



Revisiting the ideas of trees as templates and the competition paradigm in pairwise analyses of ground-dwelling ant species occurrences in a tropical forest

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ABSTRACT

A challenge for studies on the organization of ant assemblages in forest ecosystems is to disentangle the causal effects of species occurrences. The structural and functional attributes of trees can act as environmental filters for ground-dwelling ant species influencing resource availability and the microclimate. The biotic interactions, especially competition, can work together with plant characteristics influencing ant species occurrences. To test the importance of tree traits and species interactions on co-occurrence patterns of ants, we collected ground-dwelling ants, with pitfalls and litter sampling, beneath the canopies of four tree species during the rainy and dry seasons in a restored forest. We used five predictors (tree identity, crown size, trunk circumference, litter depth, and leaves density) to model the presence probabilities of ants. Hence, we applied habitat constrained null models in pairwise analyses to disentangle the causal effects of ants co-occurrences. The random pattern predominated in the assemblages, making up 96% of all possible species pairs combinations. Overall, 50% of the species pairs that showed nonrandomness in the ant occurrences were interpreted as resulting from environmental filters, 36% as negative associations and 14% as positive associations. Additionally, we found that the effects of season and the sampling technique on the ant assemblages were also important. We suggest that the ideas of the trees as templates and the paradigm of competition are both useful for understanding pairwise occurrence patterns in ant assemblages, and can be tested using tree traits as predictors in ant species distribution models for running constrained null models.

Introduction

The influence of vegetation characteristics and biotic interactions are usually considered two important factors behind the organization of local ant assemblages. In forest ecosystems, the trees can be used as templates due to the variations in their structural and functional attributes that affects the resource supplies and modify abiotic conditions for the ant species (Andersen, 1995; Ríos-Casanova et al., 2006; Lopes and Vasconcelos, 2008; Donoso et al., 2010, 2013). The litter is a resource for the ground-dwelling ant species, providing food and shelter (Blüthgen and Feldhaar, 2010), and its composition and quantity can vary depending on the tree species (Ehrenfeld, 2003; Donoso et al., 2010; Mejía-Domínguez et al., 2011; Donoso et al., 2013). The differences among tree species in the crown characteristics, such as size, leaves density, and leaf phenology can create microsite variations in solar radiation and soil humidity (Mejía-Domínguez et al.,

2011), and is capable of influencing ant assemblages (Levings, 1983; Kaspari and Weiser, 2000). Therefore, the tree traits can be important for the environmental filtering of ground-dwelling ant species in local assemblages.

On the other hand, interspecific competition has been considered as the most important biotic interaction structuring ant assemblages (Parr and Gibb, 2010; Cerdá et al., 2013). Ants can compete for nesting space and food, with the species sometimes displaying a hierarchy of dominance (Pisarski and Vepsäläinen, 1989). Bait experiments have been used to measure the abundance and aggressiveness of ant species in order to assess the degree of species dominance (LeBrun, 2005; Baccaro et al. 2012). However, the presence of dominant species excluding other species from baits does not necessarily result in an assemblage structured by competition (Baccaro et al., 2012). Moreover, the competitive interactions between ant species can be affected by plants directly through differences in plant attractiveness for ant species

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(Dejean et al., 2019) and indirectly by the effects on microclimate (Fellers, 1989; Bestelmeyer, 1997, 2000). It is not simple to control experimentally the environmental factors that affect the community structure of organisms with a complex life cycle, such as ants. Therefore, a challenge for studies on the organization of ant assemblages is to disentangle the causal effects of species occurrences. Thus, the use of statistical modeling has become an acceptable approach to address the challenges of studying the rules that govern the assembly of communities (Gotelli, 2000; Gotelli and McCabe, 2002).

A null model is a statistical tool that randomizes data on species occurrences representing a control situation in the absence of a particular ecological mechanism (Gotelli and Graves, 1996). However it is challenging to associate nonrandom patterns of ant occurrences with particular ecological mechanisms. For example, differences in the abundance of ant species related to environmental conditions, such as temperature, and negative biotic interactions, such as competition, can produce the same segregated species distribution patterns (Ribas and Schoereder, 2002). The habitat constrained null models, introduced by Peres-Neto et al. (2001), involve modeling the presence/absence of species according to environmental variables to estimate probabilities of species occurrence; these estimates are then used in the generation of null communities and can facilitate the distinction between the processes responsible for the species co-occurrence patterns (e.g. Azeria et al., 2012; D'Amen et al., 2018; Le Borgne et al., 2018; Denis et al., 2019). Despite the fact that almost two decades have passed since the publication of Peres-Neto's work and that at the same time the number of studies with ant communities has greatly increased, the approach of constrained null models to elucidate distribution patterns in ant assemblages remains poorly evaluated. As far as we know, only Camarota et al. (2016) adopted the habitat constrained null models to the study of ant species co-occurrences. But, while they applied the approach to the study of canopy ants, here we will study the ground-dwelling ants, since it is still largely unknown which is the most important factor behind the organization of their assemblages.

In the studies of ground-dwelling ants, the sampling methods and seasonality can affect patterns of ant occurrences. Pitfall traps and Winkler extraction are among the most used techniques to sample ground-dwelling ants (Casimiro et al., 2019). We need to consider that the two techniques operate differently and tend to work better for different species and conditions. Pitfall traps capture surface-active ants, while the Winkler method can sample whole colonies in the litter and slow moving foragers (Bestelmeyer et al., 2000). The Winkler extraction method tends to collect species of a smaller size than those captured by the pitfall traps (Parr and Chown, 2001; Vargas et al., 2009). The pitfall traps are indicated for habitats with less litter, whereas the Winkler method can produce better results in habitats with more litter and not too dry (Bestelmeyer et al., 2000; Parr and Chown, 2001). Although the differences between the sampling methods are well recognized, some studies have stated that the co-occurrence patterns in ant assemblages was not affected by the sampling technique (Gotelli and Ellison, 2002; King, 2007), but others suggested that different techniques can lead to contradictory conclusions (Baccaro et al., 2012). On the other side, ant assemblages can be affected directly by daily and seasonal variations in temperature and rainfall (Levings, 1983; Silva et al., 2011). In tropical forests, heavy rainfall can inhibit the foraging of most ant species (Levings, 1983). Differences in ant abundance and richness throughout the year have been observed in neotropical habitats, with more species being sampled during the dry season (e.g. Jacquemin et al., 2016), during the wet season (e.g. Vargas et al., 2007) or with no differences between seasons (e.g. Pereira et al., 2016). Therefore, seasonality can potentially affect the co-occurrence patterns of ant species through effects on species richness and abundance.

In general, biotic interactions would be expected to be more important to community assembly on a local scale, while environmental filters would operate on a regional scale. This is because environmental conditions tend to be spatially structured (Kristiansen et al., 2012). Here, we focus on ground-dwelling ant co-occurrence patterns in a reforested area. Ecological restoration projects are a great opportunity to investigate patterns and mechanisms in the process of assembling communities (Young et al., 2001). Reforestation areas represent environments with a known history, which can facilitate the design of field experiments. We hypothesized that even in a small spatial scale, ground-dwelling ant assemblage would be structured by both, biotic interactions and environmental filters. We worked with the premise that the species distribution on a small spatial scale would not be limited by dispersion, because most ground-dwelling species in tropical forests can frequently relocate their nests (McGlynn, 2006). We hypothesized that the differences in the attributes of the trees, and canopy openness, in a forest habitat, would act as environmental filters influencing ant species co-occurrence. We also hypothesized that competition between ant species would be detected through segregated species distribution patterns on a local scale, because competition tends to be an important factor structuring local ant assemblages (Cerdá et al., 2013). We adopted the Peres-Neto et al. (2001) approach to disentangle the biotic interactions and environmental filters, applying habitat constrained null models. Instead of analyzing the structure of the entire community, we adopted the pairwise analysis (Veitch, 2014; Ellwood et al., 2016). This choice occurred because the analyses based on the entire data matrix patterns can hide important interactions between pairs of species (Camarota et al., 2016). We also checked if the results were dependent on the ant sampling techniques and seasons.

Material and methods

Study area

The study was carried out at the Reserva Ecológica de Guapiaçu (REGUA), located in the municipality of Cachoeiras de Macacu, State of Rio de Janeiro, southeastern Brazil (22° 27'3.41" S, 42° 46'28.17" W) (Fig. 1). The region has a rainy summer and a dry winter. Annual rainfall is 2,050 mm, ranging between 60 mm in July and 338 mm in February, with January being the hottest month (25.3 °C) and July the coldest (17.9 °C) (Azevedo et al., 2018). REGUA is in the Atlantic Forest biome, presenting phytophysiognomies characteristic of Dense Ombrophilous forest, with altitudes varying from 20 to 2,000 m above sea level (Veloso et al., 1991). The forest cover of the low areas is quite fragmented, being interspersed with pastures, agricultural areas and areas under forest restoration (Rocha et al., 2007; Almeida-Gomes and Rocha, 2014; Azevedo et al., 2018).

Experimental design

Our study was conducted in a 12-years aged forest restoration area, 3.4 hectares in size (Azevedo et al., 2018). Herbicide pasture grass suppression was carried out before plowing the soil for planting forest seedlings. Pioneer, secondary and climax tree species were used in the forest restoration. Chemical control of leaf-cutting ants was carried out for two years since the tree planting (Azevedo, 2012). Three years after the planting the studied area had 73 tree species (Azevedo et al., 2018).

We chose four tree species that are common in the Atlantic Forest of Rio de Janeiro state and are frequently used in forest restoration (Nóbrega et al., 2008; Sansevero et al., 2011; Azevedo et al., 2018). Our samples were distributed in 37 trees, with a minimum distance of 10 m

between individuals and with little or no crown overlap with surrounding trees. The tree individuals were distributed in the following species: *Guarea guidonia* (L.) Sleumer (GG) (n = 11); *Inga edulis* Mart (IE) (n = 9); *Nectandra membranacea* (Swartz) Giseb. (NM) (n = 8); and *Piptadenia gonoacantha* (Mart) J. F. Marc. (PG) (n = 9) (Appendix 1, Supplementary material). We chose trees with a circumference at breast height greater than 30 cm to ensure that we measured the species-specific effects of trees on ant colonies, reducing the possibility of neighboring trees effects (Lameira et al., 2019). To ensure that most of the litter was composed of material from the focal tree, all ant samples were collected at a distance of 1 m from the tree trunk (Donoso et al., 2010). In the study area, the tree species had contrasting attributes (Table 1).

The sampling of ants

Ant collections were carried out in 2018, in periods of high and low rainfall (rainy and dry season, respectively). We used two techniques to sample ground-dwelling ants: pitfall traps and litter collection for

mini-Winkler extraction. The pitfalls consisted of plastic cups with a capacity of 400 ml, 10 cm in height and 7.5 cm in diameter. The cups were buried with the upper part at ground level and half-filled with salt water (24 g/L) with a drop of detergent. Each sampling point received a pitfall for 48 hours. The litter samples were collected within a quadrat of 0.25 m² placed on the ground. The sifted litter samples were suspended in mini-Winkler sacks, during 48 hours, to extract the ants (following Bestelmeyer et al., 2000).

All the collected ants were taken to the laboratory for identification. The identification was carried out using the keys found in the literature (Kempf, 1951; Mayhé-Nunes and Meneguete, 2000; Longino, 2007; Sosa-Calvo and Schultz, 2010; Dash, 2011; Feitosa, 2011; Baccaro et al., 2015; Johnson, 2015; Ješovnik and Schultz, 2017).

The environmental variables

The circumference of the tree trunk and the diameters of the crown of each focal tree were measured with a tape; for the diameters, we took

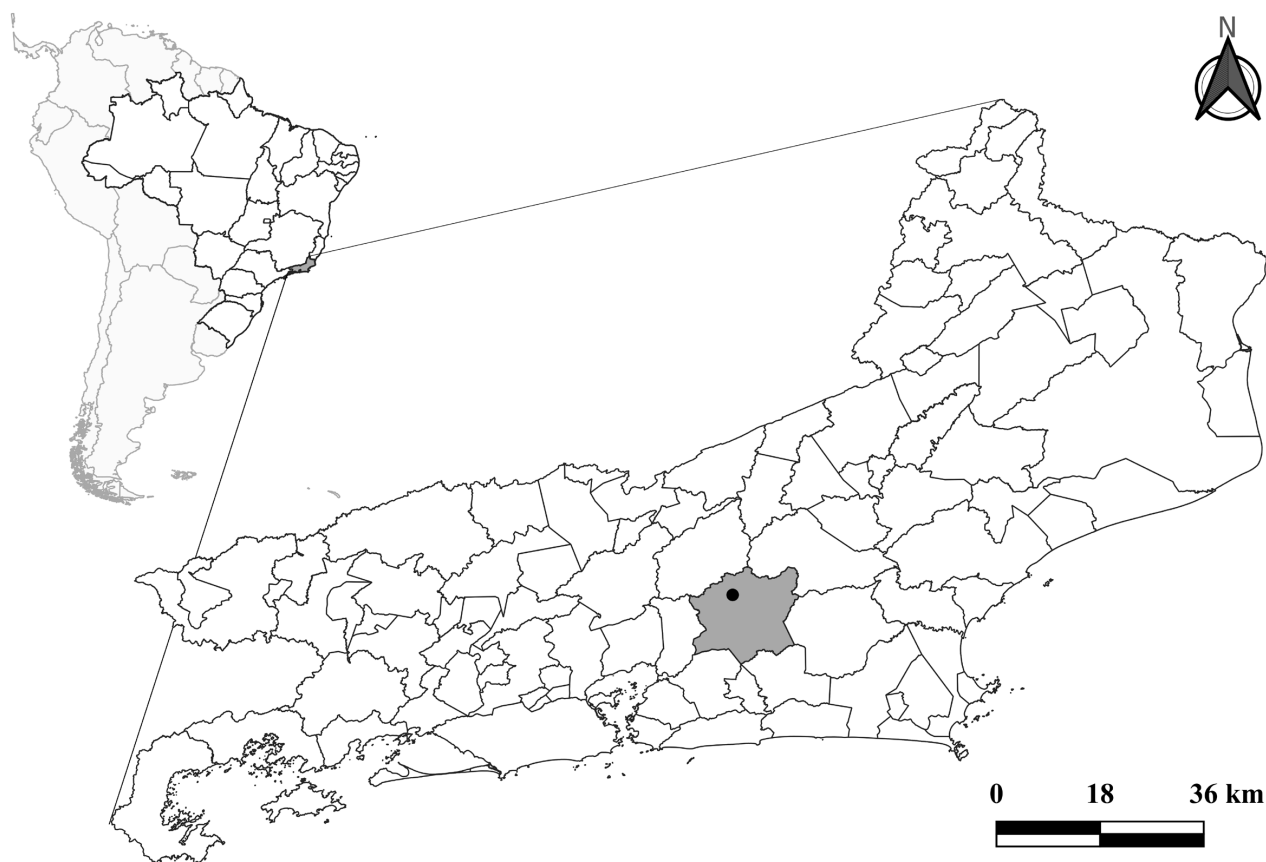


Figure 1 Location of REGUA in the State of Rio de Janeiro, Brazil.

Table 1

Attributes of tree species in the study area at REGUA.

Tree identity	Family	Leaf-loss strategy	Crown size	Trunk circumference	Litter height
<i>Guarea guidonia</i>	Meliaceae	Perenifolius	Small	Small	Medium
<i>Inga edulis</i>	Fabaceae	Perenifolius	Large	Large	High
<i>Nectandra membranacea</i>	Lauraceae	Perenifolius	Small	Large	High
<i>Piptadenia gonoachanta</i>	Fabaceae	Deciduous	Small	Large	Low

two measurements for each crown. The size of the crown was estimated using the ellipse model, which consists of multiplying the minor axis by the major axis and then by the value of pi. The leaf area indexes (LAI) were calculated through hemispheric photography, which consisted in the photograph of the canopy at the height of one meter with the camera facing north (Denslow and Guzman, 2000). Subsequently, the photos were analyzed in Hemiview 2.1 software, to obtain the LAI values. At each sampling site, the litter height was measured at four points with the aid of a ruler. The litter depth values used in the subsequent analyzes was an average of the four values. The measuring of LAI and litter depth was carried out in the rainy and dry seasons.

Data preparation

Ant species presence/absence matrices were built to analyze the community's assembly pattern according to sampling technique and season; this means that we built four different matrices (2 techniques x 2 seasons). We analyze only the most frequent ground-dwelling ant species sampled in the area, i.e. those which were recorded in more than 10% of the samples for each technique and season. Fungus-growing ants were also removed because of their food diet specialization. The presence/absence matrices are generally used to avoid overestimating the abundance of certain species that may have a large representation of individuals in certain samples (Longino et al., 2002; Gotelli et al., 2011).

SDM analysis

We obtained probability values for each ant species in the samples for the four presence/absence matrices by fitting species distribution models (SDM). We used five environmental predictors (tree identity, crown size, trunk circumference, litter depth, and the leaf area index-LAI). We applied two modeling techniques indicated for presence/absence data (Qiao et al., 2015): Generalized Linear model and Random Forest, with 10 replications each. The resulting 20 projections were averaged based on AUC (area under the curve) statistics (values are provided in Appendix 2, supplementary material) to implement a single ensemble estimate of the probability of presence for each ant species in each sample. The analyses were performed using the SDM package for R (Naimi and Araújo, 2016). We used the probability values of ant presence in the constrained null models described below.

Null model analysis

The matrices containing ant species presence/absence data were subjected to co-occurrence analysis using the pairwise C-score index. The pairwise C-score index quantifies the degree of co-occurrence of each pair of species for the locations in the matrix (Stone and Roberts, 1990); it is appropriate for detecting non-randomness in the community, in addition to being less susceptible to type I and II errors (Gotelli, 2000). The pairwise C-score index is calculated for each pair of species using the equation $C_{ij} = (r_i - S_{ij})(r_j - S_{ij}) / (r_i * r_j)$, where r_i represents the total sites that species i occurs in the matrix, r_j is the total sites for species j , S_{ij} is equivalent to the total number of sites that both species occur simultaneously. One C-score value is calculated for each species pair (observed index). The observed pairwise C-score is compared to a mean of simulated indices. To calculate the mean and variance of simulated indices, each of the four presence/absence matrices was randomized 1000 times.

We followed the methodology proposed by D'Amen et al. (2018) running unconstrained and constrained null models. The unconstrained model uses only presence/absence matrix, whereas the constrained null

model uses the presence/absence matrix with the probability values of ant presence calculated by the SDMs. Given that we worked in a small spatial scale, in an area relatively homogeneous, we considered that ant species wouldn't be limited by dispersion. We adapted the approach of D'Amen et al. (2018) in the following way: if we find a result for the pairwise occurrence that is nonrandom for the unconstrained model and random for the constrained model, it is interpreted as caused by the environmental filter. If the result is segregation for both models, it is interpreted as a negative association; and if the result is aggregation for both models, it is interpreted as a positive association. We used the algorithm fixed-equiprobable (FE) for the randomizations because it is less prone to Type I and II errors (Gotelli, 2000). In the FE algorithm, the number of ant species remains (fixed lines), because co-occurrence tests are very sensitive to variation in species occurrence frequencies (Gotelli, 2000), but ant occurrence in each sampling point is equiprobable (equiprobable columns) (Gotelli and Ellison, 2002). This algorithm is indicated for samples collected in homogeneous habitats (Gotelli, 2000), and can be biased towards detecting positive associations (Azeria et al., 2012). The analyzes were performed using the Open Source R version 3.6.1 statistical software (R Core Team, 2019), following the scripts provided by D'Amen et al. (2018).

Results

In the study, 82 ant species/morphospecies were collected, corresponding to 22 genera. However, after the removal of rare (less than 10% of records), arboreal species, and fungus-growing ants, the co-occurrence analyzes, by null models, were restricted to 459 records of 27 species/morphospecies of ants, corresponding to 11 genera and two trophic guilds (predators and omnivores). The most frequent ant species included *Pheidole subarmata*, *Hypoponera parva*, *Linepithema neotropicum*, *Rogeria scobinata*, and several species of *Solenopsis* and *Strumigenys*. *Pheidole subarmata* was collected either with pitfalls and Winkler samples, in the rainy or dry seasons. *Hypoponera parva* and *Rogeria scobinata* were both more frequent in Winkler samples, with the former increasing their presence in the dry season. *Linepithema neotropicum* was predominantly collected by pitfalls in the dry season; and species of *Solenopsis* or *Strumigenys* showed varied patterns. Pitfalls in the rainy season had the lower proportion of species records and richness (15.7%, 8 species), followed by pitfalls in the dry season (24.8%, 13 species), Winkler samples in the rainy season (25.9%, 12 species), and Winkler samples in the dry season (33.6%, 17 species). The species records increased by 58% for pitfall samples and 29% for Winkler samples during the dry season. Five ant species were present only in the pitfall data matrices, whereas nine species were exclusive of Winkler data matrices. Four species were exclusive of rainy season data matrices, whereas 11 species were exclusive of dry season data matrices (Table 2).

The random pattern of distribution of species pairs predominated in the assemblages, making up 96% of all possible species pairs in the four data matrices ($n = 351$ pairs) for the unconstrained null model. Fourteen species (52%) were involved in at least one nonrandom pairwise species association (significant C-scores values are provided in Appendix 3, supplementary material). The presence probabilities for the 14 species involved in nonrandom associations varied within and among tree species (Fig. 2). There was a reduction of 50% in the number of nonrandom species pairs for the habitat constrained null model in relation to the unconstrained model. The proportions of nonrandom pairs varied between seasons and sampling techniques (Table 3).

The numbers of nonrandom pairs in the unconstrained model were seven aggregated and seven segregated. For the constrained model, we observed two aggregated and five segregated pairs. All the segregated pairs, under either the constrained or unconstrained models, involved

Table 2

Species composition and records (>10% of the samples) of ant assemblages sampled at REGUA during the rainy and dry season, with two sampling techniques, and used in null model analyses.

Species	Pitfall		Winkler		Total
	Rainy season	Dry season	Rainy season	Dry season	
<i>Brachymyrmex admotus</i> Mayr, 1887		7			7
<i>Ectatomma permagnum</i> Forel, 1908		4			4
<i>Hypoponera parva</i> (Forel, 1909)			6	18	24
<i>Hypoponera</i> sp3				5	5
<i>Linepithema neotropicum</i> Wild, 2007	5	14		6	25
<i>Nylanderia fulva</i> Mayr, 1862	5			4	9
<i>Nylanderia steinheili</i> Forel, 1893		8		5	13
<i>Pachycondyla harpax</i> (Fabricius, 1804)				6	6
<i>Pachycondyla striata</i> (Smith, 1858)	4				4
<i>Pheidole</i> sp1		4			4
<i>Pheidole</i> sp2		5			5
<i>Pheidole</i> sp5				4	4
<i>Pheidole subarmata</i> Mayr, 1884	24	26	20	14	84
<i>Rogeria germani</i> Emery, 1894				7	7
<i>Rogeria scobinata</i> Kluger, 1994			19	19	38
<i>Solenopsis invicta</i> Buren, 1972	5	9		5	19
<i>Solenopsis</i> sp1			7		7
<i>Solenopsis</i> sp2		4	5	21	30
<i>Solenopsis</i> sp3	11		14		25
<i>Solenopsis</i> sp4		16		9	25
<i>Solenopsis</i> sp5	14	6	6		26
<i>Solenopsis</i> sp7		7	12		19
<i>Strumigenys denticulata</i> Mayr, 1887			4	14	18
<i>Strumigenys eggersi</i> Emery, 1890	4	4	10	7	25
<i>Strumigenys louisianae</i> Roger, 1863				5	5
<i>Strumigenys subdentata</i> Mayr, 1887			9		9
<i>Wasmannia auropunctata</i> Roger, 1863			7	5	12
Total	72	114	119	154	459

Table 3

Percentages of nonrandom species pairs in the unconstrained (unc) and constrained (cons) models.

Sampling method	Season	Model	% of nonrandom species pairs
Pitfall	Rainy	unc	10.7
		cons	3.6
	Dry	unc	1.3
		cons	1.3
Winkler	Rainy	unc	1.5
		cons	0.0
	Dry	unc	5.9
		cons	3.7

species in the same genus or in the same trophic guild, whereas the aggregated pairs involved species in the same trophic guild in 56% of the cases and species from different trophic guild in 44% of the cases. *Linepithema neotropicum* and *Pheidole subarmata* was the only species pair exhibiting different patterns of association: negatively associated in the rainy and positively associated in the dry season. Overall, 50% of the species pairs that showed nonrandomness in the occurrences were interpreted as resulting from environmental filters, 36% as negative associations and 14% as positive associations (Tables 4 and 5).

Discussion

The ant assemblages we analyzed were composed by species belonging to the genera that are commonly reported for ground

samples in neotropical forests (McGlynn and Kirksey, 2000; Ward, 2000; Baccaro et al., 2012). Given that our study area was in the process of recovery from disturbance by active restoration with native trees (Azevedo, 2012) and because we have restricted the analyzes to the most frequent ant species, we worked with habitat generalists which are very common in disturbed areas elsewhere. For example, *Pheidole subarmata*, the most frequent species in our studied area, can be found in areas of pastures and crops (Perfecto and Vandermeer, 2011; Munhae et al., 2014). *Linepithema neotropicum*, *Solenopsis invicta* and *Wasmannia auropunctata* are recorded in different types of agroecosystems or urban areas (Munhae et al., 2014; Sinisterra et al. 2016; Santos et al., 2019). *Hypoponera parva* can be found in both mature and disturbed forests (California Academy of Sciences, 2020). The *Strumigenys* species are also found in disturbed ecosystems (Mentone et al., 2011; Souza-Campana et al., 2016). Moreover, several species, although nesting and foraging on the ground, are also known to forage on trees, such as *L. neotropicum*, *W. auropunctata*, and species of *Solenopsis* (e.g. Schoederer et al., 2010; Sinisterra et al., 2016; Qin et al., 2019). Therefore, we believe that the results found here may apply to other generalist ant assemblages in forests with some degree of disturbance, such as naturally recovered secondary forests.

Our analyses showed that species pair co-occurrence patterns for ground-dwelling ants are influenced by environmental filter, even on a small spatial scale, although other studies have suggested that environmental filter operate more frequently at larger spatial scales or between different types of ecosystems (Gotelli and Ellison, 2002; King, 2007; Fichaux et al., 2019). When we ran the analyzes using the predictions of the ensemble models to create restricted null models (Peres-Neto et al., 2001), 50% of the pairs with nonrandom distribution

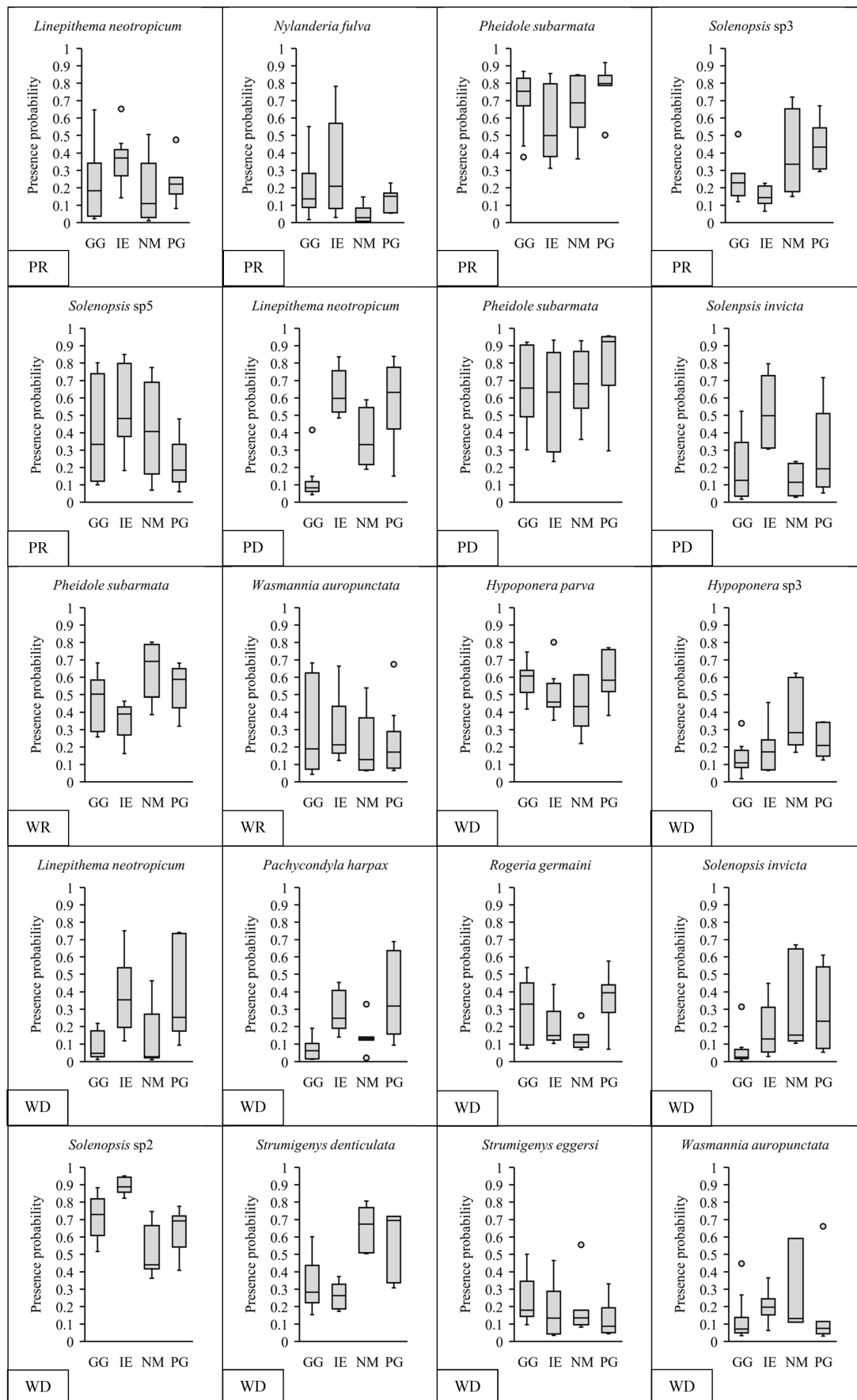


Figure 2 Presence probabilities of ant species in each tree species only for those one involved in nonrandom pairs for the two sampling techniques and seasons. PR: pitfall, rainy season; PD: pitfall, dry season; WR: Winkler, rainy season; WD: Winkler, dry season. GG= *Guarea guidonia*; IE= *Inga edulis*; NM= *Nectandra membranacea*; PG= *Piptadenia gonoachanta*.

Table 4

Species pair co-occurrence patterns of ants sampled with pitfall traps and the interpreted process behind the observed patterns. Food habits indicated in parentheses; om=omnivore, pr=predator.

Sp1	Sp2	Season	Model	P	Pattern	Interpretation
<i>Pheidole subarmata</i> (om)	<i>Linepithema neotropicum</i> (om)	rainy	unc	0.04	Segregated	Negative association
			cons	0.05	Segregated	
<i>Solenopsis</i> sp3 (om)	<i>Solenopsis</i> sp5 (om)	rainy	unc	<0.01	Segregated	Environmental filter
			cons	>0.05	Random	
<i>Solenopsis</i> sp5 (om)	<i>Nylanderia fulva</i> (om)	rainy	unc	0.05	Aggregated	Environmental filter
			cons	>0.05	Random	
<i>Pheidole subarmata</i> (om)	<i>Linepithema neotropicum</i> (om)	dry	unc	0.02	Aggregated	Positive association
			cons	0.04	Aggregated	
<i>Solenopsis invicta</i> (om)	<i>Linepithema neotropicum</i> (om)	dry	unc	0.02	Aggregated	Environmental filter
			cons	>0.05	Random	

Table 5

Species pair co-occurrence patterns of ants sampled with Winkler method and the interpreted process behind the observed patterns. Food habits indicated in parentheses; om=omnivore, pr=predator.

Sp1	Sp2	Season	Model	P	Pattern	Interpretation
<i>Pheidole subarmata</i> (om)	<i>Wasmannia auropunctata</i> (om)	rainy	unc	0.04	Segregated	Environmental filter
			cons	>0.05	Random	
<i>Pachycondyla harpax</i> (pr)	<i>Solenopsis invicta</i> (om)	dry	unc	0.04	Aggregated	Environmental filter
			cons	>0.05	Random	
<i>Pachycondyla harpax</i> (pr)	<i>Linepithema neotropicum</i> (om)	dry	unc	0.01	Aggregated	Environmental filter
			cons	>0.05	Random	
<i>Hypoponera</i> sp3 (pr)	<i>Hypoponera parva</i> (pr)	dry	unc	0.01	Segregated	Negative association
			cons	0.01	Segregated	
<i>Hypoponera</i> sp3 (pr)	<i>Wasmannia auropunctata</i> (om)	dry	unc	0.02	Aggregated	Environmental filter
			cons	>0.05	Random	
<i>Hypoponera parva</i> (pr)	<i>Strumigenys eggersi</i> (pr)	dry	unc	0.01	Segregated	Negative association
			cons	0.03	Segregated	
<i>Strumigenys denticulata</i> (pr)	<i>Strumigenys eggersi</i> (pr)	dry	unc	0.01	Segregated	Negative association
			cons	0.04	Segregated	
<i>Strumigenys eggersi</i> (pr)	<i>Solenopsis</i> sp2 (om)	dry	unc	0.04	Aggregated	Positive association
			cons	0.04	Aggregated	
<i>Rogeria germani</i> (om)	<i>Solenopsis</i> sp2 (om)	dry	unc	0.03	Segregated	Negative association
			cons	0.03	Segregated	

became non-significant, being interpreted as environmental filter effects (D'Amen et al., 2018). Therefore, using the habitat restricted null models, we were able to confirm our hypothesis that the tree attributes act as environmental filters for ant species occurrences. We also demonstrated that the biotic interactions were as important as environmental filters in the observed nonrandom distribution of a small number of species pairs. However, despite the small spatial scale and the relatively homogeneous area where the work was done, our interpretation of the processes causing the co-occurrence pattern is dependent on our premise related to species dispersion. We have to consider that there is little information on the dispersion capacity of ant species (Hakala et al., 2019) and some of them may not fit the general pattern. Moreover, although model performances for most species were quite acceptable, there is room for substantial model improvement for a few species. We found that the effects of environmental filters generate more convergence than divergence on the occurrence of ant

species. The results suggested that the selected tree species are able to cause patchiness in the distribution of a small number of generalist ants in the area, favoring the idea that trees can be used as templates for the study of ground-dwelling ants. For example, *L. neotropicum*, *Pachycondyla harpax*, and *S. invicta* showed an aggregated distribution with higher presence probabilities under the canopies of *I. edulis* and *P. gonoacantha*. Although the tree species have some differences in attributes (see Table 1), both have extrafloral nectaries (EFN) which can be tended by ants. Indeed, *L. neotropicum* is known to visit EFN of *I. edulis* in agroecosystems (Sinisterra et al., 2016) and perhaps the same can occur with *S. invicta*. Ant species that are frequent users of EFN are known to nest closer to EFN-producing plants (Silva et al., 2019). *Pachycondyla harpax* usually nests underground in the vicinity of trees (Grüter et al., 2018), and although workers forage primarily on the ground, they can also be found on trees (Rocha et al., 2015). However, if *P. harpax* attends EFNs on *I. edulis* and *P. gonoacantha* trees needs to

be further investigated. *Nylanderia fulva* and *Solenopsis* sp5 is another case of aggregated distribution associated with *Inga edulis*, possibly involving the frequent visit to EFN (Wang et al., 2016). Among the few pairs that were segregated in distribution by environmental filters, the pair of *Solenopsis* species can also be explained by the divergence in the probabilities of occurrence under the canopies of the different trees. *Hypoponera* sp3 and *W. auropunctata* had aggregated distribution because both had the occurrence more associated with trees where the litter was deeper, such as *Nectandra membranacea*. However, the segregation by environmental filters between *P. subarmata* and *W. auropunctata* cannot be linked to tree identity, because there was no clear trend on the probability of species presence among the tree species. Based on our results we can predict that other pairs of ant species occurring in the area, but not evaluated here, will also show nonrandom distributions caused by either aggregation or segregation in relation to tree species.

We found more negative than positive associations between species, thus differing from the pairwise analysis of Ellwood et al. (2016) with tropical ant assemblages. In relation to the species pairs negatively associated, but not by environmental filters, we had two pairs in the same genus and three pairs in the same guild (omnivores). Therefore, our results agree with Camarota et al. (2016), who also demonstrated that species pairs with segregated distribution had higher trait similarity. The first case of negative association was between *Strumigenys eggersi* and *S. denticulata*. *Strumigenys* species are very common in litter samples, and although, on some occasions, we have collected more than one *Strumigenys* species per sample, we never found *S. eggersi* and *S. denticulata* together. *Strumigenys denticulata* is the most frequent species in litter samples in the Atlantic Forest and has a morphology very similar to that of *S. eggersi* (Silva, 2014). We do not know of studies on interactions between species of this genus, but this may be because they are small in size and cryptic (Lattke et al., 2018). Species of *Strumigenys* nest in the litter and are recognized by their specialized habit of preying on Collembola (Silva, 2014; Lattke et al., 2018). The second case is between *Hypoponera parva* and *Hypoponera* sp3; *Hypoponera parva* was one of the most frequent species in the studied site, whereas *Hypoponera* sp3 was much less frequent. The case of *Hypoponera* is similar to *Strumigenys* in that the species are cryptic, nesting in the litter or soil, and although they are considered more generalist than *Strumigenys*, they can also be predators of collembolans (Baccaro et al., 2015). Although it is not uncommon to find *Hypoponera* species at the same spot (Soares and Schoederer, 2001; Brandão et al., 2012), we never found *H. parva* and *H. sp3* together in our samples.

The segregation we observed between species from different genera, involved pairs in the same feeding guild. *Hypoponera parva* and *S. eggersi* were negatively associated and, because they both share feeding and nesting habits, it is possible to suggest competition as the cause of the segregated distribution. Our results suggested that *Rogeria germaini* could be displaced by *Solenopsis* sp2, because the latter was much more frequent than the former. There is little information about *Rogeria* species besides they are cryptic omnivores, nesting in soil or litter (Baccaro et al., 2015; Koch et al., 2019). However, *Solenopsis* species are usually involved in competition with different ant groups (Hölldobler and Wilson, 1990; LeBrun et al., 2007; Perfecto and Vandermeer, 2011), since their colonies are bigger, being generalist in feeding and nesting habits, and dominant over resources (Baccaro et al., 2012, 2015).

The last case of segregation is somewhat dubious, and perhaps more interesting. *Pheidole subarmata* and *L. neotropicum* showed segregation in the rainy season and aggregation in the dry season for the pitfall sampling. *Linepithema neotropicum* was much less frequent in the rainy season, agreeing with the results of Munhae et al.

(2014), whereas *P. subarmata* was highly frequent in both seasons. It is possible to assume that the rainy season negatively affects the foraging of *L. neotropicum* but not *P. subarmata*, perhaps because the latter forages predominantly on the ground. The negative effect of rainfall should be greater on vegetation than on the ground, since vegetation intercepts raindrops in the forest (Park and Cameron, 2008). Indeed, Hahn and Wheeler (2002) found that the effect of the rainy season on ant activity was higher for ants on plants than on the ground. If the foraging of *L. neotropicum* became more restricted to the ground during the rainy season, it is possible that it suffered more from competition with *P. subarmata*, thus explaining the segregated distribution during this season. Conceição et al. (2014) observed that *L. neotropicum* is behaviourally dominant at baits. On the other side, Perfecto and Vandermeer (2011) observed that *P. subarmata* discover fastly food resources on the ground, when nest density is high, but it is not dominant in interactions with *S. geminata*. Therefore, the rainy season may favor *P. subarmata* in the dispute with *L. neotropicum* for resources on the ground. During the dry season, the foraging of *L. neotropicum* workers is increased, and without the negative effects of raindrops on vegetation, the foraging on trees might be higher. Hence, in the dry season, the competitive interactions between *P. subarmata* and *L. neotropicum* on the ground would be reduced. We found only one other species pair with positive association between *Strumigenys eggersi* and *Solenopsis* Sp2. Positive associations between species can be generated by consumer-resource and mutualistic interactions, but Ellwood et al. (2016) suggested that even competition between ant species may result in positive associations between species. Therefore, we need more information on species biology to uncover the mechanisms behind these positive associations.

Similar to our results, other studies, done in ecosystems with different levels of precipitation, have observed a negative effect of the rainy season on the abundance and richness of ants (e.g. Basu, 1997; Anu et al., 2009; Neves et al., 2010; Jacquemin et al., 2016). Different reasons were suggested to explain the seasonal pattern in ant assemblages, including the direct effect of rain (Anu et al., 2009) and the change in food supply (Neves et al., 2010; Jacquemin et al., 2016). Our results suggest that the direct effects of rain can be more important for the species that forage on the surface, e.g. *L. neotropicum*, and that are most captured in pitfall traps (Bestelmeyer et al., 2000). The drop in the number of records for pitfall traps was much greater than that observed in Winkler samples during the rainy season. Perhaps the rainy season affects less those species that forage in the interstices of the litter, and are sampled with the Winkler method, e.g. *Strumigenys* species (Parr and Chown, 2001; Farji-Brener et al., 2004). Seventy-one and sixty-four percent of the pairs with nonrandom distribution was observed, respectively, in the dry season and in the Winkler samples. The effects of both the season and the sampling technique on the co-occurrence patterns of ground-dwelling ants suggest that they should not be ignored in this kind of study.

Our work showed that most pairs of ant species were randomly distributed. The random distribution of ground-dwelling ant species also prevailed in studies done in the Amazon rainforest (Baccaro et al., 2012), the Atlantic forest (Hanisch et al., 2018), and Asian tropical forests (Fayle et al., 2013). Even for canopy ants the random pattern can prevail (e.g. Camarota et al., 2016). The predominance of random distribution of species pairs in ant assemblages may suggest that stochastic processes would be more important than deterministic processes (Ribas and Schoederer, 2002) or that we do not know the nonrandom causes behind the observed patterns (Vellend, 2017). For the species pairs showing nonrandom distribution, we suggest that the ideas of the trees as templates (Donoso et al., 2010, 2013) and the paradigm of competition (Cerdá et al., 2013) are both useful for understanding pairwise occurrence

patterns in ant assemblages. Advances in myrmecology, especially in Brazil, are providing increasingly taxonomic and ecological knowledge of neotropical ant species (e.g. Baccaro et al., 2015). This will allow a substantial improvement in species distribution models that can be used in the approach reported here to uncover the deterministic causes of species' co-occurrence patterns.

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Conflicts of interest

The authors declare no conflicts of interest.

Author contribution statement

Conceived and designed experiments: JMQ; data sampling: RCN, ECBF, JMQ; organism identifications: ECBF, RMF; data analysis: RCN, JMQ; manuscript writing: RCN, JMQ; manuscript review: ECBF, RMF.

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Supplementary material

The following online material is available for this article:

Appendix 1 - Map of georeferenced distribution of ant sampling points in the studied area at REGUA, Cachoeiras de Macacu-RJ, Brazil. GG= *Guarea guidonia*; IE= *Inga edulis*; NM= *Nectandra membranacea*; PG= *Piptadenia gonoachanta*

Appendix 2 - AUC statistics for the SDMs of ant species used in the constrained null models.

Appendix 3 - SES_Cscore and p values for each nonrandom species pair