus hyperrhynchus</i>	0	0	0	0	0	1	1	1	0	0	0	0	1	0	0	0
<i>Notharchus pectoralis</i>	0	1	1	1	1	1	1	1	0	1	1	1	1	1	1	1
<i>Malacoptila panamensis</i>	0	0	0	1	0	1	1	1	1	1	1	1	1	1	1	1
<i>Ramphastos ambiguus</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Ramphastos sulfuratus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Pteroglossus torquatus</i>	0	1	0	0	1	1	1	1	1	0	1	0	1	0	1	0
<i>Campephilus melanoleucos</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Celeus loricatus</i>	0	1	1	0	0	0	0	1	1	0	0	0	0	0	1	1
<i>Hylatomus lineatus</i>	0	0	0	1	1	1	1	1	0	1	1	1	1	1	1	1
<i>Melanerpes pucherani</i>	0	1	1	1	1	0	1	1	1	1	1	1	1	1	1	1
<i>Melanerpes rubricapillus</i>	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Micrastur mirandollei</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	0	1	1
<i>Micrastur semitorquatus</i>	0	0	1	0	1	0	0	1	1	1	1	1	1	1	0	1

<i>Falco ruficularis</i>	0	1	0	0	0	0	0	1	0	1	0	0	0	0	0	1
<i>Brotogeris jugularis</i>	0	0	1	1	1	1	0	1	0	0	0	0	0	0	0	1
<i>Pyrilia haematotis</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	1
<i>Pionus menstruus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Amazona autumnalis</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Amazona farinosa</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Microrhopias quixensis</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Epinecophylla fulviventris</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Myrmotherula ignota</i>	0	1	0	1	0	1	0	0	0	0	0	0	0	0	0	1
<i>Myrmotherula axillaris</i>	0	1	1	1	1	0	1	1	1	1	1	1	1	1	1	1
<i>Dysithamnus puncticeps</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	1	1	1
<i>Cymbilaimus lineatus</i>	0	0	1	0	0	0	0	0	1	0	0	0	0	0	1	1
<i>Taraba major</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Thamnophilus atrinucha</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Cercomacroides tyrannina</i>	0	0	1	1	1	1	0	1	1	0	0	0	0	0	1	1
<i>Phaenostictus mcleannani</i>	0	0	1	1	0	0	0	0	1	0	0	0	0	0	1	1
<i>Gymnopithys bicolor</i>	0	1	1	1	1	1	1	1	1	1	0	1	1	1	1	1
<i>Hylophylax naevioides</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Poliocrania exsul</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Myrmeciza longipes</i>	0	0	0	1	0	0	0	0	0	1	0	0	1	0	0	0
<i>Hylopezus perspicillatus</i>	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0
<i>Formicarius analis</i>	0	1	1	1	1	1	1	1	1	0	0	0	0	0	1	1
<i>Sclerurus mexicanus</i>	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0

<i>Sclerurus guatemalensis</i>	0	1	1	1	0	1	0	1	1	0	0	0	1	1	1	1
<i>Deconychura typica</i>	0	1	0	0	0	0	0	1	1	0	0	0	0	0	0	0
<i>Sittasomus griseus</i>	0	0	0	0	0	0	0	1	1	0	0	0	0	0	1	0
<i>Dendrocincla fuliginosa</i>	0	0	1	1	0	1	1	1	1	1	1	1	1	1	1	1
<i>Glyphorhynchus spirurus</i>	0	0	1	0	1	0	1	1	1	1	1	1	1	1	1	1
<i>Dendrocolaptes sanctithomae</i>	0	0	1	0	0	1	1	0	1	0	1	1	1	1	0	1
<i>Xiphorhynchus susurrans</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Xiphorhynchus lachrymosus</i>	0	0	1	1	0	0	0	0	0	1	1	1	1	1	1	1
<i>Lepidocolaptes souleyetii</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
<i>Xenops genibarbis</i>	0	1	1	1	0	1	0	1	1	0	1	1	1	1	1	1
<i>Automolus ochrolaemus</i>	0	0	1	1	0	0	0	0	0	0	0	0	0	0	1	1
<i>Manacus vitellinus</i>	0	0	1	1	1	1	0	0	0	0	0	0	0	0	0	0
<i>Ceratopipra mentalis</i>	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Lepidothrix coronata</i>	0	1	1	1	0	0	1	1	1	0	0	0	0	0	1	1
<i>Querula purpurata</i>	0	1	0	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Lipaugus unirufus</i>	0	0	1	0	0	0	0	0	1	0	0	0	1	0	1	0
<i>Onychorhynchus coronatus</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Terenotriccus erythrurus</i>	0	1	1	1	1	1	1	0	1	1	1	1	1	1	1	1
<i>Tityra semifasciata</i>	0	0	0	1	1	0	1	1	1	1	0	1	1	1	1	1
<i>Pachyramphus polychopterus</i> ^a	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
<i>Schiffornis veraepacis</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	0	1	0
<i>Platyrinchus coronatus</i>	0	1	0	0	0	0	0	0	1	0	0	0	0	0	1	1
<i>Mionectes oleagineus</i>	0	0	1	1	1	0	1	0	1	0	1	1	1	1	1	1

<i>Cnipodectes subbrunneus</i>	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1
<i>Rhynchocyclus aequinoctialis</i>	0	1	1	1	0	0	0	1	1	1	1	1	1	1	1	1
<i>Tolmomyias sulphurescens</i>	0	0	0	1	0	0	1	1	0	0	0	0	0	0	0	0
<i>Tolmomyias assimilis</i>	0	1	0	0	1	1	1	1	1	1	1	1	1	1	1	1
<i>Myiornis atricapillus</i>	0	0	1	0	0	0	0	1	1	1	0	1	1	1	1	1
<i>Oncostoma olivaceum</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Todirostrum nigriceps</i>	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
<i>Zimmerius vilissimus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Ornithion brunneicapillus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Camptostoma obsoletum</i>	0	0	1	0	1	1	1	0	0	1	0	0	0	0	0	0
<i>Tyrannulus elatus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	0
<i>Myiopagis gaimardii</i>	0	1	1	1	1	1	1	1	1	1	0	1	1	1	1	0
<i>Myiopagis viridicata</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
<i>Attila spadiceus</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Legatus leucophaeus</i>	1	0	1	0	0	0	0	0	0	0	0	0	0	0	1	0
<i>Pitangus sulphuratus</i>	0	0	1	0	1	1	0	1	1	1	0	0	0	0	0	0
<i>Megarynchus pitangua</i>	0	0	1	0	0	0	0	0	0	0	0	0	1	0	0	0
<i>Myiodynastes maculatus</i>	0	0	0	1	1	1	1	0	1	1	1	1	0	0	0	0
<i>Myiozetetes cayanensis</i>	0	0	0	0	1	1	0	0	0	1	0	0	0	0	0	0
<i>Myiozetetes similis</i>	0	0	1	0	1	1	1	0	1	1	0	0	0	1	0	0
<i>Tyrannus melancholicus</i>	0	0	0	0	1	1	0	0	1	1	1	0	0	0	0	0
<i>Rhytipterna holerythra</i>	0	1	1	0	0	0	0	0	1	1	0	0	0	1	1	0
<i>Myiarchus tuberculifer</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1

<i>Myiarchus crinitus</i>	1	0	1	1	1	1	1	1	1	1	1	1	0	1	1	1	1
<i>Empidonax virescens</i>	1	0	1	1	1	1	1	1	0	1	1	1	1	1	1	1	1
<i>Contopus cooperi</i>	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Contopus virens</i>	1	0	1	1	0	0	0	0	0	0	0	0	0	0	0	1	1
<i>Vireolanius pulchellus</i>	0	0	1	1	0	0	1	0	1	0	0	0	0	0	0	1	1
<i>Tunchiornis ochraceiceps</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1
<i>Pachysylvia decurtata</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Pachysylvia aurantiifrons</i>	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
<i>Vireo olivaceus</i>	1	1	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Microbates cinereiventris</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	1	1
<i>Ramphocaenus melanurus</i>	0	1	1	1	1	1	1	1	1	1	0	0	1	0	1	1	1
<i>Polioptila plumbea</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Microcerculus marginatus</i>	0	1	1	1	0	0	0	0	0	0	0	0	0	0	0	1	1
<i>Pheugopedius fasciatoventris</i>	0	0	1	1	1	0	0	1	1	0	0	0	0	0	0	1	1
<i>Thryophilus rufalbus</i>	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1	0
<i>Cantorchilus leucotis</i> ^a	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Cantorchilus nigricapillus</i>	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	0	0
<i>Henicorhina leucosticta</i>	0	1	1	1	0	0	0	0	1	0	0	0	0	0	0	1	1
<i>Cyphorhinus phaeocephalus</i>	0	1	1	1	1	1	0	0	1	1	0	0	0	0	0	1	1
<i>Hylocichla mustelina</i>	1	1	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
<i>Turdus grayi</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Euphonia luteicapilla</i>	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Euphonia laniirostris</i>	0	0	1	1	0	0	0	1	0	0	0	0	0	0	0	0	0

<i>Euphonia fulvicrissa</i>	0	0	0	1	0	0	1	0	1	0	0	0	0	1	1	1
<i>Euphonia minuta</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Rhodinocichla rosea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
<i>Arremonops conirostris</i>	0	0	1	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Arremon aurantirostris</i>	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
<i>Amblycercus holosericeus</i>	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0
<i>Psarocolius wagleri</i>	0	0	1	1	0	1	0	1	1	1	1	0	0	0	1	1
<i>Cacicus microrhynchus</i>	0	0	1	1	1	0	1	1	1	0	0	0	0	0	1	1
<i>Cacicus cela</i>	0	0	0	0	1	1	1	1	1	1	0	0	0	1	1	0
<i>Icterus chrysater</i>	0	0	0	1	0	1	1	0	1	1	0	0	0	0	0	0
<i>Parkesia noveboracensis</i>	1	0	0	0	0	0	0	1	1	0	0	0	0	0	0	0
<i>Mniotilta varia</i>	1	0	1	0	0	0	0	0	0	0	0	0	0	1	0	0
<i>Leiothlypis peregrina</i>	1	0	1	1	0	0	0	0	1	0	0	0	0	0	0	0
<i>Setophaga ruticilla</i>	1	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1
<i>Setophaga castanea</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
<i>Setophaga pensylvanica</i>	1	0	1	1	0	0	0	0	0	1	1	1	0	0	0	1
<i>Cyanoloxia cyanooides</i>	0	0	1	1	1	1	1	1	1	1	0	0	0	0	1	1
<i>Habia fuscicauda</i>	0	0	1	1	1	1	1	1	1	1	0	0	1	1	1	1
<i>Habia carmioli</i> ^a	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
<i>Piranga rubra</i>	1	0	0	1	1	1	0	0	0	1	0	0	0	1	1	0
<i>Chlorophanes spiza</i>	0	0	1	1	1	1	1	1	1	1	1	0	0	0	0	1
<i>Cyanerpes cyaneus</i>	0	0	0	1	1	1	1	1	1	1	1	1	1	1	0	0
<i>Cyanerpes lucidus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0

<i>Dacnis cayana</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	0	1
<i>Saltator grossus</i>	0	0	1	1	0	1	1	1	1	0	0	0	0	0	1	1
<i>Eucometis penicillata</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Islerothraupis luctuosa</i>	0	0	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>Chrysocorypha delatrii</i>	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0
<i>Tangara palmarum</i>	0	0	0	0	1	0	0	1	0	0	0	0	0	0	0	0
<i>Tangara inornata</i>	0	0	0	1	0	1	0	1	0	1	1	0	1	1	0	0

936

937 ^a Species detected only in encounters, and not in fixed radius point counts.

938 ^b Forest Specialists: species that are scored as having high forest-dependence (BirdLife

939 International 2018).

940 ^c Site details: 1 and 2 = mainland 20-yr-old SF; 3 and 4 = peninsula 40-yr-old SF; 5 and 6 =

941 peninsula 60-yr-old SF; 7 = mainland 90-yr-old SF; 8 = island 90-yr-old SF; 9 and 10 = island

942 120-yr-old SF; 11 and 12 = island PF; 13 and 14 = mainland PF.

943 TABLE S2. Most abundant bird species in each forest age category in Central Panama and their
 944 primary foraging substrate, feeding guild and forest specialist status.

Forest age	Scientific name	No. individuals	% of all observations	Foraging Height ^a	Feeding Guild ^b	Forest Specialist ^c
20 (n = 2640 individuals)	<i>Oncostoma olivaceum</i>	155	5.9	L	I	0
	<i>Thamnophilus atrinucha</i>	154	5.8	L	I	0
	<i>Cercomacroides tyrannina</i>	140	5.3	L	I	0
	<i>Ramphocaenus melanurus</i>	104	3.9	L	I	1
	<i>Poliocrania exsul</i>	99	3.8	L	I	1
40 (n = 2239 individuals)	<i>Thamnophilus atrinucha</i>	248	11.1	L	I	0
	<i>Pachysylvia decurtata</i>	183	8.2	M/U	I	0
	<i>Xiphorhynchus susurrans</i>	121	5.4	L	I	0
	<i>Oncostoma olivaceum</i>	115	5.1	L	I	0
	<i>Tyrannulus elatus</i>	107	4.8	L/M	I	0
60 (n = 2223 individuals)	<i>Thamnophilus atrinucha</i>	264	11.9	L	I	0
	<i>Pachysylvia decurtata</i>	169	7.6	M/U	I	0
	<i>Xiphorhynchus susurrans</i>	108	4.9	L	I	0
	<i>Poliocrania exsul</i>	101	4.5	L	I	1
	<i>Oncostoma olivaceum</i>	93	4.2	L	I	0
90	<i>Pachysylvia decurtata</i>	161	6.4	M/U	I	0

(n = 2534 individuals)	<i>Poliocrania exsul</i>	151	6	L	I	1
	<i>Amazona autumnalis</i>	133	5.2	U	F	0
	<i>Thamnophilus atrinucha</i>	115	4.5	L	I	0
	<i>Xiphorhynchus susurrans</i>	100	3.9	L	I	0
<hr/>						
	<i>Thamnophilus atrinucha</i>	299	13.9	L	I	0
120	<i>Poliocrania exsul</i>	203	9.4	L	I	1
(n = 2149 individuals)	<i>Pachysylvia decurtata</i>	162	7.5	M/U	I	0
	<i>Xiphorhynchus susurrans</i>	149	6.9	L	I	0
	<i>Hylophylax naevioides</i>	134	6.2	L	I	1
<hr/>						
	<i>Amazona autumnalis</i>	272	9.3	U	F	0
PF (isolated)	<i>Thamnophilus atrinucha</i>	243	8.3	L	I	0
(n = 2922 individuals)	<i>Poliocrania exsul</i>	237	8.1	L	I	1
	<i>Pachysylvia decurtata</i>	221	7.6	M/U	I	0
	<i>Hylophylax naevioides</i>	135	4.6	L	I	1
<hr/>						
	<i>Amazona autumnalis</i>	159	6.6	U	F	0
PF (extensive)	<i>Thamnophilus atrinucha</i>	129	5.4	L	I	0
(n = 2391 individuals)	<i>Pachysylvia decurtata</i>	119	5	M/U	I	0
	<i>Microrhopias quixensis</i>	91	3.8	L/M	I	1
	<i>Zimmerius vilissimus</i>	72	3	C	O	0

945 ^a Foraging height, classified using information in Ridgely & Gwynne (1989): L = "lower levels"
946 (from ground to about 10 feet above the ground); M = "middle levels" (from about 10 to 30 feet
947 above the ground); U = "upper levels" (everything above 30 feet including the canopy); C =

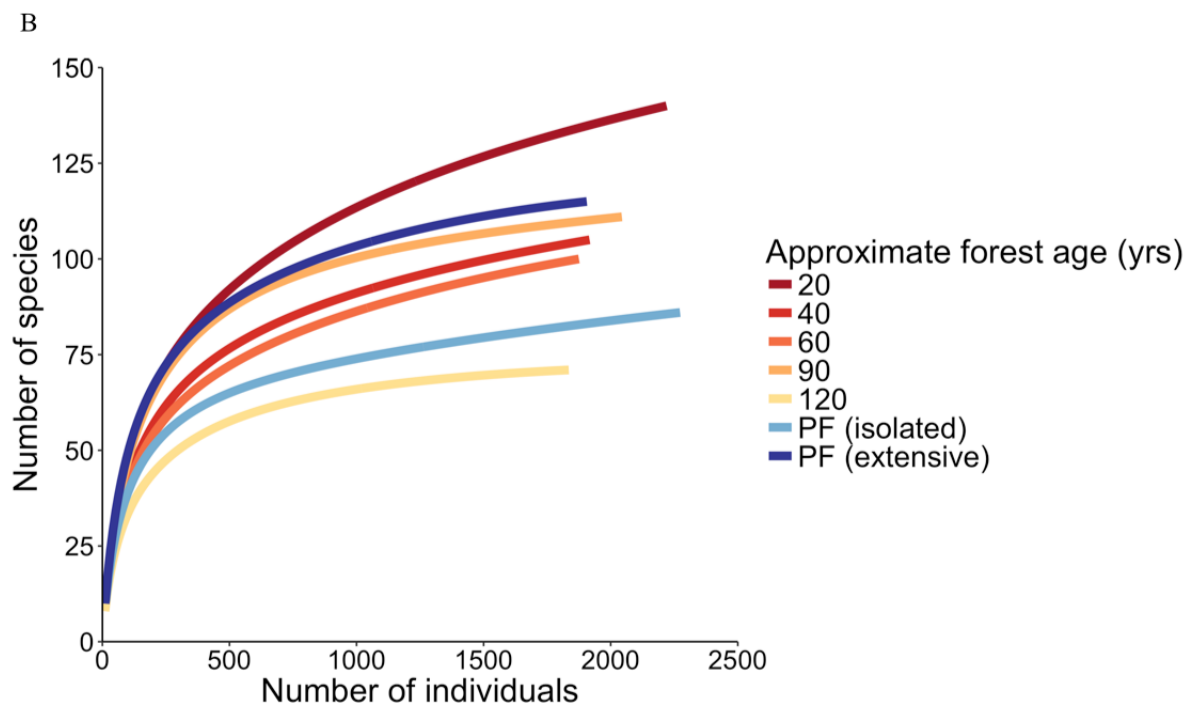
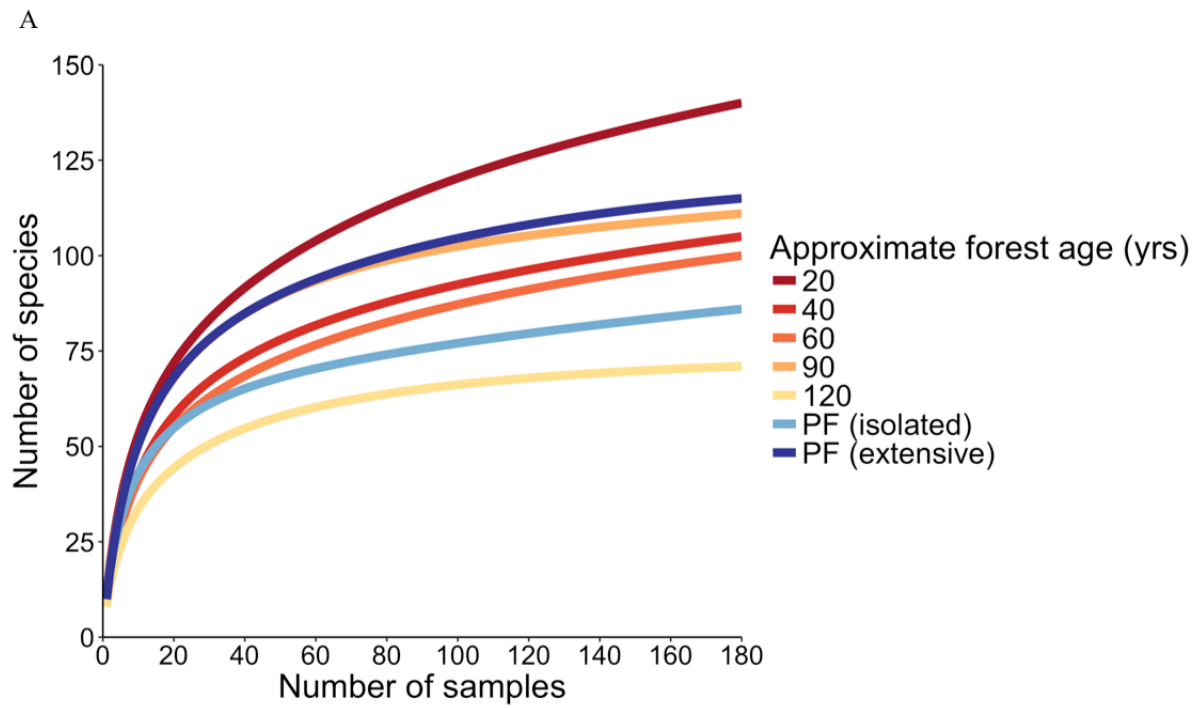
948 "only canopy" (the very top layer of the forest, within about 10ft of the upper level of leaves
949 only).

950 ^b Feeding Guild, classified using information in Wilman *et al.* (2014): F = frugivore; I =
951 insectivore; O = omnivore.

952 ^c Forest Specialist: species that are scored as having high forest-dependence (BirdLife
953 International 2018).

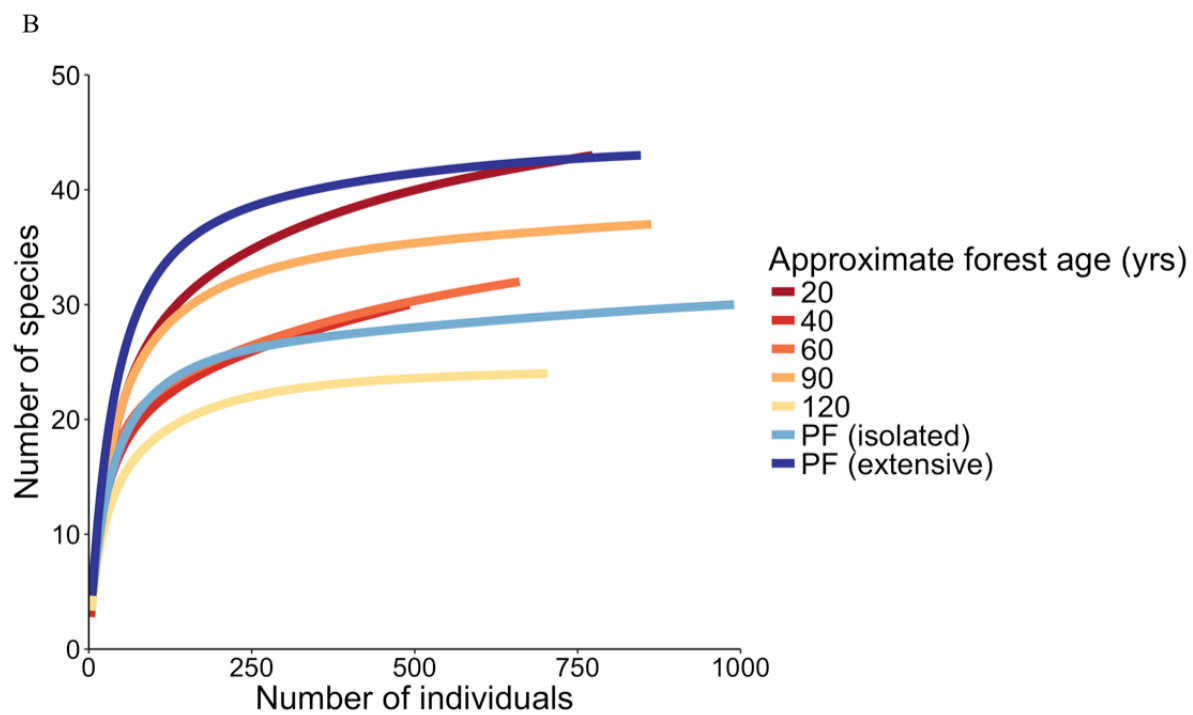
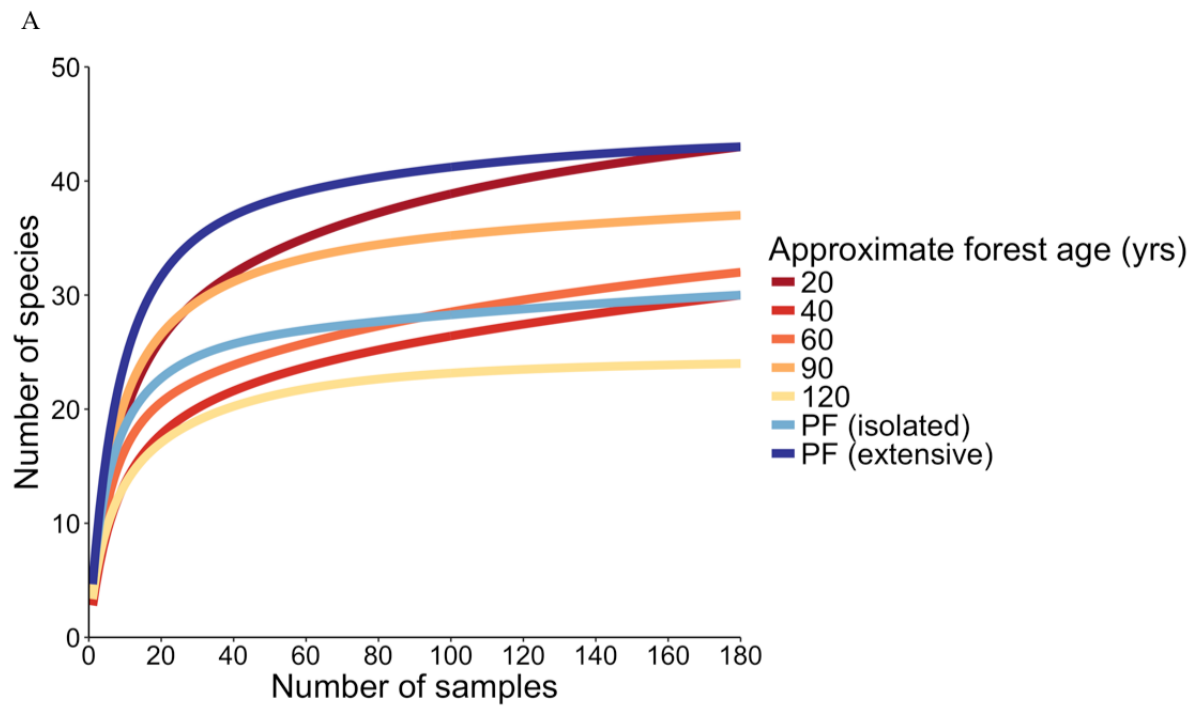
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957 FIGURE S1

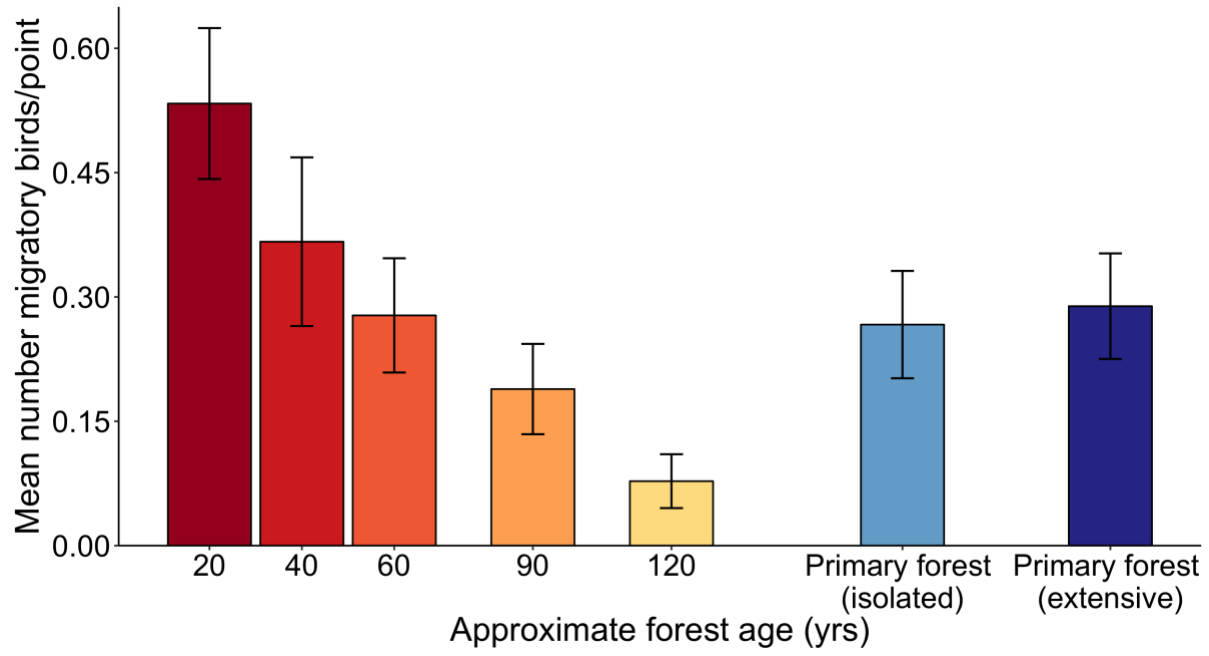


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959 FIGURE S2

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963 FIGURE S3.

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