# Short- and long-term temporal changes in the assemblage structure of Amazonian dung beetles

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#### 2 dung beetles

3

# 4 Abstract

5 Species diversity varies in space and time. Temporal changes in the structure and 6 dynamics of communities can occur at different scales. We investigate the temporal 7 changes of dung beetle assemblages in the Amazonian region along seasons, years, and 8 successional stages. We evaluated if assemblage structure changes between temporal 9 scales and whether such changes affect the functional structure of communities. To 10 achieve these goals, we sampled dung beetles using linear transects of baited pitfall 11 traps during the dry and rainy seasons at two natural reserves in the Amazon region, 12 each representing different time scales: one covering successional variations (80, 30, 13 five, and one years of recovery from logging) and the other one encompassing three consecutive years at two successional stages (20 and 10 years from logging). We used 14 15 Generalized Linear Models to analyze interannual and successional changes in 16 diversity, described assemblage structure with a NMDS, and examined compositional 17 variation by partitioning beta diversity into its nestedness and turnover components. 18 Abundance and richness decrease from the rainy to the dry season and towards earlier 19 successional stages but do not differ between years. Assemblage diversity changes 20 differently in interannual and successional scales. During succession, dung beetle 21 assemblages change drastically, following a nested structure due to the appearance of 22 species and functional groups in later successional stages. In contrast, functional group 23 composition does not show consistent changes between years, displaying a turnover 24 structure. This pattern supports non-deterministic changes in dung beetle assemblage 25 structure along forest succession.

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Keywords Beta diversity, Functional groups, Interannual variations, Nestedness,
Scarabaeinae, Species turnover

29

## 30 Introduction

31 Species diversity varies across both space and time (Rosenzweig 1995). The structure 32 and dynamics of communities vary at multiple temporal scales, from daily and seasonal 33 changes to variations throughout geological periods (Villéger et al. 2011; Fritz et al. 34 2013; Grøtan et al. 2014). Within short ecological time scales, local communities show 35 regular and, to some extent, predictable seasonal variations, coupled with seemingly stochastic interannual changes (e.g., Grimbacher and Stork 2009; Labidi et al. 2012). 36 37 Such stochasticity is produced by phenological and populational processes (e.g., Tanner 38 et al. 2009; Hodgson et al. 2010) that are in turn dependent on interannual changes in 39 climate and temporal variations in resource availability (e.g., Voss et al. 2009; Encinas-40 Viso et al. 2012). Over longer time periods, the effects of large-scale processes, such as 41 climate change, biological invasions, or land transformation, also promote different kinds of responses and changes to community dynamics (see Forister et al. 2010; 42 43 Dijkstra et al. 2011).

Long-term changes in community composition have been traditionally studied under the umbrella of ecological succession. In general, successions can be described as the non-random changes in the structure and composition of an ecological community that take place over time after a disturbance event (e.g., Walker and Del Moral 2003). Currently, many successions take place after habitat perturbations caused directly or indirectly by human activities (e.g., logging, agricultural intensification, or cattle expansion), and they are thought to have negative effects on biodiversity and its related 51 ecological functions, as well as on ecosystem services (Nichols et al. 2007; Horgan
52 2008; Barragan et al. 2011; Braga et al. 2013).

53 The classical view of ecological successions hypothesizes that communities in a 54 new or post-disturbed habitat are formed by a few pioneering species that are 55 subsequently replaced by competitively dominant ones (Connell and Slatyer 1977). This 56 hypothesis, initially proposed by Clements (1916), has received considerable support. 57 However, its view of temporal community dynamics as a directional deterministic 58 process has also been traditionally questioned (Walker and Del Moral 2003). Since the 59 first criticisms of Gleason (1927), detractors of this idea argue that community 60 dynamics are not completely deterministic, giving more relevance to historical 61 contingencies and large-scale processes, such as the individualistic responses of species 62 to the environment (Hortal et al. 2012). Under this paradigm, communities assemble 63 from the dispersal of species that are present and/or arrive in the landscape, with local 64 interactions playing a comparatively less important role. Here, local assembly becomes 65 a density-dependent phenomenon, where the first arrivals correspond to species that are 66 frequent and/or abundant in the landscape, receiving comparatively less rare species. 67 Following this idea, the core-satellite species hypothesis (Hanski 1982) states that the 68 first colonizers would become the core species (i.e., species that make up the bulk of the 69 individuals of the local community), whereas the satellite species (i.e., species that hold 70 small local abundances) would be the ones arriving to the community at a later stage. 71 Although both the pioneering species and the core-satellite species hypotheses predict 72 similar trends of temporal increase in the richness and abundance of communities, their 73 predictions are markedly different in terms of temporal beta diversity patterns. The 74 pioneering hypothesis predicts a temporal turnover between pioneer and replacement 75 (i.e., non-pioneer) species (Connell and Slatyer 1977; Denslow 1980). Whereas the

76 core-satellite hypothesis expects that temporal changes in species composition should 77 be mostly driven by a non-random gain of species leading to increasing richness, since satellite species would be sequentially added to the community (Menéndez 1994). 78 79 Therefore, a preeminence of species replacement with time will be in accordance with 80 the pioneering hypothesis, whereas a preeminence of nestedness will provide evidence 81 for the core-satellite hypothesis. It follows that by studying temporal trends in beta 82 diversity components (i.e., turnover and nestedness sensu Baselga 2010), it should be 83 possible to elucidate the main process governing the (re)assembly of communities along 84 short and long time periods.

85 The Amazon rainforest is the largest contiguous and most biodiverse tropical 86 rainforest in the world, hosting a large proportion of known diversity (Peres et al. 2010). The spatial heterogeneity, seasonal regimes (e.g., pluviosity, river pulse, seasonal 87 88 humidity and temperature), and anthropic disturbances that characterize the Amazonian 89 region result in a complex mosaic of temporal and spatial changes in its communities 90 (Andresen 2002; Noriega et al. 2007; Korasaki et al. 2013). During the last 50 years, the 91 Amazon may have lost up to 29% of its forest cover due to an increase in deforestation 92 for the creation of cattle ranches, agriculture, and logging (Peres et al. 2010; INPA 93 2017). These activities have led to the appearance of a mosaic of habitat fragments 94 holding communities ongoing different successional stages, which provide an ideal 95 scenario for understanding how communities reassemble after disturbance (Braga et al. 96 2013; Franca et al. 2016; Cajaiba et al. 2017).

97 Dung beetles (Coleoptera: Scarabaeidae: Scarabaeinae) are an excellent study 98 system to explore temporal variations in biodiversity. They are sensitive to 99 environmental changes and respond quickly to habitat destruction, fragmentation or 100 isolation (Halffter and Arellano 2002; Barlow et al. 2007; Nichols et al. 2007). Indeed,

101 dung beetle abundance and richness are affected by human disturbance, usually 102 following a gradient of increasing negative effects when moving from natural forests to 103 secondary forests, plantations, and pastures (e.g., Howden and Nealis 1975; Barragan et 104 al. 2011; Braga et al. 2013). By burying and using dung as both a food and nesting 105 resource, dung beetles provide key ecological functions and services like nutrient 106 cycling, soil fertilization and aeration, seed dispersal, and biological pest control 107 (Andresen 2002; Bang et al. 2005; Nichols et al. 2008), which makes them important 108 for economy and human welfare (Losey and Vaughan 2006). Therefore, they have been 109 widely used as indicators for evaluating and monitoring spatial and temporal changes 110 and the impact of disturbances on natural communities (e.g., Davis et al. 2001; 111 McGeoch et al. 2002; Gardner et al. 2008; Otavo et al. 2013; Da Silva and Hernández 112 2018; Noriega et al. 2020).

113 In this study we evaluate both interannual and successional trends in the 114 diversity and composition of dung beetle assemblages in the Amazonian rainforest. To 115 do this, we use data from two dung beetle surveys with different temporal extents: one 116 including four successional stages with a temporal extent of 80 years (using a space-for-117 time substitution) and another including three consecutive years of sampling with two 118 successional stages. We address the following specific questions: 1) Do diversity and 119 assemblage structure change in short (interannual) and long (successional) temporal 120 scales? 2) How different are beta diversity patterns between these two temporal scales? 121 and 3) How do temporal changes affect the functional structure of dung beetle 122 assemblages at these two scales?

123

#### 124 Materials and methods

125 *Study sites* 

126 Surveys were conducted in two nearby Natural Reserves of the Amazon basin, Palmari 127 (Brazil) and Monilla (Colombia) (Fig. 1). The climate is humid with a mean rainfall of 2500 mm and a mean annual temperature of 25 °C. The area has a monomodal 128 pluviometric regime with two seasons: rainy – November to May (≥300 mm/month 129 130 average), and dry – June to October (≤100 mm/month average). The two Natural Reserves are contiguous to indigenous human settlements, so they are subject to 131 132 significant anthropic pressures that vary with their distance from the villages. These 133 human settlements affect the forest in different ways and intensities: from cutting small-134 medium areas for "slash and burn" shifting cultivation (see below), to extracting big 135 trees for construction and wood, to the collection of fruits and seeds, and occasionally 136 hunting for food.

137 The first study area, used to characterize long-term successional changes, was 138 located in the Palmari Natural Reserve (home of the Marubo indigenous community) in the municipality of Atalaia do Norte (4°17'1" S - 70°17'0" W, 77 m a.s.l.), 22 km from 139 140 the city of Benjamin Constant, in the Brazilian state of Amazonas (Fig. 1). Here we 141 sampled four types of habitats, representing a long-term successional recovery from human-induced disturbances: primary forest, with ~80 years of recovery after logging; 142 143 secondary forest, with ~30 years of recovery after logging; an old chagra, abandoned 144 approximately 5 years before sampling; and a *new chagra*, abandoned the same year of 145 the sampling. Here, "chagras" are forest areas that were completely cut and burn in a 146 "slash and burn" shifting cultivation regime used for different types of crops like 147 cassava, maize, rice, sugarcane, soybean, and plantain (see van Vliet et al. 2013).

148 The second study area, used to characterize both short-term yearly variations and 149 short-term successional changes, was located in the Monilla Amena Natural Reserve 150 (herein Monilla; home of Ticuna indigenous community), in the municipality of Leticia

151 (4°06'46" S - 69°55'52" W, 60 m a.s.l.), 9.5 km from the city of Leticia, in the Colombian state of Amazonas (Fig. 1). We assessed differences in short-term temporal 152 153 variations (i.e., between sampling years, see below) by sampling this locality during 154 three consecutive years. In addition, we evaluated whether these between-year 155 differences are similar in different moments of long-term community evolution by 156 doing these surveys in two different successional stages, secondary old forest and 157 secondary new forest, with approximately 20 and 10 years of recovery after logging, 158 respectively. This also allows us to evaluate whether the successional changes 159 eventually identified in the Palmari dataset hold up in other similar communities, rather 160 than responding to local idiosyncrasies.

161

# 162 *Dung beetle sampling and processing*

163 Dung beetle assemblages were sampled in Palmari in April (herein dry season) and 164 September (herein rainy season) 2009. Surveys were conducted using one linear 165 transect of 10 pitfall traps in each type of habitat (primary forest, secondary forest, old 166 chagra, and new chagra) with 50 m between traps (following Larsen and Forsyth 2005). 167 The pitfall traps were baited with 30 g of a 1:1 mixture of human and pig dung (a 168 combination that allows high quantities of bait with a high attraction level). In Monilla, 169 dung beetle assemblages were sampled during three consecutive years (2002, 2003, and 170 2004), in the same months and seasons than in Palmari (April, dry season; and 171 September, rainy season). In this case, because of the large extension of the forest, the 172 sampling was conducted using 10 linear transects of 10 pitfall traps in each type of 173 habitat (secondary old forest and secondary new forest) with 10 m between each trap 174 and 20 m between transects (see Noriega et al. 2007). The pitfall traps were baited with 175 30 g of carrion or human dung. Due to the short distance between traps, and in order to

make both datasets more comparable, the sampling unit was considered to be the trap in
the case of Palmari and the entire transect in Monilla (n=10 per habitat type and
season).

179 The model of the pitfall trap (plastic containers with the bait hanging above the 180 trap; see Noriega and Fagua 2009) and the time that the traps were active in the field (48 181 hours) were the same in both localities. The dung beetle specimens collected were 182 stored in 70% alcohol and identified to species level using several taxonomic keys 183 (Edmonds 1994; Genier 1996; Cook 2002; Edmonds and Zidek 2004, 2010; Camero 184 2010; Vaz-de-Mello et al. 2011; Cupello and Vaz-de-Mello 2013) and expert support. 185 After identifying and labelling all the specimens, the ones from Palmari were deposited 186 at the Natural History Museum of Los Andes University, Bogotá (EANDES), and those 187 from Monilla at the Natural History Museum of the Pontificia Javeriana University, 188 Bogotá (MPUJ).

189 All individuals were assigned to functional groups based on the dung beetle 190 functional classification proposed by Doube (1990), which combines their main food 191 relocation strategies (i.e., guilds) with the size of the individuals. We inferred dung 192 beetle guilds from the food relocation behavior known for each genus, assigning each 193 species to one of the three distinct guilds: paracoprids (or tunnelers), which dig under 194 the dung pile and make tunnels where they relocate a brood mass of dung; telecoprids 195 (or rollers), which construct a brood ball in the dung pile, roll it apart from the main 196 source, and bury it; and endocoprids (or dwellers), which nest and feed exclusively 197 inside the dung pats or build their nests right beneath the resource in the dung-soil 198 interface (Halffter and Matthews 1966; Halffter and Edmonds 1982; Cambefort and 199 Hanski 1991). Average body size was measured from the captured individuals as the 200 total length from the external border of the clypeus to the pygidium for each species,

201 with recourse to the literature for species with low abundances. These measurements 202 were performed using an electronic digital caliper (Powerfix - Z22855,  $\pm$  0.01 mm). 203 Each species was assigned to one of the three size categories (defined by the gaps in 204 species' body size; see Appendix S1): Large (>18 mm), Medium (10-18 mm), and 205 Small (<10 mm). We combined guilds and average body size into nine functional 206 groups (Appendix S1): large paracoprids, medium paracoprids, small paracoprids, large 207 telecoprids, medium telecoprids, small telecoprids, large endocoprids, medium 208 endocoprids and small endocoprids.

209

#### 210 Data analysis

211 We described dung beetle assemblages for each year, season, and type of forest by total 212 abundance, richness, and the abundance of each species and functional group. We 213 assessed inventory completeness as the percentage of the estimated total species 214 richness that were observed in each locality (Lobo 2008) to ensure that the surveys 215 attained a fair description of the studied assemblages. We estimated total species 216 richness with four nonparametric estimators (ACE, ICE, Chao 1, and Jacknife 1) that 217 are commonly used to characterize dung beetle communities for the ease of comparison 218 with other studies. All estimators were calculated with EstimateS v 9.1.0 (Colwell 219 2016).

We explored whether dung beetle species richness and abundance vary through time using generalized linear models (GLMs) on each one of these diversity attributes at the sampling unit level, as a function of successional stages in the Palmari dataset, and as a function of sampling year and successional stage in the Monilla dataset. In both datasets we also included the season as a predictor variable. We conducted this analysis separately for the diversity attributes related with species diversity and functional group diversity. We assumed a Poisson error distribution to fit richness and abundance
models, following a backward model selection based on Akaike Information Criterion
corrected by sample size (Burnham and Anderson 2002; AICc).

229 We described temporal variations in assemblage composition with a non-metric 230 multidimensional scaling ordination (NMDS) performed on matrices of sampling sites 231 by species and by functional groups. To avoid an excessive influence of rare taxa, 232 species with less than 5% occurrences were excluded from the NMDS analysis (see 233 McCune and Grace 2002). Data were subject to Wisconsin double standardization, 234 where each value is first standardized by the column maximum (i.e., for each species or 235 functional group), abundance is divided by its largest value in the surveys before being 236 standardized by the row total (i.e., for each trap), and species or functional group 237 abundance is divided by the total abundance of all species or functional groups in that 238 sampling unit. We used the Bray-Curtis index to measure the dissimilarity in species or 239 functional groups between the two datasets. We assessed the goodness of fit of the 240 ordination through the percentage of variance represented (see McCune and Grace 2002 241 for details). To select the main factor affecting assemblage composition, we performed 242 Spearman correlations between the NMDS axes resulting from previous analyses and 243 the potential explanatory variables (Matos et al. 2015). We also analyzed the 244 relationship between the NMDS ordination and the explanatory variables through vector 245 fitting. Then, those variables presenting significant correlations were overlaid in the 246 NMDS ordination (McCune and Grace 2002; Oksanen 2009).

247 Compositional variations in assemblage structure were evaluated based on the 248 nestedness and turnover components of beta diversity patterns using the indices 249 proposed by Baselga (2010):  $\beta_{\text{SIM}}$  (Simpson dissimilarity – spatial turnover),  $\beta_{\text{NES}}$ 250 (nestedness dissimilarity), and  $\beta_{\text{SOR}}$  (Sørensen dissimilarity – total Beta diversity).

These dissimilarity measures are additive fractions, so  $\beta_{SOR}=\beta_{SIM}+\beta_{NES}$ . Specifically, we measured nestedness and turnover for each pair of sampling units belonging to different time periods. Then, we measured beta diversity between consecutive periods and for each season and successional stage (in the case of Monilla dataset) independently. Finally, we used a Wilcoxon matched-pairs test to explore if nestedness and turnover measures differ between different pairs of time periods.

All analyses were performed in R v. 3.1.1 environment (R Development Core Team 2016), through different packages: *lsmeans* package (Lenth 2016) for the leastsquares means; the functions metaMDS and envfit of *vegan* package (Oksanen et al. 2013) for NMDS; and *betapart* package (Baselga and Orme 2012) in the case of beta diversity components.

262

## 263 **Results**

264 A total of 1,073 individuals were collected in Palmari, representing 34 species from 12 265 genera and six tribes (Table 1). The most abundant species in this site were Dichotomius 266 cf. boreus (Olivier, 1789), Dichotomius cf. fortestriatus (Luederwaldt, 1923), and 267 Onthophagus haematopus Harold, 1875, which altogether encompass about a third of 268 total abundance. In Monilla surveys rendered 822 individuals from 35 species belonging 269 to 13 genera and six tribes (Table 1). In this case, the most abundant species were 270 Sylvicanthon aequinoctialis (Harold, 1868), O. haematopus, and Canthon luteicollis 271 Erichson, 1847, together constituting more than 40% of all individuals. The percentages 272 of total species covered by the surveys were between 85.2 and 100% for Palmari and 273 73.6–98.9% for Monilla, indicating that sampling effort was sufficient to cover most of 274 the assemblage (Appendix S2).

## 276 Abundance and species richness

277 Regarding successional variations, both richness and abundance increased along successional stages. In Palmari, primary forest held the highest recorded species 278 279 richness and abundance (33 species, 14 of them unique, and 699 individuals), followed 280 by secondary forest (19 species, one of them unique, 249 individuals), old chagra (10 species, 99 individuals) and new chagra (7 species, 26 individuals); neither of the two 281 282 chagras had unique species (Table 1). In Monilla, richness and abundance were also 283 higher at the secondary old forest (31 species, 17 of them unique, and 671 individuals), compared with the secondary new forest (18 species, five of them unique, and 151 284 285 individuals) (Table 1).

When considering yearly variations, in general, abundance did not differ substantially between years within neither habitat (i.e., successional stage) nor season in the Monilla dataset; however, it varied in the rainy season of one of the years for both habitats, remaining nonetheless similar between years for both habitats during the dry season (Fig. 2b, Appendix S3). Richness did not vary significantly between years for each combination of habitat and season (Fig. 2d, Appendix S3).

292 Both seasons show similar patterns of variation in abundance and species 293 richness in both datasets (Fig. 2). In general, abundance and species richness decreased 294 from rainy to dry season, when dung beetle faunas seemingly tend to homogenize along 295 successional stages. During the dry season all stages presented low abundances in 296 Palmari, and the two chagras did not differ in neither abundance nor richness (Fig. 297 2a,c). These patterns were similar in Monilla, where the differences in abundance and 298 richness between the old and the new forest were lower, though significant during the 299 dry season (Figs. 2b,d, Appendix S4). Indeed, the interaction between season and 300 successional stage was the best supported model for richness, and a model describing 301 the joint effects of seasons and successional stages was in both datasets the best model302 for abundance (Appendix S3).

303

# 304 Assemblage composition

305 The main variations in species composition described by NMDM ordinations can be 306 summarized by axes related to successional and seasonal gradients in both datasets (Fig. 307 3a and 3b). At Palmari, the two first axes (final stress = 0.17, total variability explained 308 = 56%; Fig. 3a) identify a seasonal gradient between the rainy and the dry season, and a 309 successional gradient from the new chagra to the primary forest (see Appendix S4). In 310 Monilla, the obtained NMDS axes (final stress = 0.16, total variability explained = 52%; 311 Fig. 3b) were not related with variations throughout sampling years, but rather with 312 easily-identifiable successional and seasonal gradients like the Palmari dataset-313 although in this case these changes are summarized by the combination of both NMDS 314 axes (Appendix S4).

315 Nestedness and turnover (i.e.,  $\beta_{\text{NES}}$  and  $\beta_{\text{SIM}}$ ) markedly changed between long 316 and short temporal scales, showing that compositional changes are fundamentally 317 different for these two distinct time scales. Nestedness is influential in the long 318 successional gradient of Palmari (Figs. 4a,c), driven by a steep reduction of richness along the habitat disturbance gradient mainly due to the loss of rare species (while 14 319 320 species are unique to the primary forest, only one is unique for the new chagra). On the 321 contrary, in the shorter temporal scales of the interannual variations at Monilla, beta 322 diversity is mainly driven by a true turnover of species (Figs. 4b,d). Nonetheless, 323 compositional differences are attenuated in both datasets during the dry season, a period 324 when there were almost no differences of beta diversity between years (see Appendix 325 S5 for results regarding the dry season and total  $\beta_{SOR}$ ).

326

# 327 Functional diversity and structure

328 Both study sites presented all nine functional groups, defined according to resource 329 allocation behavior and body size, but their patterns of diversity along temporal changes 330 differ between datasets (Table 1). In Palmari, more than half of the functional groups 331 (small paracoprids and endocoprids, and all telecoprids) appear towards the latter stages 332 of the succession, while the composition of functional groups is maintained between 333 years in Monilla, corroborating that responses to succession are fundamentally different 334 from interannual variations. In general, paracoprids were dominant in Palmari, although 335 some functional groups appear with increasing successional maturity: large paracoprids 336 are exclusive of the primary forest, small telecoprids and small endocoprids appear only 337 on the secondary forest, and medium telecoprids and small paracoprids do so at the old 338 chagra (Table 1). This contrasts with Monilla, which is dominated by small paracoprids 339 and medium and small telecoprids, and only the former functional group appears in the 340 older successional stage, while the only large endocoprid species (Eurysternus velutinus 341 Bates, 1887) appears just in the secondary new forest (Table 1).

342 The abundance of all functional groups (except large telecoprids) increased 343 significantly with habitat succession in Palmari in the rainy season, while small and 344 large paracoprids and medium endocoprids were the only groups showing abundance 345 differences between successional stages in Monilla (Fig. 5j, 5l, 5q). Also, there was an 346 important decrease in abundance-accompanied with a functional homogenization of 347 the successional stages-from the rainy to the dry season in most functional groups in 348 both datasets, except for large telecoprids in Palmari and small telecoprids in Monilla 349 (Fig. 5f and 5j, respectively). Such successional and seasonal variability contrasts with 350 the interannual similarity in the abundance of functional groups found in the Monilla

dataset, which holds up for all groups except for medium telecoprids and endocoprids in
the secondary old forest during the rainy season (Fig. 5n and 5q). In fact, medium
telecoprids were the only functional group showing a distinct response, with changes in
abundance between years, successional stages, and seasons (Fig. 5n).

355 The NMDS ordinations describing variations in functional group composition 356 for both datasets identified similar gradients of joint successional and seasonal change, 357 from the young stages in dry season to old stages in the rainy season. In Palmari, the 358 two axes selected (final stress = 0.17, total explained variability = 61%) show an 359 increasing trend in all types of telecoprids and small endocoprids in both the rainy 360 season and the older stages of the successional gradient (Fig. 3c, Appendix S4). In 361 Monilla (two NMDS axes, final stress = 0.19, total explained variability = 52%), the 362 first axis selected was not correlated with any of the explanatory variables studied, but 363 the second was clearly related with both successional and seasonal variations, showing a 364 gradient of functional change from new forest assemblages in the dry season to old 365 forest assemblages in the rainy season (Appendix S4). Here, small and medium 366 telecoprids and small endocoprids are related with secondary old forest, while large 367 endocoprids and telecoprids and medium paracoprids are related with the rainy season 368 (Fig. 3d).

369

#### 370 Discussion

Our results show that changes in assemblage structure are fundamentally different between interannual and successional temporal scales, but they also show that there is some coherence between seasonal and successional variations. Amazon dung beetles present relatively similar trends towards higher abundance, richness, and functional diversity both in the older successional stages and during the rainy season. Another 376 important result is the almost negligible effect of interannual variations for the overall 377 diversity and structure of the assemblage, which is limited to significant species 378 turnover between years. In contrast, composition and diversity change in a more 379 estimated way throughout the succession, as variations between successional stages 380 show a clear nested structure. Indeed, changes in functional group structure are also 381 different between time scales, with drastic shifts throughout succession and seasonally 382 (with some groups disappearing from assemblages), which contrast with the 383 maintenance of functional structure at short temporal scales despite changes in the 384 identity of some of the species.

385

## 386 Successional variations in assemblage diversity, composition and structure

387 The increase of perturbation in Amazon forests generates poor dung beetle assemblages 388 in terms of both abundance and species richness (Barragan et al. 2011; Braga et al. 389 2013; Beiroz et al. 2017), a well-known trend that we also identify in both Palmari and 390 Monilla. This decay in diversity could be related to the loss of vegetation cover, 391 changes in soil texture, and the disappearance of vertebrate fauna that affects resource 392 availability and drastically changes the microclimatic conditions for dung beetles 393 (Andrade et al. 2011; Beiroz et al. 2017; Cajaiba et al. 2017; Ferreira et al. 2019). Many 394 low-intensity anthropogenic disturbances (e.g., reduced-impact logging, moderate 395 hunting, or logging roads) may also affect negatively dung beetle assemblage structure 396 (Bicknell et al. 2014; Feer and Boissier 2015; Edwards et al. 2017) and alter ecological 397 functions (Hosaka et al. 2014). This result is a clear nested pattern of compositional 398 change over the long temporal scales of succession, contrasting with the turnover that 399 defines the compositional variation at the short interannual time scales studied in the 400 Monilla dataset (Fig. 6; see below).

401 This nested pattern along successional stages suggests an effect of a non-random 402 species loss towards increasingly disturbed habitats. Indeed, some eurytopic species 403 (i.e., species with wide habitat requirements, such as Dichotomius mamillatus (Felsche, 404 1901), Dichotomius cf. boreus, or Sylvicanthon aequinoctialis) seem to be able to 405 pioneer the recovery of dung beetle assemblages after disturbance by persisting in the 406 majority of successional stages. In contrast, some stenotopic species (i.e., species with a 407 narrow habitat requirement, such as Canthon luteicollis, Deltochilum aff. pseudoparile 408 Paulian, 1938, or Onthophagus rubrescens Blanchard, 1843) are probably more 409 sensitive and restricted to the less disturbed habitats of the later successional stages. The 410 ability of colonizing habitats at different levels of disturbance may be the result of 411 several local factors that constrain assemblage structure, whose importance decreases 412 along the succession. These include loss of vegetation cover, abundance, diversity of the 413 available resources, and a significant change in microclimatic conditions related to open 414 areas (i.e., loss of humidity, increase of temperature, changes in soil texture, etc.; Beiroz 415 et al. 2017).

416 The anthropic pressure represented along the succession gradient also affected 417 the functional structure of the assemblages. Functional group richness is known to 418 decrease in disturbed areas as a result of changes in land use (Barragan et al. 2011, 419 Beiroz et al. 2018). In our study, some functional groups are absent or rare in the most 420 perturbed successional stages, especially those that include larger species. The loss of 421 large dung beetles is a common trend that occurs in several ecosystems and regions due 422 to the increase in anthropic perturbations (e.g. Lobo 2001, Tonelli et al. 2018). These 423 changes result in the existence of gradients in the composition of functional groups 424 found in our NDMS analyses. Interestingly, while the different functional groups of 425 telecoprids and endocoprids show a successional (and seasonal, see below) replacement,

426 paracoprids show no evident correlation with the succession. Under Hanski's (1982) 427 core-satellite hypothesis (see below), this could be attributed to many species with this 428 nesting behavior being "core" elements of the community that are present since the 429 beginning of assembly after the disturbance occurs.

430 Different responses of dung beetle functional groups to ecological pressures are 431 commonly found in the studies addressing the (generally negative) effects of habitat 432 disturbance on dung beetle diversity in the Neotropics (Andresen 2005; Horgan 2008; 433 Barragan et al. 2011; Braga et al. 2013; Nichols et al. 2013; Da Silva and Hernández 434 2015; Beiroz et al. 2017). Indeed, Audino et al. (2014) established that more than 18 435 years of recovery are needed to restore the functional diversity of dung beetle 436 assemblages in tropical rain forests. Here, it is essential to include the identity of species 437 and functional groups to understand restoration processes (Tonelli et al. 2020). 438 Following our results and taking into consideration the successional process in Palmari, 439 it is evident that, although some species from mature forest communities may endure 440 disturbance, full recovery of species richness, abundance, and functional group diversity 441 may take more than 30 years. Nonetheless, it is important to note that the analyses of 442 succession based on space-for-time substitution may underestimate the negative 443 consequences of human impact on local species diversity (Franca et al. 2016).

444

## 445 Interannual turnover

In general, our results indicate that the diversity, composition, and structure of dung beetle assemblages change significantly more between successional stages and seasons than between years. Indeed, beta diversity between years in Palmari is mainly due to the turnover component of compositional variation. This pattern of apparently random species replacement can be related with the existence of climatic or biotic factors (such

451 as unpredictable start of rains, strong variations in dry season rainfall, or a big reduction 452 in food resources) that might affect the phenological cycles or generate local population 453 declines and/or peaks. Besides the intrinsic environmental stochasticity of small time 454 periods, these factors can include: species of longer phenological cycles (i.e., supra-455 annual cycles; cycles that took more than a year for the larvae to become an adult), 456 explosive population outbursts (i.e., species that appear in high numbers during a brief 457 window of time associated with complex life history cycles), or the presence of rare 458 species that are active during very narrow windows of time (Wolda 1988; Ribeiro and 459 Freitas 2011; Kishimoto-Yamada and Itioka 2015). Other groups like butterflies have 460 life cycles longer than a year (Grøtan et al. 2014), evidencing the importance of 461 sampling during longer time periods.

462 The differences between interannual and successional changes in Amazonian 463 dung beetle communities have been seldom studied. The only study also addressing 464 both temporal scales that we are aware of also reports large interannual oscillations 465 during five years at an Amazonian primary forest (Beiroz et al. 2017), contrasting with 466 the relatively constant turnover between years we found in Monilla. However, in this 467 same study, dry season fauna in poorer years were comparable between primary forest 468 and disturbed areas (Beiroz et al. 2017), showing a pattern of biotic homogenization of 469 early and late successional stages with seasonality that coincides with the coherent 470 successional-seasonal gradient we found. Indeed, despite the relatively short time period 471 of our study in Monilla, the turnover between these three years could also be related to a 472 small forest recovery that translates into an increase in the number of species between 473 2002 and 2004 (from 30 to 34 species, 5 new species appear and one disappears), 474 similar to Beiroz et al. (2017) findings for their study area. Actually, the number and 475 composition of functional groups in the Monilla dataset do not vary between years,

which is congruent with the remarkable interannual stability in the relative abundance of functional groups found by these authors (Beiroz et al. 2017). Indeed, other studies exploring functional diversity trends after long temporal periods (i.e. 34 or 35 years; see Escobar et al. 2008 and Cuesta & Lobo 2019) did not find significant shifts in the composition of functional groups, reporting only changes in the proportion of individuals of each group, possibly connected with the effect of human activities.

482

## 483 Similarities between seasonal and successional gradients

484 Strikingly, the importance of the community enrichment along the successional process 485 is relatively similar to that of the seasonal gradient towards more abundant and diverse 486 assemblages in the rainy season. This common trend is particularly consistent for 487 functional structure and is likely related with the biotic homogenization promoted by 488 the low productivity characteristic of both disturbed habitats (see above) and the dry 489 season (see Hernández and Vaz-de-Mello 2009). Dung beetle assemblages are 490 characterized by a strong seasonality (e.g., Hernández and Vaz-de-Mello 2009; Andrade 491 et al. 2011; Lopes et al. 2011; Labidi et al. 2012; Viega et al. 2014), where the species 492 active during the dry season represent an impoverished sample (i.e., a subsample) of 493 those found during the rainy season (Agoglitta et al. 2012). In tropical rain forests, fruit 494 and green leaf production peaks during the hot-humid season, increasing the amount of 495 resources available for mammals and consequently the amount of excrements available 496 for dung beetles (Estrada et al. 1993). In contrast, during the dry season, dung 497 production is lower and its distribution is less spatially aggregated, with fewer dung 498 pads being available and becoming drier faster due to climatic conditions (Andresen 499 2005). However, some studies have found little seasonal variation in dung beetle 500 abundance, richness, and assemblage structure in the humid forests of the Amazon

region (Gardner et al. 2008; Korasaki et al. 2013). Following the results of Berioz et al.
(2017), such small seasonal variations may be an effect of the stochastic interannual
variations in climate, where climatically milder years allow the maintenance of richer
faunas throughout the whole year (see also Ferreira et al. 2019).

505

## 506 *A matter of core and satellite species and functional groups*

507 Our results point to the importance of core rather than pioneering species along the 508 successional gradients. The pioneering hypothesis predicts a strong temporal turnover 509 between early successional stages and later stages, following a sequence of species over 510 time (i.e., a deterministic process from few pioneer species to competitively dominant 511 non-pioneering species: "Pioneer hypothesis"; Connell and Slatyer 1977). In contrast, 512 dung beetle assemblages from Palmari indicate a negligible turnover and strong 513 nestedness along the successional gradient, with early stages acting as species-poor sub-514 samples of the later and richer stages (Fig. 6). This nested structure is consistent with 515 the core-satellite species hypothesis proposed by Hanski (1982) and found in the few 516 studies that analyzed this hypothesis in dung beetle assemblages (e.g., Menendez 1994). 517 This hypothesis states that communities are primarily formed by an initial assembly of 518 core species, which are widely distributed, locally frequent and/or abundant, and better 519 adapted to environmental changes; satellite species, which present a patchy distribution 520 and are locally rare and less adapted to environmental changes, are sequentially added 521 to the assemblage (Hanski 1982). This consistency with a core-satellite pattern suggests 522 that changes in dung beetle assemblage composition in the Amazon are not only a 523 consequence of deterministic successional processes but also of context-dependent 524 historical contingencies and neutral assembly of the species that are more abundant in 525 the landscape.

526 Although patterns of successional change are seemingly not deterministic, they 527 may have consequences for the functional structure of the assemblages. The species that 528 could be playing the role of "core species" are medium-large paracoprids (such as D. 529 mamillatus, D. cf. boreus, or D. cf. fortestriatus in the Palmari dataset) and medium-530 large endocoprids (e.g., Eurysternus spp.), whereas those seemingly acting as "satellite species" are small-large telecoprids (e.g., Canthon smaragdulus (Fabricius, 1781), 531 532 Deltochilum amazonicum Bates, 1887, or D. aff. pseudoparile) and small paracoprids 533 (e.g., Ateuchus and Uroxys spp.). Large paracoprids show the highest performance in 534 terms of dung removal in the ecosystem (Slade et al. 2007), so their presence from the 535 beginning would guarantee the maintenance of an important part of ecosystem functionality. In contrast, large telecoprids, which are more important for seed dispersal 536 537 (Andresen 2002; Vulinec 2002), may be more vulnerable to habitat transformations 538 (Lobo 2001; Nichols et al. 2013), appearing only in the later successional stages and 539 well-preserved patches. This is congruent with our results which show a perceptible 540 sequence in the dominance of different functional groups along successional stages, 541 shifting first from paracoprids to endocoprids, and then towards large telecoprids. These 542 sequences may relate to either species' specific ecological and physiological 543 requirements or some type of guild facilitation in terms of assemblage structuring, 544 which may affect ecosystem functionality. However, these hypotheses need to be tested 545 in the field with a mesocosm experimental design. Regardless of the existence of a 546 succession or not, a complete set of all functional groups is required in order to 547 maximize ecosystem functioning (Slade et al. 2007; Braga et al. 2013; Milotic et al. 548 2018).

549

550 Caveats

551 Although Palmari and Monilla are located nearby (approx. 30 km, see Fig. 1), they 552 show small differences in species composition. However, we believe that despite such 553 differences these two localities provide accurate representations of the variability in 554 central Amazonian dung beetle assemblages at different temporal scales. Note that 555 Korasaki et al. (2013) collected more species than us in a compositionally similar 556 locality near Palmari, possibly due to the lower numbers of traps (80 vs. ~450), sampling points (4 vs. ~14-18), and habitats (4 vs. 6) sampled in our study. In any case, 557 558 the differences between Palmari and Monilla study sites could be related to eventual 559 variations in the soil and forest composition, because they belong to different 560 geomorphological plates (Pebas formation at Palmari and Nauta formation at Monilla) 561 with different evolutionary histories (Higgins et al. 2011). Differences could also be 562 related with some distinctive methodological aspects between the two datasets, such as 563 sampling years and types of bait. We believe that the effects of these differences are 564 small, so the dissimilarities in the assemblages captured in this study correspond to the 565 different temporal factors that we studied, allowing the comparison in terms of the 566 magnitude of compositional change in the two datasets. Note that the differences 567 between sampling designs in the two study sites are due to the specific questions 568 addressed in each site (either successional or interannual changes). Despite these 569 differences, the clear contrast between successional and interannual scales found in this 570 study give us confidence about the reliability of our results.

571

## 572 Conclusions

573 In summary, dung beetle richness, abundance, and composition of functional groups
574 differ between short and long temporal scales. In long-term successional scales, dung
575 beetle assemblage structure changes significantly through time, following a nested

576 structure and causing many species and some of the functional groups to disappear in 577 highly disturbed areas. Seasonal changes may follow the same structure, at least to some 578 extent. On the contrary, in short interannual temporal stages richness and functional 579 group composition remain stable, as the turnover between years only affects the identity 580 of some species. Indeed, our results show that dung beetle assemblages in the Amazon 581 region are highly dynamic in time but with contrasting structural beta diversity patterns 582 depending on the processes involved in temporal changes. This evidence stresses the 583 importance of promoting long-term studies (especially long-term trapping) that include 584 temporal beta diversity analysis in order to elucidate insect community dynamics. The 585 inclusion of temporal dynamics in sampling protocols and monitoring studies could 586 favor the confidence and completeness of biodiversity inventories, positively affecting 587 conservation planning strategies (Hewitt et al. 2016). Moreover, it is necessary to study 588 the recovery process of ecological functions in restoration chronosequences, 589 incorporating functional trait data to fully understand the re-establishment and resilience 590 capacity of Amazon rain forests. This enhanced knowledge on the functional dynamics 591 of this hyperdiverse biome will eventually aid specific management and long-term 592 conservation strategies.

593

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Fig. 1 Map of the study region. Location of the two sampling localities: Palmari (BrazilNatural Reserve Palmari) and Monilla (Colombia - Natural Reserve Monilla Amena)in the Amazon region

**Fig. 2** Abundance (a,b) and richness (c,d) of the two sampling localities in the Amazon region: Palmari (Brazil - successional) and Monilla (Colombia - interannual) in each year (2002, 2003, 2004), sampling season (R: Rainy and D: Dry) and type of habitat (ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest, SNF: secondary new forest and SOF: secondary old forest). Different letters, above bars, indicate statistically significant differences (Wilcoxon's test; p<0.05) among years/seasons/forests. Error bars represent average±SE

**Fig. 3** Non-metric multidimensional scaling (NMDS) analyses of species (a,b) and functional groups (c,d) composition in Palmari (Brazil) and Monilla (Colombia) in the Amazon region. Season (Rainy and Dry) and type of habitat (ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest, SNF: secondary new forest and SOF: secondary old forest). Vectors represent significant correlations between assemblage composition and significant explanatory variables: Successional and Rainy (rainy season of the year). Functional groups and species code correspond to ones listed in Table 1 (P: paracoprids, T: telecoprids, E: endocoprids and Small (S): < 10 mm, Medium (M): 10-18 mm and Large (L): > 18 mm)

**Fig. 4** Beta diversity components ( $\beta$ sim [a,b] and  $\beta$ nes [c,d]) of Palmari (Brazil – successional) and Monilla (Colombia - interannual) during the rainy season, between types of habitat (ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest, SNF: secondary new forest and SOF: secondary old forest), and sampling years (2002, 2003, 2004) in the Amazon region. Different letters, above bars, indicate statistically significant differences (Wilcoxon's test; p<0.05) among years/seasons/forests

**Fig. 5** Abundance of each functional group (grouped by relocation food behavior -Paracoprids, Telecoprids and Endocoprids, and body size - Small < 10 mm, Medium 10 to 18 mm and Large > 18 mm) for Palmari (Brazil – successional) and Monilla (Colombia - interannual) in each year (2002, 2003, 2004), season (R: Rainy and D: Dry) and type of habitat (ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest, SNF: secondary new forest and SOF: secondary old forest) in the Amazon region. N/A: no species collected for that functional groups Different letters, above bars, indicate statistically significant differences (Wilcoxon's test; p<0.05) among years/seasons/forests

**Fig. 6** Beta diversity components: a) nestedness pattern in successional habitats (Palmari – Brazil, ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest); and b) turnover pattern in interannual comparisons (Monilla – Colombia) of dung beetles assemblage structure in the Amazon region. Numbers indicate the total number of species that were found only in each habitat or year

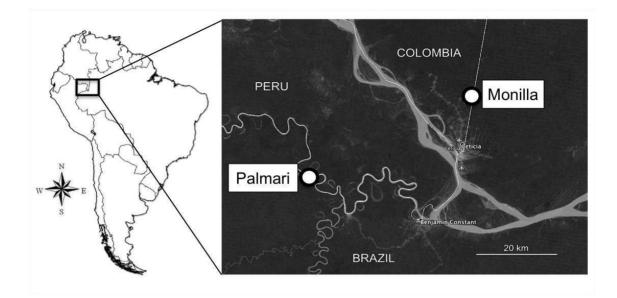
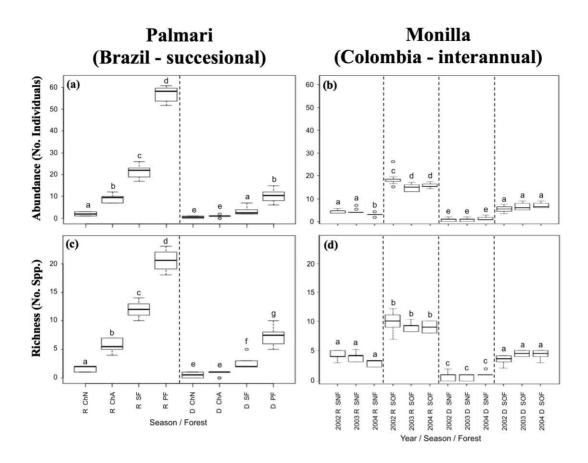
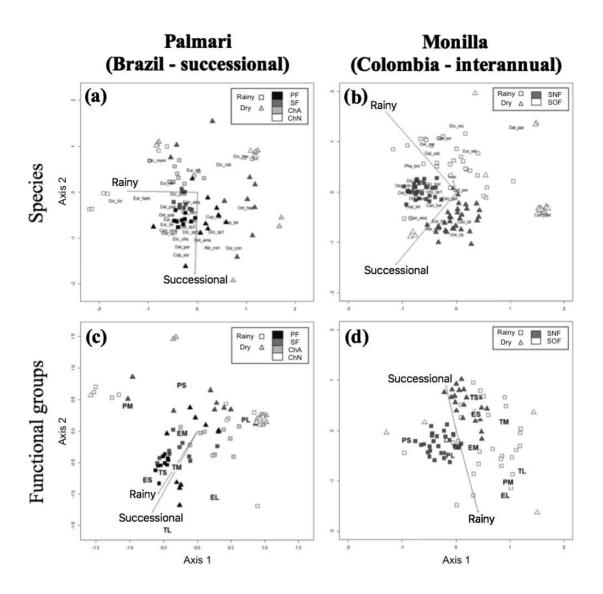


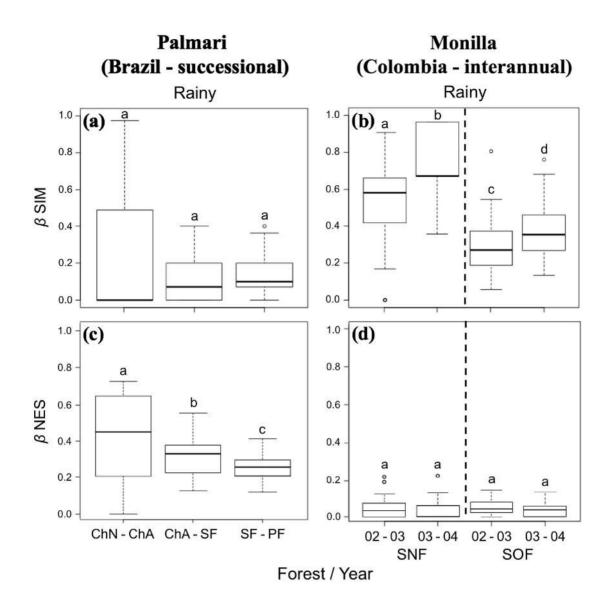
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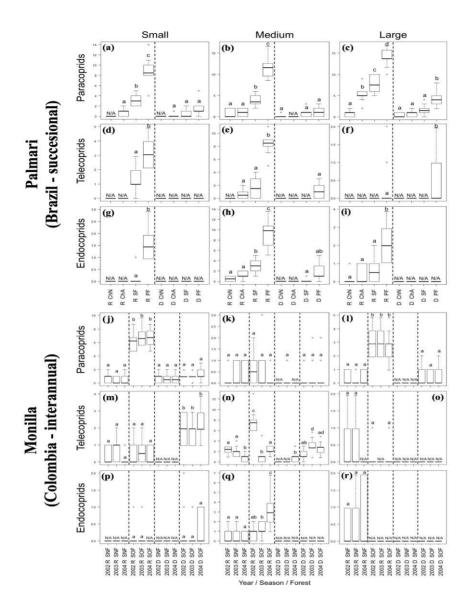
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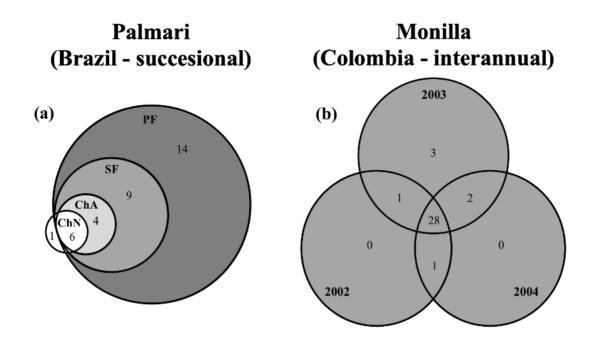
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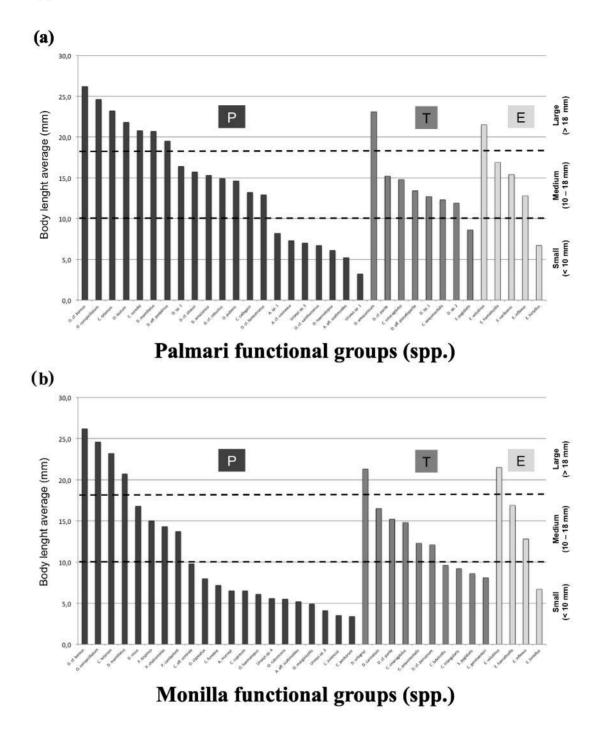
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**Table 1** Species list and abundance in Palmari (Brazil - successional) and Monilla (Colombia - interannual) in each type of habitat sampled (ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest) and year (2002, 2003, 2004) in the Amazon region. FG corresponds to functional groups built based on relocation food behavior and body size (P: paracoprids, T: telecoprids, E: endocoprids and Small (S): < 10 mm, Medium (M): 10-18 mm and Large (L): > 18 mm; Appendix S1); AS is the average individual length in mm; and Code is the species code (used in Fig. 5)

Tribe	Conus	Species	FG	AS	Code	Palmari			Monilla				
The	Genus	Species	гu	AS	Code	ChN	ChA	SF	PF	2002	2003	2004	Total
Ateuchini	Ateuchus	A. murrayi (Harold, 1868)	PS	6.5	Ate_mur	0	0	0	0	2	3	1	6
		A. cf. connexus (Harold, 1868)	PS	7.3	Ate_con	0	0	0	7	0	0	0	7
		A. aff. scatimoides (Balthasar,											
		1939)	PS	5.2	Ate_sca	0	0	0	3	0	1	1	5
		A. sp. 1	PS	8.2	Ate_sp1	0	0	0	4	0	0	0	4
	Uroxys	Uroxys sp. 1	PS	3.2	Uro_sp1	0	0	0	11	0	0	0	11
		Uroxys sp. 2	PS	4.1	Uro_sp2	0	0	0	0	12	13	15	40
		Uroxys sp. 3	PS	7.0	Uro_sp3	0	0	0	5	0	0	0	5
		Uroxys sp. 4	PS	5.6	Uro_sp4	0	0	0	0	0	2	0	2
Coprini	Canthidium	C. cupreum (Blanchard, 1843)	PS	6.5	Can_cup	0	0	0	0	1	1	2	4
		C. funebre Balthasar, 1939	PS	7.2	Can_fun	0	0	0	0	8	9	5	22
		C. gerstaeckeri Harold, 1867	PS	8.1	Can_ger	0	0	0	0	5	2	4	11
		C. aff. centrale Boucomont, 1928	PS	9.8	Can_cen	0	0	0	0	9	6	9	24
	Dichotomius	D. mamillatus (Felsche, 1901)	PL	20.7	Dic_mam	5	29	29	18	11	9	8	109
		D. nisus (Olivier, 1789)	PM	16.8	Dic_nis	0	0	0	0	1	3	3	7
		D. cf. boreus (Olivier, 1789)	PL	26.2	Dic_bor	7	20	29	81	14	12	16	179
		D. cf. fortestriatus (Luederwaldt,											
		1923)	PM	12.9	Dic_for	7	10	23	73	0	0	0	113
		D. cf. ohausi (Luederwaldt, 1923)	PM	15.7	Dic_oha	0	0	0	9	0	0	0	9
		D. cf. robustus (Luederwaldt, 1935)	PM	14.9	Dic_rob	0	0	7	12	0	0	0	19
		D. aff. podalirius (Felsche, 1901)	PL	19.5	Dic_pod	0	8	21	37	0	0	0	66
		D. sp. 1	PM	16.4	Dic_sp1	0	0	0	9	0	0	0	9
	Ontherus	O. pubens Génier, 1996	PM	14.6	Ont_pub	0	0	6	12	0	0	0	18
Deltochilini	Canthon	C. aequinoctialis (Harold, 1868)	ТМ	12.3	Can_aeq	0	6	10	58	81	35	48	238

		C. juvencus (Harold, 1868)	TS	3.5	Can_juv	0	0	0	0	3	7	1	11
		C. luteicollis Erichson, 1847	TS	9.6	Can_lut	0	0	0	0	23	19	24	66
		C. smaragdulus (Fabricius, 1781)	ТМ	14.8	Can_sma	0	0	0	4	1	0	1	6
		C. triangularis (Drury, 1770)	TS	9.2	Can_tri	0	0	0	0	0	1	0	1
	Cryptocanthon	C. peckorum Howden, 1973	TS	3.4	Cry_pec	0	0	0	0	0	2	1	3
	Deltochilum	D. amazonicum Bates, 1887	TL	23.1	Del_ama	0	0	0	8	0	0	0	8
		D. carinatum (Westwood, 1837)	TM	16.5	Del_car	0	0	0	0	9	9	1	19
		D. orbignyi (Blanchard, 1846)	TL	21.3	Del_orb	0	0	0	0	7	5	1	13
		D. cf. parile Bates, 1887	ТМ	15.2	Del_par	0	0	5	9	2	1	5	22
		D. cf. peruanum Paulian, 1938	ТМ	12.1	Del_per	0	0	0	0	15	14	9	38
		D. aff. pseudoparile Paulian, 1938	ТМ	13.4	Del_pse	0	0	0	14	0	0	0	14
		D. sp. 1	ТМ	12.7	Del_sp1	0	0	0	6	0	0	0	6
		D. sp. 2	ТМ	11.9	Del_sp2	0	0	0	6	0	0	0	6
	Scybalocanthon	S. pygidialis (Schmidt, 1922)	TS	8.6	Scy_pyg	0	0	13	30	2	1	2	48
Oniticellini	Eurysternus	E. caribaeus Herbst, 1789	EM	15.4	Eur_car	0	0	6	19	0	0	0	25
		E. hamaticollis Balthasar, 1939	EM	16.9	Eur_ham	3	3	7	34	14	13	35	109
		E. hirtellus Dalman, 1824	ES	6.7	Eur_hir	0	0	2	16	5	4	3	30
		E. inflexus (Germar, 1824)	EM	12.8	Eur_inf	2	8	20	57	0	3	0	90
		E. velutinus Bates, 1887	EL	21.5	Eur_vel	1	3	7	20	3	4	7	45
Onthophagini	Onthophagus	O. clypeatus Blanchard, 1846	PS	8.0	Ont_cly	0	0	0	0	4	2	2	8
		O. haematopus Harold, 1875	PS	6.1	Ont_hae	0	9	32	60	37	37	39	214
		O. marginicollis Harold, 1880	PS	4.9	Ont_mar	0	0	0	0	2	1	2	5
		O. rubrescens Blanchard, 1843	PS	5.5	Ont_rub	0	0	0	0	12	16	12	40
		O. cf. xanthomerus Bates, 1887	PS	6.7	Ont_xan	0	0	4	13	0	0	0	17
Phanaeini	Coprophanaeus	C. callegarii Arnaud, 2002	PM	13.2	Cop_cal	0	0	10	16	0	0	0	26
		C. suredai Arnaud, 1996	PL	20.8	Cop_sur	0	0	0	8	0	0	0	8
		C. telamon (Erichson, 1847)	PL	23.2	Cop_tel	0	0	11	15	6	6	8	46
	Gromphas	G. amazonica Bates, 1870	PM	15.3	Gro_ama	1	0	0	0	0	0	0	1
	Oxysternon	O. conspicillatum (Weber, 1801)	PL	24.6	Oxy_con	0	3	7	21	2	3	1	37
		O. lautum (Macleay, 1819)	PL	21.8	Oxy_lau	0	0	0	4	0	0	0	4
	Phanaeus	P. bispinus Bates, 1868	PM	15.0	Pha_bis	0	0	0	0	5	3	1	9
		P. cambeforti Arnaud, 1982	PM	13.7	Pha_cam	0	0	0	0	1	3	3	7
		P. chalcomelas (Perty, 1830)	PM	14.3	Pha_cha	0	0	0	0	2	3	0	5
				I	Abundance	26	99	249	699	299	253	270	1895
					Richness	7	10	19	33	30	34	31	55

## **Supplementary material**



Appendix S1 Definition of the functional groups based on guilds (P: paracoprids, T: telecoprids and E: endocoprids) and body size (Small (S): < 10 mm, Medium (M): 10-18 mm and Large (L): > 18 mm) for each locality: a) Palmari (Brazil - successional) and b) Monilla (Colombia - interannual) in the Amazon region

## Appendix S2 Observed (Sobs) and estimated percentage of species richness completeness calculated by four estimators (ACE, ICE, Chao 1 and Jack 1) for each locality (Palmari: Brazil and Monilla: Colombia), season (Rainy and Dry), habitat type (80, 30, 5 and 0 years and SOF: secondary old forest and SNF: secondary new forest), and years (2002, 2003, 2004)

C:4-	Casaar	II.ah.:4a4	Year	C		Estin		
Site	Season	Habitat	1 Cal	$S_{ m obs}$	ACE	ICE	Chao 1	Jack 1
		80 years		16	86.1	91.9	88.3	94.7
	Ś	30 years		8	87.6	93.5	89.5	95.0
(liz	Dry	5 years		4	89.2	94.7	90.2	98.7
Palmari (Brazil)		0 years		3	91.3	96.0	93.1	100
ari		80 years	-	33	85.2	90.0	87.5	91.7
Palm	γı	30 years		19	85.7	90.6	87.9	92.4
	Rainy	5 years		10	86.3	91.3	88.4	95.6
		0 years		6	88.9	93.4	89.1	99.2
			2002	11	78.3	92.1	87.5	95.6
		SOF	2003	13	77.5	91.7	86.5	97.5
	Ś		2004	14	79.2	92.5	88.3	98.1
-	Dry		2002	2	78.5	92.7	88.4	98.3
Monilla (Colombia)		SNF	2003	3	78.3	92.5	88.1	98.0
olor			2004	4	79.0	93.9	89.3	98.6
a (C			2002	23	73.6	91.8	85.4	94.5
onill		SOF	2003	21	74.0	91.9	85.7	95.7
M	γι		2004	21	73.9	91.4	85.5	94.4
	Rainy		2002	17	74.5	90.3	86.3	97.5
		SNF	2003	14	75.1	91.9	87.2	98.1
			2004	10	76.2	92.1	88.1	98.9

**Appendix S3** Richness and abundance models (Poisson) and AICc values for Palmari (Brazil - successional) and Monilla (Colombia - interannual). The explained deviance of the best model in terms of AICc is also provided

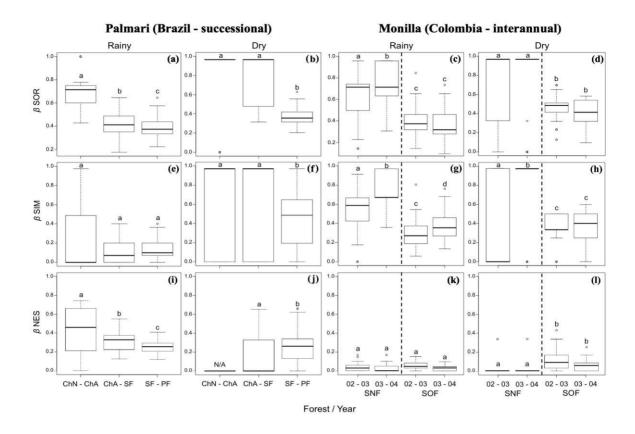
			Models	AICc		
	Richness	Mod 1	Season:ForestType	281.85*		
	models	Mod 2	283.52			
Palmari						
(Brazil)		Mod 1	Season:ForestType	322.11		
	Abundance	Mod 2 Season+ForestType		320.78*		
	models	Mod 3	Season	1300.24		
		Mod 4	4 ForestType			
			Explained deviance: 97.41%			
		Mod 1	Year:Season:ForestType	411.51		
		Mod 2	Year+Season:ForestType	405.73		
	Distance	Mod 3	Year:Season+ForestType	410.28		
	Richness	Mod 4	Year+Season+ForestType	409.73		
	models	Mod 5	Mod 5Season:ForestType			
M '11 .		Mod 6	Year	652.13		
Monilla			Explained deviance: 88.68%			
(Colombia)		Mod 1	Year:Season:ForestType	456.28		
		Mod 2	Year+Season:ForestType	458.59		
	A.1	Mod 3	Year:Season+ForestType	453.94*		
	Abundance	Mod 4	Year+Season+ForestType	461.24		
	models	Mod 5	Year:Season	807.10		
		Mod 6	ForestType	653.51		
			Explained deviance: 92.83%			

\* = lower AICc value

**Appendix S4** Spearman correlation (r) between non-metric multidimensional scaling ordination (NMDS) axes based on Palmari (Brazil - successional) and Monilla (Colombia - interannual) species and functional group records and explanatory variables. Squared correlation coefficient ( $R^2$ ) between the NMDS ordinations and explanatory variables are also presented for additional confirmation

	Palmari	Rainy	Successional	Monilla	Rainy	Successional	Year
NMDS based	r (Axis 1)	-0.49***	ns	r (Axis 1)	-0.53***	-0.38***	ns
on species	r (Axis 2)	ns	-0.65***	r (Axis 2)	0.69***	-0.40***	ns
	R <sup>2</sup> (NMDS)	0.27***	0.34***	R <sup>2</sup> (NMDS)	0.55***	0.42***	ns
NMDS based	r (Axis 1)	-0.31**	-0.40***	r (Axis 1)	ns	ns	ns
on functional	r (Axis 2)	-0.43***	-0.51***	r (Axis 2)	-0.63***	0.22*	ns
groups	R <sup>2</sup> (NMDS)	0.20***	0.38***	R <sup>2</sup> (NMDS)	0.33***	0.10**	ns

ns = non significant, \* *p*<0.05, \*\* *p*<0.01, \*\*\**p*<0.001



**Appendix S5** Beta ( $\beta$ sor,  $\beta$ sim,  $\beta$ nes) analysis for Palmari (Brazil – successional) and Monilla (Colombia - interannual) between types of habitat (ChN: new abandoned "chagra", ChA: old abandoned "chagra", SF: secondary forest, PF: primary forest, SNF: secondary new forest and SOF: secondary old forest), years (2002, 2003, 2004), and seasons (Rainy and Dry) in the Amazon region. N/A: no species available for comparisons. Different letters, above bars, indicate statistically significant differences (Wilcoxon's test; p<0.05) among years/seasons/forest