How forest amount and bioclimatic factors shape small mammal communities in Atlantic Forest fragments?

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Abstract:

Understanding how the mammalian diversity responds to anthropogenic disturbances on local and landscape scales is an urgent task. The Atlantic Forest biome, with only 12% of its original area remaining, still harbors great diversity of small mammals (Rodentia and Didelphimorphia), a key group that responds quickly to disturbances. Here, using the largest dataset of Atlantic Forest small mammals, we evaluate how forest amount and bioclimatic variables affect the non-volant small mammal diversity. For this purpose, we use 214 small mammal assemblages across the Atlantic Forest domain. Our results show that forest amount, with a positive relation, was the most important predictor explaining the diversity of small mammals in Atlantic Forest remnants. We also found that the bioclimatic variables (temperature and precipitation) can positively and/or negatively affect small mammal biodiversity, depending on the region analyzed. This is the first study that has assessed diversity across the entire Atlantic Forest biome, showing the importance of large-scale assessment and of forest amount and bioclimatic variables in shaping the diversity of small mammals regardless of the biogeographic context.

Keywords: precipitation, temperature, diversity, fragmentation, marsupials, rodents.

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Short title

Forest amount and bioclimatic shapes small mammal assemblies

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¹⁹ 19 Introduction

20	20	Community assembly at multiple spatial scales has been an essential topic in community ecology
21	21	(Fukami and Morin, 2003). Assessments of how species diversity is distributed has become
22	22	relevant due to provision of ecosystem services (Tscharntke et al., 2012; Whittaker et al., 2001).
23	23	Changes in environmental variables and habitat amount can occur together with anthropogenic
24	24	disturbances at multiple spatial and temporal scales (Chase et al., 2019). These have been shown
25	25	to influence community composition (Dambros et al., 2015; Delciellos et al., 2016; Gaston,
26	26	2000; Melo et al., 2017). The native Atlantic Forest in Brazil is experiencing significant habitat
27	27	loss and deforestation due to human activities such as cultivation of crops, grazing of livestock
28	28	and general overexploitation of natural resources (FAO, 2018). As a result, habitat loss and
29	29	deforestation have been identified as the main drivers of the decline in biodiversity in Atlantic
30	30	Forest regions (Cassano et al., 2014; Galetti et al., 2021; Pardini et al., 2017; Rios et al., 2021).
31	31	The Brazilian Atlantic Forest covers 1.5 million km ² along the country's coast to northeastern
32	32	Argentina and eastern Paraguay (Tabarelli et al., 2005). After hundreds of years of deforestation
33	33	due to anthropogenic disturbances (Joly et al., 2014; Lapola et al., 2014; Marcilio-Silva and
34	34	Marques, 2017), the Atlantic Forest has lost around 77% of its original vegetation cover
35	35	(Vancine et al., 2024). Today, remnants surrounded by several different anthropogenic matrices
36	36	can be found throughout the original Atlantic Forest biome (Lira et al., 2012). Even so, the
37	37	remaining Atlantic Forest fragments are noted for high biodiversity, including endemic species
38	38	(Mittermeier et al., 2011), and are considered global hotspots of biodiversity (Myers et al., 2000)
39	39	Within its rich biodiversity, these Atlantic Forest remnants harbor substantial diversity of
40	40	mammals, comprising more than 41% of all Brazilian species and most of the non-volant





41 41 mammals belong to the orders Rodentia and Didelphimorphia (Bovendorp et al., 2017; Paglia et
42 42 al., 2012).

43	43	Rodents and Didelphimorphs are composed of more than 200 species within Atlantic Forest
44	44	remnants (Bovendorp et al., 2017). They occupy several habitat strata (underground, ground and
45	45	arboreal) and have different habitat requirements (Abreu and De Oliveira, 2014; Estavillo et al.,
46	46	2013; Umetsu and Pardini, 2007). Compared to medium and large mammals, rodents and
47	47	marsupials are less vulnerable to extinction from anthropogenic disturbances due to their small
48	48	size and rapid reproductive rates (Gardner, 2008; Keesing, 2000; Pardini et al., 2010; Smith,
49	49	2001). However, many species are sensitive to fragmentation and loss of landscape connectivity
50	50	(Banks-Leite et al., 2014; Pardini et al., 2010), land use changes (Cassano et al., 2014; Ferreira et
51	51	al., 2020; Silva et al., 2019; Vieira et al., 2009), variation in food availability (Taitt, 1981; Taitt
52	52	and Krebs, 1981) and climate change (Loyola et al., 2012). Anthropogenic disturbances in
53	53	natural habitats lead to defaunation (Bovendorp et al., 2019; Cardillo et al., 2008; Fritz et al.,
54	54	2009; Galetti et al., 2015; Lira et al., 2012). Therefore, small mammals are ideal models for
55	55	investigating how diversity is shaped by bioclimatic variables and forest amount in the
56	56	biogeographic context.
57	57	The diversity of small mammals has been associated with the amount and composition of the
58	58	landscape elements (forest remnants and matrices) (Palmeirim et al., 2019; Pardini et al., 2010;
59	59	Vieira et al., 2018). Large forest remnant is associated with greater small mammal richness in
60	60	different biomes as well as for some assemblages in the Atlantic Forest (Palmeirim et al., 2019;
61	61	Pardini et al., 2010; Vieira et al., 2018). In addition, the matrix composition can positively or
62	62	negatively influence small mammals' richness and abundance, depending upon the species'

- ⁶³ 63 response to the matrix composition (Brady et al., 2011; Delciellos et al., 2016; Paise et al., 2020).





Studies have shown that factors such as temperature and precipitation can influence small mammal assemblages locally (Bergallo and Magnusson, 1999; Dambros et al., 2015; Ferreguetti et al., 2021; Graipel et al., 2006). Thus, the spatial variation of climate can shape small mammal assemblages by imputing dispersion limits (Dambros et al., 2015). Precipitation can also shape small mammal assemblages due to the positive effects on primary productivity (i.e., food availability) (Ferreguetti et al., 2021). Using the largest dataset of Atlantic Forest small mammals, we evaluate how bioclimatic variables and forest amount affect non-volant small mammal diversity in Atlantic Forest fragments. Furthermore, this is the first study aiming to ascertain how the effects of these factors differ among the tree main Atlantic Forest regions (northeast, southeast and south) (see detailed explanation in Material and Methods - Small mammal assemblages) because of geographic barriers (rivers and mountains) and heterogeneous species pools (Costa et al., 2000; DaSilva et al., 2015; Prance, 1982; Leite et al., 2016). The physical characteristics of remnants (size, forest amount and landscape heterogeneity) are frequently mentioned as the primary drivers of species diversity (da Fonseca 1989; Hortal et al., 2008; Banks-Leite et al., 2014; Magioli et al., 2021), and based on studies in the Atlantic Forest (Pardini 2004; Pardini et al., 2005, 2010; de la Sancha et al., 2020), we hypothesized that small mammal diversity would be more affected by forest amount than by bioclimatic variables. Thus, we expect small mammal diversity to increase as the forest amount expands due to the area effect, since a larger total forest amount will contain more individuals and for a given abundance distribution, this will imply more species available to colonize the forest area (Fahrig, 2013). Nonetheless, previous studies have demonstrated this relationship locally for small mammals in the Cerrado, Amazon Forest and some areas in the Atlantic Forest biome (Melo et al., 2017; Palmeirim et al., 2018; Pardini et al., 2005, 2010).



87	87	Also, we hypothesized that factors that affect the diversity of small mammals in the whole
88	88	Atlantic Forest will be different from ones that affect regional pools (de la Sancha et al., 2020),
89	89	since the remaining Atlantic Forest fragments (ranging from 3° S to 31° S in latitude) contain
90	90	diverse habitats and environmental conditions at, both on large and small scales, and this
91	91	heterogeneity can lead to factors influencing small mammal assemblages at different scales
92	92	(Fuentes-Montemayor et al., 2020).
93	93	Materials and Methods
94	94	Small mammal assemblages
95	95	We gathered information about small mammals (rodents and marsupials) available in the most
96	96	recent and complete studies (Bovendorp et al., 2017 and Figueiredo et al., 2017). We only
97	97	included studies providing appropriate geographic coordinates, richness, abundance, taxonomic
98	98	identification, sampling year and trapping effort (Fig. 1), and we removed any duplicated data.
99	99	We classified the dataset into three subgroups (northeast, southeast and south), creating a non-
100	100	homogeneous pool due to geographic barriers (mainly rivers, in particular to Rio Doce (Leite et
101	101	al., 2016)). These three main regions are widely accepted for systematic studies of small
102	102	mammals by taxonomists and also identify different areas of Atlantic Forest endemism,
103	103	highlighting the different and unique species composition across Atlantic Forest fragments
104	104	(Dalapicolla et al., 2021) and the factors that affect small mammals in these three main regions
105	105	(see Bovendorp et al., 2019).
106	106	Forest amount
107	107	For the landscape variables, we obtained land use information from MapBiomas (Souza et al.,
108	108	2020) (https://mapbiomas.org/en), which covers the period from 1985 to 2017 for Brazil. Given





that our small mammal assemblage data span several years of sampling (from 1986 to 2015) and that land use has changed substantially over time, we tried to obtain the most accurate information by extracting land use data for the average years of sampling. By matching the time of both sampling and land use data we reduced potential bias due to the temporal trends in land use change. Land use data for each year of sampling was extracted using QGIS 3.18 (QGIS Development Team, 2020). Our analysis focused on forest, pasture and agriculture land uses, which were the predominant categories of our sampling sites. We combined pastures and cropped areas into one category (farming) since they are both unnatural plantations with a production objective. We did not consider other land cover types, such as urban infrastructure, rivers, lakes, mangroves, beaches, dunes and mining areas, due to their low prevalence in our dataset. Using the geographic coordinates and sampling year provided in each study, we quantified the proportion of forest and farming amount around each sampling site within multiple spatial scales (100, 250, 500, 1000, 1500, 2000, 3000, 4000 and 5000 m radius). For example, a single sampled site can have a large amount (e.g., 90%) of forest cover at a 100 m radius because it is located in the middle of a fragment. However, this fragment can be isolated from other forest patches, so at 5000 m radius, the forest cover can be only 10%. Then, we selected the appropriate spatial scale for the subsequent analyses using the AIC value with the 'multifit' R package (Huais, 2018). For each variable, the package fits multiple models, i.e., one model for each scale. With a set of candidate models across all scales, the appropriate scale is selected according to the lowest AIC value (Huais, 2018). In our analyses, the forest amount was expressed as percentages. Since the amount of forest cover and farmed areas were strongly negatively correlated (-0.95), we only maintained forest amount. We only included assemblages with complete land cover use



132	132	information for the largest radius (5km), which we determined using Diva-Gis version 7.5
133	133	(Hijmans et al., 2012).
134	134	Bioclimatic variables
135	135	Unfortunately, bioclimatic variables were not available for each sampling site due to the lack of
136	136	information in published studies. In most cases, climatic information for sampling sites simply
137	137	does not exist. To standardize the metric for all sampling sites, we used the bioclimatic variables
138	138	from the nearest pixel accessed from the WorldClim database (www.worldclim.org) as a proxy
139	139	for local (site) conditions. To ensure that the variables used in our analyses were not highly
140	140	correlated and were statistically independent, we performed a correlation test with the 19
141	141	bioclimatic variables using the 'ENMwizard' package (Heming et al., 2019). After that step, we
142	142	just used seven bioclimatic variables (mean diurnal range, isothermality, maximum temperature
143	143	of warmest month, annual precipitation, precipitation seasonality, precipitation of wettest
144	144	quarter, and precipitation of coldest quarter).
145	145	Data analysis
146	146	We calculated the Shannon index for each small mammal assemblage using the function
147	147	'diversity' of the vegan package (Oksanen, 2017). In addition, we conducted Mantel correlation
148	148	analysis using the 'mantel' function of the vegan package (Oksanen, 2017) to certify that the
149	149	assemblages were not correlated spatially (t=0.969; df=3445; p=0.33) and/or temporally (t=-
150	150	0.338; df=3445; p=0.73). Note that the data assembled by us only allow a single diversity
151	151	estimate to be obtained for each study site.
152	152	We utilized the iNEXT package (Hsieh et al., 2016) to estimate the completeness of assemblages
153	153	through interpolation/extrapolation analysis. This approach was necessary due to variations in





sampling efforts and species pools in the available literature (Hsieh et al., 2016). For each assemblage, we provided species richness as input to the package, which then extrapolated the estimates to predict the total number of species and the percentage of the dataset represented. This rarefaction/extrapolation curve-based analysis allowed us to estimate the expected number of species as sampling effort increased and the proportion of the assemblage that was sampled. The analysis involves systematically increasing the number of individuals or samples collected from the community and calculating the cumulative number of observed species at each sampling effort level. By analyzing the rarefaction/extrapolation curves, we were able to make inferences about the species diversity and richness of the community. We estimated how many additional species would likely be observed if more sampling effort was applied and assessed the level of completeness of the actual sampling. Following these steps, were included 214 assemblages were included, with sample completeness of 80% or higher the analysis. Because diversity indices combine species richness and evenness in a single number, frequently they do not reflect differences in species richness, so biological or statistical interpretations can be misleading (Gotelli and Colwell 2001). Therefore, we tested the correlation between species richness and the Shannon index (r(212)=0.837, t=22.3, p<0.005) to make sure that our interpretation of the Shannon diversity index was straightforward (Fig. 2). To understand the relationship between the Shannon index and various predictor variables (forest amount, mean diurnal temperature range, isothermality, max temperature of the warmest month, annual precipitation, precipitation seasonality, precipitation of wettest quarter and precipitation of coldest quarter), we built linear models (LMs) using the function 'lm', because the variance of the residuals was constant. We generated a model selection table using the 'dredge' function and limited the number of predictor variables to two per model. We ranked the models using second





order AICc weights and built an average model (Anderson, 2008; Burnham and Anderson, 2002) using the 'model.avg' function, based on the weights of the models (>0.001). To check for uninformative parameters, we used the 85% confidence interval of each variable in the averaged models (Arnold, 2010). For model building comparison and averaging, we used the R package "MuMIn" (Barton 2019). Due to the different species pools in the Atlantic Forest fragments, we split the dataset into three subsets representing the regions (south, southeast and northeast - see above) and analyzed them separately. We used the average model for the subsets as well, under the same terms, to verify if the factors that affected the diversity of small mammals differed among all the Atlantic Forest remnants in the three regions. The landscape variable (forest amount) was used in linear and polynomial (quadratic) form (Edwards, 2002). The consideration of a quadratic effect can help identify potential threshold effects in the relationship between forest cover and the amount of forested area. Thresholds represent points at which the impact of forest cover on the variable undergoes a qualitative change. This can be crucial to understand critical levels of habitat for certain species of as small mammals (Pardini et al., 2010; Banks-Leite et al., 2014). The model was fitted using the normal distribution. We also assessed the statistical evidence of the variables (variable importance) by summing the AICc weights (W) of the model in which the explanatory variables were present (Burnham and Anderson, 2002). All analyses were performed with the R platform version 4.2.3 (R Core Team, 2023). **Results**

197 197 The final dataset included 214 assemblages, with 45,195 individuals from 104 species, and the
198 198 average number of individuals per assemblage was 186 (2 to 5,188 individuals; SD=429). The
199 199 most abundant species in these assemblages were *Akodon montensis* (n=6401), *Oligoryzomys*





200	200	nigripes (n=4826), Akodon cursor (n=3650), Didelphis aurita (n=3582) and Nectomys

²⁰¹ 201 *squamipes* (n=2236).

202 The appropriate spatial scale of our data was 5km radius, and all variables in the model selection 202 203 203 analysis were performed within this size. The most informative variables were forest amount 204 (quadratic, w=0.58, estimate=0.00004, p=0.001 and linear, w=0.41, estimate=0.005, p=0.003) 204 205 and maximum temperature of warmest month (w=0.42, estimate=-0.004, p=0.004), with negative 205 effects on diversity of small mammals, and precipitation of coldest quarter (w=0.27, estimate=-206 206 207 (0.001, p=0.007) and precipitation of wettest quarter (w=0.25, estimate=(0.001, p=0.008)) having 207 208 negative and positive effects on diversity of small mammals, respectively (Tab. 1; Fig. 3). The 208 209 209 other variables had importance of less than 0.05. 210 210 In the northeast region, none of the variables had statistical significance. In the southeast region, precipitation of wettest quarter (w=0.42, estimate=0.001, p=0.012), forest amount (linear, 211 211

- 212 212 w=0.34, estimate=0.005, p=0.004; quadratic, w=0.40, estimate=0.00004, p=0.003) and annual
- 213 213 precipitation (w=0.09, estimate=0.0005, p=0.019) had positive effects on the diversity of small
- 214 214 mammals (Tab. 2 and 3). On the other hand, maximum temperature of warmest month (w=0.50,
- 215 estimate=-0.006, p=0.001), mean diurnal temperature range (w=0.19, estimate=-0.009, p=0.014)
- 216 216 and isothermality (w=0.4, estimate=-0.025, p=0.046) had negative effects (Tab. 2 and 3).
- 217 217 In the south region, forest amount (linear, w=0.46, estimate=0.007, p=0.022; quadratic, w=0.54,
- 218 estimate=0.0007, p=0.016) and mean diurnal temperature range (w=0.29, estimate=0.008,
- ²¹⁹ 219 p=0.046) had a positive effect on the diversity of small mammals (Tab. 2 and 4).
- 220 220 **Discussion**





As hypothesized, our findings reveal that forest amount exerted a significant influence on the small mammal diversity in the of Atlantic Forest fragments studied. Additionally, our analyses demonstrate that bioclimatic variables (temperature and precipitation), have a discernible impact on the biodiversity of small mammals in this region. This is the first study assessing diversity across the entire Atlantic Forest biome, showing the importance of large-scale assessment and forest amount and bioclimatic variables in shaping the diversity of small mammals, regardless of the biogeographic context.

The diversity of small mammals increases in proportion to the amount of forest cover in the landscape (5km radius), across all levels of analysis, including per region and in all the Atlantic Forest remnants. This relationship has also been observed in previous studies of small mammals in the Cerrado and Amazon Forest biomes (Melo et al., 2017; Palmeirim et al., 2019), where amount of forest cover was associated with high species diversity (Pardini et al. 2010) based on the variety of habitat types and levels of habitat complexity (Umetsu et al. 2007). It is important to note that this positive correlation is not universal, and small mammal assemblages inhabiting small fragments (Pardini et al., 2005, 2010) or an inhospitable matrices (Estavillo et al., 2013) may be negatively impacted. But the significant relationships between the amount of forest cover and species diversity of small mammals in Atlantic Forest patches have also been observed in various studies in the last 20 years (Pardini et al. 2005, Püttker et al. 2011, Estavillo et al. 2013).

²³⁹ 239 Our analyses indicate that bioclimatic variables, specifically temperature and precipitation, can
²⁴⁰ 240 have a significant impact on small mammal diversity. Indeed, changes in temperature and
²⁴¹ 241 precipitation are expected to induce species range shifts and alter diversity and distribution
²⁴² 242 patterns in the Atlantic Forest, since the region is projected to become warmer and drier (Brown





et al., 2020, Almazroui et al., 2021). Based on this unique dataset, we observed a negative correlation between the maximum temperature of the warmest month and precipitation in the coldest quarter with small mammal diversity, while precipitation in the wettest quarter was found to have a positive relationship with small mammal diversity. Some studies investigating the relationship of temperature and precipitation with small mammal abundance and richness have reported contrasting findings (Barros-Battesti et al., 2000; Ferreguetti et al., 2021). Bioclimatic variables can indirectly influence small mammal diversity and abundance by increasing the availability of food resources (Bergallo and Magnusson, 1999; Graipel et al., 2006; Passamani and Ribeiro, 2009), leading to higher reproduction rates (Barros, 2013; Bonecker et al., 2009; Graipel et al., 2006). Such indirect impacts can help explain the complex and varied relationships between bioclimatic variables and small mammal biodiversity observed in different studies. Upon analyzing the three regions of the Atlantic Forest, we observed that none of the variables investigated in this study were statistically significant in the northeast region. The protected areas in this region are considered to have the "worst state of conservation" compared to other regions where Atlantic Forest fragments are present (SOS Mata Atlântica and INPE, 2023). The northeast region also has the lowest number of forest fragments (Dalapicolla et al., 2021), which are typically small (<50 ha) and isolated from one another (Ribeiro et al., 2009; Vancine et al., 2024). Due to these landscape characteristics, it is likely that much of the original mammal diversity in these areas has become extinct, since there are not enough large forest patches left to support their populations (de la Sancha et al., 2020). Furthermore, our investigation of the bioclimatic variables found no effect on small mammal diversity in the northeast Atlantic Forest region. This lack of effect may be due to the low variance in precipitation and temperature in the area (Carvalho et al., 2020).





For the southeastern region, we discovered that the maximum temperature of the warmest month and precipitation of coldest quarter had a negative impact on small mammal diversity. Conversely, precipitation of wettest quarter, and quadratic and linear forest amount were positively correlated. The temperature and precipitation are closely related to the geographic and latitude conditions of regions. In this region, the summer is generally characterized by high temperatures and abundant rainfall, which enhances food availability, and consequently small mammal abundance (Corrêa et al., 2017). Warmer temperatures and precipitation in the coldest periods can affect the availability of suitable microhabitats, food resources and water sources for small mammals, potentially leading to negative impacts on their population and diversity (de Castro and Fernandez 2004). Despite historical disturbances caused by agriculture and urban encroachment (Joly et al., 2014), the southeast region has the largest Atlantic Forest fragments (Vancine et al., 2024), which can promote small mammal conservation and reinforce the importance of forest amount in this group (Pardini et al., 2010). In the south region, mean diurnal temperature and the quadratic and linear forest amount has a positive relationship with small mammal diversity. The southern region has the lowest temperatures in the Atlantic Forest biome, reflecting also the geographic and latitude conditions of the region (Oliveira-Filho and Fontes 2000; Colombo and Joly 2010). Indeed, the mean diurnal temperature can have positive impacts on small mammal populations and diversity, not only on microhabitat, food and water resources (de Castro and Fernandez 2004), but also on furthermore in the individuals' activities and foraging (Kenagy 1973; Milling et al., 2017). Moreover, the southern region contains the second and third largest Atlantic Forest fragments (Vancine et al., 2024), contributing to small mammal conservation in these areas. The positive relationship between small mammal diversity and forest amount can be attributed to the legally





289 289 protected forest remnants in the region (protected areas), which have the capacity to support a
290 290 high richness of non-volant small mammal species within the southern Atlantic Forest. This
291 291 underscores the presence of areas characterized by high endemism among small mammal species
292 292 (Dalapicolla et al., 2021).
293 293 Conclusions

This is the first study assessing diversity in the entire Atlantic Forest biome (214 small mammal 294 294 295 295 assemblages), showing the importance of large-scale assessment. The results of this study also 296 reveal the importance of forest amount in the landscape for small mammal diversity in the 296 297 Atlantic Forest, regardless of the biogeographic region. Future research should focus on 297 298 298 investigating the variation in bioclimatic variables (temperature and precipitation), climate 299 299 change and the characteristics of the surrounding land cover, since these factors can also 300 300 contribute to small mammal diversity. By better understanding the factors that influence this 301 301 diversity, conservation efforts can be targeted more effectively to protect these important and 302 302 often overlooked members of the ecosystem (Asfora and Pontes, 2009; Dalmagro and Vieira, 303 303 2005; Lima et al., 2010; Püttker et al., 2008).

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628 628 **TABLES**

629 629

630	630	Table 1 Variables, model averaged estimates (Estimate), standard errors (SE), z values (z) an
631	631	p-value given by the Z-statistic (p) across the model-average linear mixed model (LM) and the
632	632	importance of each variable (W) to testing the variation of the small mammal Shannon index
633	633	with data from studies carried out in the Atlantic Forest. The importance value (W) was obtained
634	634	by the sum of the weights of each variable in the global models. The spatial scale used was 5km
635	635	radius.

636			Estimate	SE	Z	р	W
637		(Intercept)	1.671	0.750	2.227	0.026	
638		Max temperature of warmest month	-0.004	0.001	2.855	0.004*	0.42
639		Forest amount quadratic	0.00004	0.00001	3.265	0.001*	0.58
640		Precipitation of coldest quarter	-0.001	0.0002	2.695	0.007*	0.27
641		Forest amount	0.005	0.002	3.014	0.003*	0.41
642		Precipitation of wettest quarter	0.001	0.0002	2.673	0.008*	0.25
643		Isothermality	0.012	0.006	1.762	0.078	0.03
644		Mean diurnal temperature range	-0.001	0.003	0.193	0.847	0.01
645		Annual precipitation	-0.00001	0.0001	0.065	0.948	0.01
646	636	Significance code: *p<0.05					



- 647 637 **Table 2 -** Variables and importance of each variable (*W*) for each region. The importance value
- ⁶⁴⁸ 638 for small mammal diversity (Shannon index) was obtained by the sum of the weights of each
- ⁶⁴⁹ 639 variable in the global models. The spatial scale used was 5km radius.

650	Variables	W	W	W
651		Northeast	Southeast	South
652	Annual precipitation	0.13	0.09	0.04
653	Isothermality	0.44	0.04	0.25
654	Max temperature of warmest month	0.11	0.50	0.10
655	Mean diurnal temperature range	0.14	0.19	0.29
656	Precipitation of coldest quarter	0.17	0.01	0.12
657	Precipitation of wettest quarter	0.22	0.42	0.08
658	Forestry amount	0.16	0.34	0.46
659	Forestry amount quadratic	0.16	0.40	0.54

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662	642	Table 3 - Variables, model averaged estimates (Estimate), standard errors (SE), z values (z) and
663	643	the p-values given for the Z-statistic (p) across the model-average linear mixed model (LM)
664	644	testing the variation of the Shannon index for small mammals with data from studies carried out
665	645	in the Atlantic Forest (for the southeast dataset). The spatial scale used was 5km radius.

666		Variable	Estimate	SE	Z	р
667		Intercept	2.076	1.373	1.511	0.131
668		Max temperature of warmest month	-0.006	0.002	3.176	0.001*
669		Forest amount quadratic	0.00004	0.00002	2.934	0.003*
670		Forest amount	0.005	0.002	2.849	0.004*
671		Precipitation of wettest quarter	0.001	0.0004	2.515	0.012*
672		Mean diurnal temperature range	-0.009	0.004	2.446	0.014*
673		Annual precipitation	0.0005	0.0002	2.355	0.019*
674		Isothermality	-0.025	0.013	1.993	0.046*
675		Precipitation of coldest quarter	0.0008	0.0013	0.639	0.523
676	646	Significance code: *p<0.05				

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Mean diurnal temperature range

Precipitation of coldest quarter

Precipitation of wettest quarter

Significance code: *p<0.05

Annual precipitation

Max temperature of warmest month

Isothermality

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677 647 Table 4 - Variables, model averaged estimates (Estimate), standard errors (SE), z values (z) and 678 648 the p-values given for the Z-statistic (p) across the model-average linear mixed model (LM) 649 testing the variation of the Shannon index for small mammal with data from studies carried out 679 680 650 in the Atlantic Forest (for the south dataset). The spatial scale used was 5km radius. 681 Estimate SE Ζ р (Intercept) 1.019 0.394 682 0.405 0.694 683 Forestry amount quadratic 0.0007 0.0003 2.407 0.016* Forestry amount 0.007 0.003 2.288 0.022* 684

0.008

0.028

0.001

-0.004

-0.0005

0.0002

0.004

0.016

0.0007

0.003

0.0005

0.0003

1.995

1.737

1.434

1.275

0.957

0.073

0.046*

0.082

0.152

0.202

0.338

0.942

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⁶⁹³ 653 Figure 1 - Geographic distribution of the 214 small mammal assemblages in the Atlantic Forest
⁶⁹⁴ 654 fragments in Brazil. Gray shows the historical Atlantic Forest distribution, green shows the
⁶⁹⁵ 655 remaining patches.



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⁶⁹⁷ 657 Figure 2 - Relationship between the Shannon index estimates and the species richness of small
⁶⁹⁸ 658 mammals across sampled sites in northeast (NE), southeast (SE) and south (S) regions of the
⁶⁹⁹ 659 Atlantic Forest.







⁷⁰¹ 661 **Figure 3** - Relationship between Shannon index (small mammal diversity) and (A) forest



⁷⁰³ 663 precipitation of wettest quarter. The spatial scale used was 5km radius.

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