






# Wood-decay fungi (Agaricomycetes, Basidiomycota) in three physiognomies in the Savannah region in Brazil<sup>1</sup>

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Received: 27 July 2018; accepted: 18 February 2019

**How to cite:** Abrahão, M.C., Pires, R.M., Gugliotta, A.M., Gomes, E.P.C. & Bononi, V.L.R. 2019. Wood-decay fungi (Agaricomycetes, Basidiomycota) in three physiognomies in the savannah region in Brazil. *Hoehnea* 46: e692018. <http://dx.doi.org/10.1590/2236-8906-69/2018>.

**ABSTRACT** - (Wood-decay fungi (Agaricomycetes, Basidiomycota) in three physiognomies in the Savannah region in Brazil). This study was carried out aiming at determining whether the following factors are related to the diversity of wood-decay fungi: cerrado physiognomies, vegetation diversity (availability of substrates), preservation status of the area and its management backgrounds, and the existence of rivers nearby (humidity). 796 specimens were collected in four areas: one in the cerrado *sensu stricto*, one in the semi-deciduous forest and two in the riparian forest (one more preserved and the other less preserved). A total of 92 species of fungi were identified. Eleven species are common in the four areas sampled and 30 species occurred only once during the two-year sampling. Higher richness and abundance were observed in the more preserved area of riparian forest. This study showed that in the same macroclimate, the Agaricomycetes diversity is influenced by the vegetation's physiognomy and diversity related to the soil characteristics, humidity, and conservation level of the area. **Keywords:** Cerrado, fungal community, ligninolytic fungi, riparian forest, semi-deciduous forest

**RESUMO** - (Fungos decompositores de madeira (Agaricomycetes, Basidiomycota) em três fisionomias no Cerrado do Brasil). Este estudo foi realizado visando Cerrado do Brasil). Visando entender se há correlações entre fisionomia e diversidade da vegetação (disponibilidade de substratos), grau de preservação, história de manejo da área e a proximidade de cursos de água com a diversidade de fungos decompositores de madeira. 796 espécimes foram coletados em quatro áreas: uma no cerrado *sensu stricto*, uma na floresta estacional semidecidual e duas de floresta ripária (mais e menos preservada). Do total de 92 espécies de fungos, onze são comuns a todas as áreas amostradas e 30 ocorreram uma única vez. Maiores riqueza e abundância de Agaricomycetes foram observadas na floresta ripária mais preservada. Este estudo mostrou que, sob o mesmo macroclima, a diversidade de Agaricomycetes é influenciada por fatores como fisionomia e diversidade da vegetação determinadas pelo tipo de solo, umidade e grau de preservação da área.

**Palavras-chave:** Cerrado, floresta estacional semidecidual, fungos lignolíticos, mata ciliar, micota

## Introduction

Wood-inhabiting Agaricomycetes has an enzymatic system with extracellular activity to degrade complex structural materials, such as lignin, wood cellulose and hemicellulose, which makes them a group of great ecological value for recycling nutrients in ecosystems (Gilbertson 1980, Alexopoulos

*et al.* 1996, Lundell *et al.* 2010). The total diversity of fungi is estimated to range from 0.7 to 5.1 million species (Blackwell 2011, Hawksworth & Lücking 2017), but so far only about 1.9% (97,330 species) have been described. A total of 538 genera and 6,384 species of wood-decay Agaricomycetes are known worldwide (Kirk *et al.* 2008).

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The Cerrado is the second largest area in Brazil (21% of the Brazilian territory, around 1.8 to 2.0 million km<sup>2</sup>), after the Amazon Forest (IBGE 2004, Coutinho 2016). Studies regarding Agaricomycetes focused on taxonomy mention the occurrence of 134 species (Sampaio 1916, Fidalgo *et al.*, 1965, Bononi 1984, Gugliotta 1997, Baseia & Milanez 2001a, b, 2002a, b, 2003, Baseia 2005, Baseia *et al.* 2007, Abrahão *et al.* 2012, Alvarenga *et al.* 2015, Costa-Rezende *et al.* 2015). None of these authors present ecological relations between fungi diversity and the “cerrado” physiognomies *sensu stricto*.

The general term “Cerrado” has been used to designate savannah-type vegetation with a gradient of physiognomies, from the grassland (clean field) to dense forest with 15 to 18 m high vegetation (“cerradão”), and between those, there are some intermediate physiognomies. Forests or woods in the “Cerrado” Biome near watercourses are riparian forests which correspond to about 5% of the Cerrado area (Pivello & Coutinho 1996). But within the “Cerrado” Biome, patches of other vegetation physiognomies may occur interspersed among the cerrado forms, determined mainly by edaphic and geomorphological features, and semi-deciduous forests, which can be found in richer soils, with more water available and they correspond to about 15% of the “Cerrado” area. Gallery forests present more diversity than those in the Cerrado areas. There are Cerrado-preferred species, and exclusively semi-deciduous forests species and there are common species in all ecosystems (Eiten 1994, Pereira *et al.* 2011).

Satellite data showed that 55% of the Cerrado had been cleared or modified, mainly for setting up pastures and agricultural fields (Klink & Machado 2005).

Considering the importance of wood-decay Agaricomycetes in the maintenance of ecosystems and the shortage of studies (Seibold *et al.* 2015), this study aimed at understanding whether there is any correlation between the diversity of wood-decay fungi and the following factors: “cerrado” physiognomies, vegetation diversity (availability of substrates), area preservation status and its management background and existence of rivers nearby (humidity).

## Material and methods

**Study area** - The Mogi-Guaçu Biological Reserve (22°11’-22°16’S, 47°08’-47°11’W) is composed by two areas (A and B), located in the municipality

of Mogi-Guaçu, São Paulo State (figure 1). In this region, the Cerrado vegetation almost reaches its austral limit (Klink & Machado 2005). Altitudes vary from 590 to 650 m; the predominant climate is Aw with rainy summers and dry winters according to the classification of Köppen-Geiger (Peel *et al.* 2007). The soil is classified as sandy, acidic, poor in organic matter and oligotrophic (Giudice-Neto 2010). Soil abiotic factors in areas A and B during the rainy and dry seasons are presented by (Nascimento *et al.* 2012).

The areas studied (areas A and B) cover a total area of 470 ha and they have not been disturbed by fire for the last 30 years. The surroundings of both areas are mostly agricultural production units, showing diversified cultures of eucalyptus, cassava, sugar cane, corn, and oranges. The riparian forest is associated with the Mogi-Guaçu River and streams. A description of the particularities of the areas can

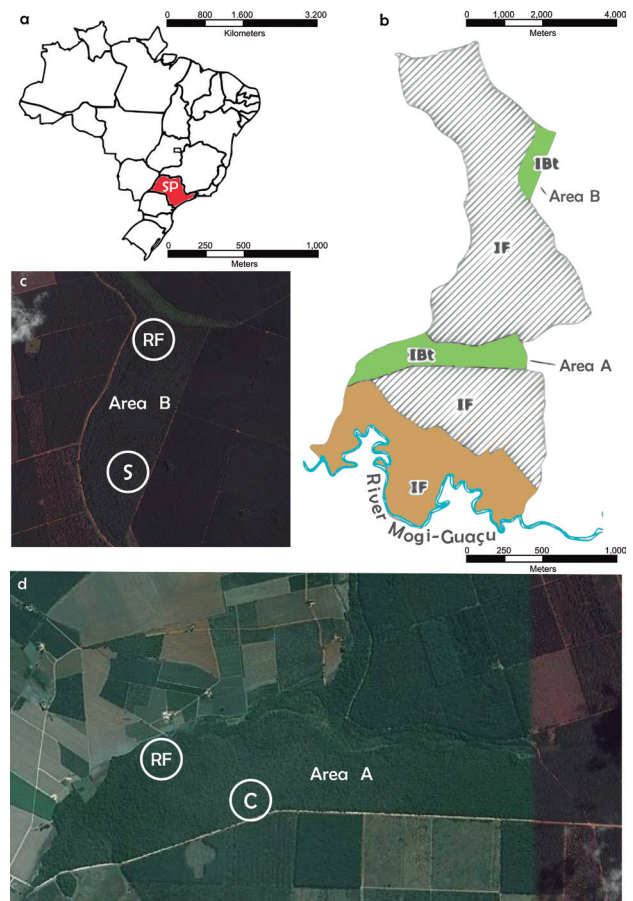


Figure 1. a. Location of the studied areas in the Mogi-Guaçu Biological Reserve (IBt), São Paulo State, Brazil. b. Mogi-Guaçu Biological Reserve (IBt: Conservation Area of the Instituto de Botânica, IF: Conservation Area of the Instituto Florestal). c. Satellite image of Area B (C: Cerrado; S: Semi-deciduous Forest; RF: Riparian Forest). d. Satellite image of Area A. (Modified from Giudice-Neto, 2010).

be found in Giudice-Neto (2010). Four areas were sampled: one in the cerrado *sensu stricto*, one in the semi-deciduous forest and two in the riparian forest (one more preserved and the other less preserved).

Area A, covering 343.42 ha, is composed of “Cerrado” vegetation at a medium stage of regeneration. According to Mantovani & Martins (1993), this area has been used for cattle grazing and logging.

Area B, covers 126.63 ha, with a semi-deciduous forest. The original forest is in good conditions (no exotic species, large gaps or herbaceous climbers). However, in some areas (mainly in the center), there are trees of around 15 m high and the herbaceous layer shows a high grass density.

Fungi sampling - Considering the recommendations of Zak & Willig (2004) and a preliminary study conducted by Abrahão *et al.* (2010), sample sets were placed randomly at 10 transects of  $4 \times 50$  m in the riparian forest in area A (A\_RF) and area B (B\_RF), in the Cerrado vegetation in area A (A\_C) and semi-deciduous forest in area B (B\_S). The entries of the transects were georeferenced. Basidiomata were collected bimonthly at those transects from August/2009 to June/2011, totaling 12 field expeditions covering all seasons. Agaricales was excluded from this study due to their ephemeral basidiomata, which requires a different methodology.

Species identification was based on observation and analysis of basidioma macro- and microscopic characteristics (Ryvarden 1991, Teixeira 1993, 1994, 1995). Evidence for dextrinoid and amyloid reactions was tested in Melzer’s reagent (Gilbertson & Ryvarden 1986, Gugliotta & Bononi 1999). Preservation and herborization of specimens were performed according to Fidalgo & Bononi (1984). Vouchers for every species were deposited at the SP Herbarium of the Instituto de Botânica (SP).

The species recorded over 15 times in all areas during the study were considered common species, while species recorded less than five times were considered rare and a species recorded between 5 and 15 times, an occasional species (Dai 2012).

Statistical analysis - To compare species richness between the sample sets (A\_C, B\_S, A\_RF, B\_RF), an estimate was made by rarefaction with a confidence interval of 95% to account for differences in the sampling effort.

Diversity was assessed by profile diversity, which allows simultaneous evaluation of richness

and abundance of every species with different weights in different diversity indices. The profile is derived from the Renyi’s general formula, the Renyi index (Tóthmérész 1995). When  $\alpha = 0$ , this function provides the total number of the species richness. For  $\alpha = 1$ , the index is proportional to the Shannon index, whereas, for  $\alpha = 2$ , the index behaves like the Simpson index assigning higher weight and relative abundance of species. The profile is represented in an alpha vs. values graph to which a curve is added, allowing checking the indices simultaneously.

The sampling units were organized according to the principal coordinate’s analysis (PCoA) to evaluate their relations in the ordering space from the distance matrix among the sampling units, using the Jaccard similarity index. Due to the sensitivity of the method to matrices with many zeros, species with a single record were excluded from the analysis.

For comparisons among groups for all transects, plant communities and area were used as grouping variables; the similarity analysis permutation test was applied (ANOSIM, Clarke & Warwick 1994) using the Bray-Curtis dissimilarity matrix. Analyses were performed with PAST software version 2.17c with 9,999 permutations for ANOSIM (Hammer *et al.* 2001).

## Results and Discussion

A total of 796 specimens of 92 species were collected throughout the study.

Higher richness and abundance were observed at area B (71 species, 503 specimens), including semi-deciduous forest and riparian forest. In the area A 54 species, 293 specimens were found, including Cerrado and riparian forest. Many species (38; 41.3%) were exclusive to area B and 21 species (38.89%) were exclusive to area A and 18 species were exclusive to the Cerrado and 31, to the riparian forest (table 1).

Considering all areas, the group sample B\_RF showed 272 specimens (34.17%), followed by B\_S (231 specimens, 29.02%), A\_C (155 specimens, 19.47%) and A\_RF (138 specimens, 17.34%) (table 1). Eleven species are common in the four areas sampled and 30 species occurred only once during the two-year sampling.

Sampling groups showed a significant difference in composition, except for A\_C and A\_RF (table 2, ANOSIM, “mean rank within” 271.6; “mean rank between” 426.2;  $p < 0.05$ ).

In the richness-based ordination analysis (figure 2), the two axes explained 23.75% of the

Table 1. Wood-decay Agaricomycetes collected in areas A and B in the Mogi-Guaçu Biological Reserve, SP. A\_C: relative abundance in Cerrado transections of area A, A\_RF: relative abundance on transections of Riparian Forest in area A, A: abundance in area A, B\_S: relative abundance of transections in Semideciduous Forest of area B, B\_RF: relative abundance of transections in the Riparian Forest of area B, B: abundance in area B, T: total abundance.

Species	A_C	A_RF	A	B_S	B_RF	B	T
<i>Fuscoporia gilva</i>	11	8	19	39	21	60	79
<i>Pycnoporus sanguineus</i>	26	27	53	2	11	13	66
<i>Tyromyces duracinus</i>	2	-	2	37	15	52	54
<i>Fomitiporia maxonii</i>	20	6	26	16	4	20	46
<i>Polyporus guianensis</i>	-	-	0	29	12	41	41
<i>Trametes villosa</i>	6	10	16	4	20	24	40
<i>Coriolopsis floccosa</i>	7	12	19	12	7	19	38
<i>Hymenochaete tenuis</i>	8	7	15	10	8	18	33
<i>Schizophyllum commune</i>	7	10	17	-	8	8	25
<i>Dichomitus setulosus</i>	5	7	12	7	1	8	20
<i>Perenniporia ochroleuca</i>	7	3	10	5	1	6	16
<i>Chondrostereum purpureum</i>	5	2	7	4	4	8	15
<i>Lentinus crinitus</i>	6	4	10	1	4	5	15
<i>Hymenochaete digitata</i>	3	4	7	1	5	6	13
<i>Polyporus tenuiculus</i>	-	-	0	6	7	13	13
<i>Coriolopsis caperata</i>	-	-	0	1	11	12	12
<i>Echinoporia aculeifera</i>	-	2	2	1	9	10	12
<i>Schizopora paradoxa</i>	3	-	3	4	5	9	12
<i>Stereum ostrea</i>	1	-	1	3	7	10	11
<i>Gloeoporus thelephoroides</i>	-	-	0	6	4	10	10
<i>Pachykytospora alabamiae</i>	2	6	8	1	1	2	10
<i>Polyporus leprieurii</i>	-	-	0	4	6	10	10
<i>Schizophyllum umbrinum</i>	-	4	4	-	5	5	9
<i>Aleurodiscus cerussatus</i>	3	2	5	3	-	3	8
<i>Grammothele subargentea</i>	2	3	5	2	1	3	8
<i>Hexagonia variegata</i>	-	-	0	6	2	8	8
<i>Hydnopolyporus fimbriatus</i>	-	-	0	-	8	8	8
<i>Hymenochaete damicornis</i>	3	-	3	-	5	5	8
<i>Podoscypha ravenelii</i>	-	-	0	-	8	8	8
<i>Scopuloides rimosa</i>	2	-	2	-	6	6	8
<i>Phylloporia chrysites</i>	-	-	0	-	7	7	7
<i>Schizopora flavipora</i>	2	-	2	4	1	5	7
<i>Hymenochaete rheicolor</i>	1	3	4	2	-	2	6
<i>Mycobonia flava</i>	-	-	0	1	5	6	6
<i>Polyporus dictyopus</i>	-	-	0	1	5	6	6
<i>Dichomitus cavernulosus</i>	2	-	2	2	1	3	5
<i>Pachykytospora papyracea</i>	-	1	1	2	2	4	5
<i>Datronia mollis</i>	-	-	0	-	4	4	4
<i>Irpex lacteus</i>	-	2	2	-	2	2	4
<i>Polyporus ciliatus</i>	-	-	0	-	4	4	4

continue

Table 1 (continuation)

Species	A_C	A_RF	A	B_S	B_RF	B	T
<i>Tyromyces leucomallus</i>	-	-	0	1	3	4	4
<i>Auricularia polytricha</i>	1	1	2	-	1	1	3
<i>Cyclomyces iodinus</i>	-	-	0	3	-	3	3
<i>Daedalea aethalodes</i>	3	-	3	-	-	0	3
<i>Fomes fasciatus</i>	-	-	0	-	3	3	3
<i>Gloeoporus dichrous</i>	-	-	0	3	-	3	3
<i>Hymenochaete opaca</i>	-	3	3	-	-	0	3
<i>Hyphoderma</i> sp. 1	1	2	3	-	-	0	3
<i>Phanerochaete</i> sp. 1	1	-	1	-	2	2	3
<i>Rigidoporus microporus</i>	-	-	0	-	3	3	3
<i>Steccherinum</i> sp. 1	1	-	1	-	2	2	3
<i>Aleurodiscus botryosus</i>	1	1	2	-	-	0	2
<i>Cymatoderma caperatum</i>	-	-	0	-	2	2	2
<i>Gloeoporus purpurascens</i>	1	-	1	-	1	1	2
<i>Hidnoide</i> sp. 1	1	1	2	-	-	0	2
<i>Hymenochaete corrugata</i>	-	-	0	-	2	2	2
<i>Hyphodontia</i> sp. 1	1	-	1	1	-	1	2
<i>Hyphodontia</i> sp. 2	1	-	1	1	-	1	2
<i>Phanerochaete</i> aff. <i>sordida</i>	-	-	0	-	2	2	2
<i>Polyporus tricholoma</i>	-	-	0	-	2	2	2
<i>Steccherinum reniforme</i>	-	-	0	1	1	2	2
<i>Trichaptum sector</i>	-	-	0	-	2	2	2
<i>Aleurodiscus exasperatus</i>	1	-	1	-	-	0	1
<i>Amauroderma praetervisum</i>	-	-	0	-	1	1	1
<i>Ceriporia spissa</i>	-	1	1	-	-	0	1
cf. <i>Brevicellicium</i> spp.	1	-	1	-	-	0	1
<i>Dichostereum</i> cf. <i>sordulentum</i>	-	1	1	-	-	0	1
<i>Hexagonia hydnoides</i>	-	-	0	1	-	1	1
<i>Hidnoide</i> sp. 2	-	-	0	-	1	1	1
<i>Hidnoide</i> sp. 3	-	1	1	-	-	0	1
<i>Hidnoide</i> sp. 4	-	-	0	-	1	1	1
<i>Hidnoide</i> sp. 5	-	-	0	-	1	1	1
<i>Hymenochaete pinnatifida</i>	1	-	1	-	-	0	1
<i>Hyphodontia</i> sp. 3	1	-	1	-	-	0	1
<i>Hyphodontia</i> sp. 4	-	-	0	1	-	1	1
<i>Hyphodontia</i> sp. 5	-	1	1	-	-	0	1
<i>Hyphodontia</i> sp. 6	-	-	0	-	1	1	1
<i>Merulius</i> sp. 1	-	-	0	1	-	1	1
<i>Phanerochaete corymbata</i>	-	1	1	-	-	0	1
<i>Phanerochaete</i> sp. 2	1	-	1	-	-	0	1
<i>Phanerochaete</i> sp. 3	1	-	1	-	-	0	1

continue

Table 1 (continuation)

Species	A_C	A_RF	A	B_S	B_RF	B	T
<i>Phanerochaete</i> sp. 4	1	-	1	-	-	0	1
<i>Phanerochaete</i> sp. 5	1	-	1	-	-	0	1
<i>Phellinus</i> cf. <i>chryseus</i>	-	1	1	-	-	0	1
<i>Phellinus punctatiformis</i>	-	-	0	-	1	1	1
<i>Phlebia</i> sp. 1	1	-	1	-	-	0	1
<i>Phylloporia spathulata</i>	-	-	0	-	1	1	1
<i>Rigidoporus lineatus</i>	-	-	0	-	1	1	1
<i>Scytinostroma duriusculum</i>	-	-	0	1	-	1	1
<i>Steccherinum</i> sp. 2	-	1	1	-	-	0	1
<i>Stereum</i> sp. 1	-	-	0	-	1	1	1
<i>Stereum</i> sp. 2	-	-	0	1	-	1	1
Total of occurrences	155	138	293	231	272	503	796
Total of species	42	32	54	41	60	71	92

Table 2. Probability of lack of differences from dissimilarity matrix of Bray-Curtis among sampling groups based on species abundance found in Mogi-Guaçu Biological Reserve, SP. Non-parametric similarity analysis of one factor (ANOSIM; 9999 permutations; R: 0.3962; Bonferroni correction). A\_C: Cerrado of area A, A\_RF: Riparian Forest in area A, B\_S: Semideciduous Forest of area B, B\_RF: Riparian Forest of area B.

	A_C	A_RF	B_RF	B_S
A_C		1,0000	0,0018	0,0006
A_RF			0,0048	0,0006
B_RF				0,0024
B_S				

variation. Axis 1 separated the areas A and B. The results of ANOSIM showed significant differences between areas (table 2, ANOSIM, “mean rank within” 271.6, “mean rank between” 426.2;  $p < 0.05$ ), except for the two plant communities in area A.

The rarefaction curve of the species (figure 3) and the diversity profile diagram (figure 4) showed that the general standard of richness is higher in B\_RF, followed by A\_C, B\_S, and A\_RF.

The results of the similarity analysis and ordination showed differences between areas A and B as well as between different physiognomies. Areas A and B, despite being close to each other and subject to the same macroclimate conditions, differ in several factors such as soil type (Nascimento *et al.* 2012), vegetation type and factors related to microclimate (Giudice-Neto 2010).

Additionally, areas A and B, although they have not been disturbed by fire for the last 30 years, differ

as to their conservation level because they have a different handling background, area B being more conserved than area A, which was more used in the past for cultivation, wood removal and cattle grazing, and later for research in general (Mantovani & Martins 1993, Giudice-Neto 2010). So, the riparian forest in area B is more preserved than that in area A, which can explain the more significant diversity of Agaricomycetes in the riparian forest of area B, and little similarity among the riparian areas. In area A, higher values for species richness were recorded for the Cerrado sampling group, probably due to the degradation of the riparian forest.

Lindner *et al.* (2006) studied the effects of forest management on fungal diversity and found significant differences for the most abundant species, indicating that the history of the site, amount and quality of the substrate and the management of the area are important factors to predict fungi species richness.

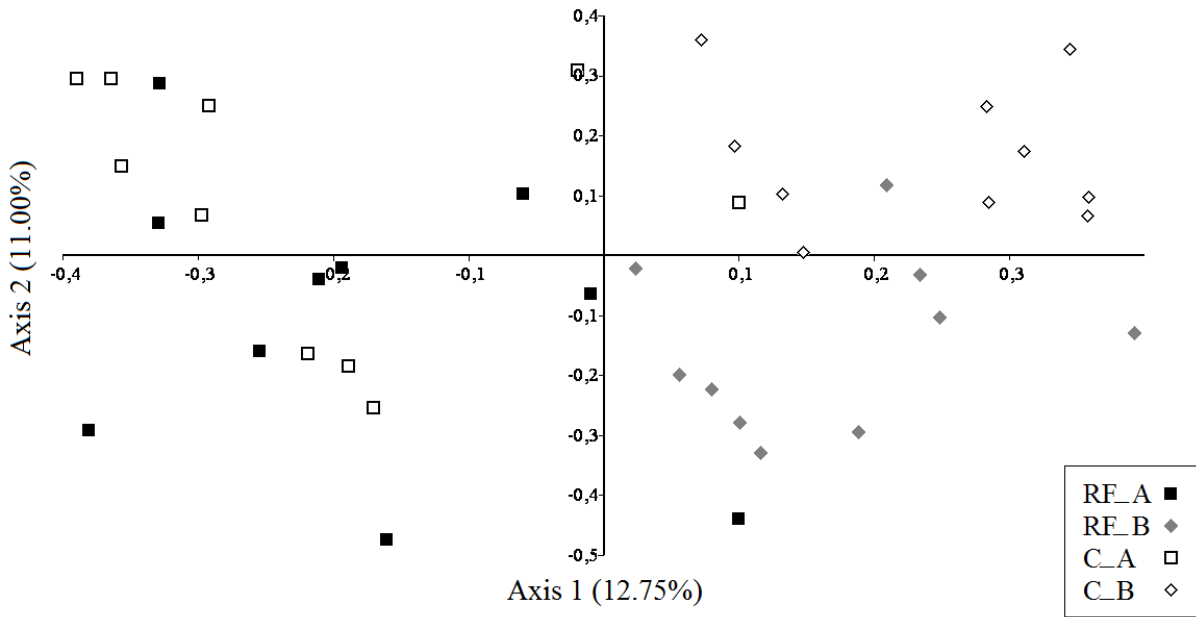


Figure 2. Ordination of 40 samples (transects  $4 \times 50$  m) in Mogi Guaçu Park, SP, Brazil. The scatter diagram is defined by ordination axes 1 and 2 generated by principal coordinates analysis from Jaccard Similarity Matrix. The percentage of total variation represented by the axes is indicated.

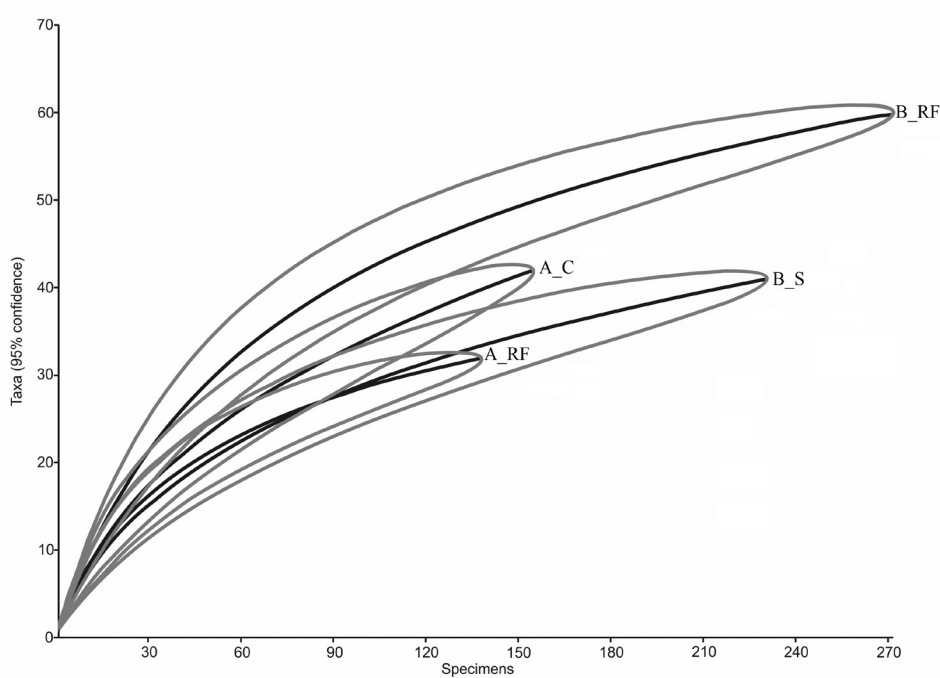


Figure 3. Rarefaction curve of sampling groups observed in the Mogi-Guaçu Biological Reserve, SP. Average of species number (+ confidence interval of 95%) for each area in relation to the function of the number of specimens sampled.

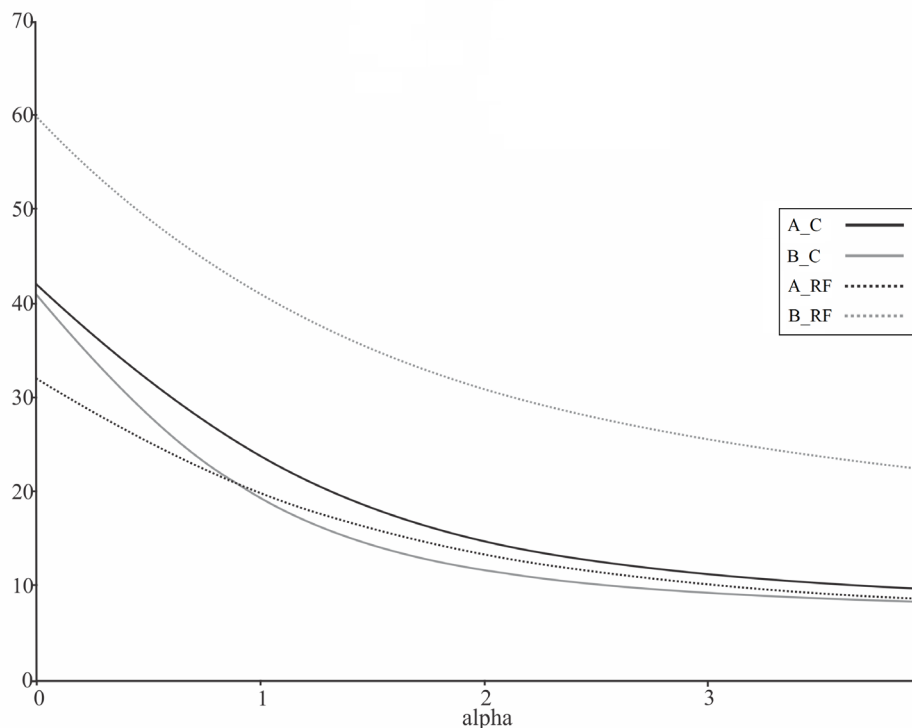


Figure 4. Diversity profile of wood-decay Agaricomycetes community in the Mogi-Guaçu Biological Reserve.

There is only one study in Neotropical region comparing basidiomycete diversity in more and less preserved areas (Gibbertoni *et al.* 2007) conducted in the northeast Atlantic rain forest. They concluded that the number of fungal species is positively affected by the level of conservation of the vegetation. However, the results of Penttilä *et al.* (2006) for polypores in northern Europe showed that species number, the incidence of occurrence and abundance of the rarest, most threatened and near-threatened species are much lower in old-growth fragments with a long history of intensive forestry. In our study, higher species richness in the riparian forest was expected because most fungi species are susceptible to humidity variations. Papers dealing with vegetation (Pereira *et al.* 2011) show that more richness and diversity can be found in riparian forests than in other physiognomies in Cerrado areas. Consequently, substrates' diversity there is higher than in non-riparian areas, which eases wood-degrading fungi's growth.

Using next-generation metagenomic sequencing (NGS) and soil physicochemical properties, Araujo *et al.* (2017) evaluated the fungal diversity across a gradient of vegetation that included Graminoid Field", sensu stricto "Cerrado", "Cerradão", and Deciduous Forest. The results showed that Deciduous Forest

presented the highest richness. Agaricomycetes was one of the most abundant classes of fungi found across the gradient of "Cerrado" area in north-eastern Brazil. The contrast in fungal communities between Deciduous Forest and Graminoid Field evidenced the differences in vegetation and soil properties of these sites. The riparian forest was not considered.

According to Castro *et al.* (2016), who observed a distinction in the fungal communities between Graminoid Field and Riparian Forest from *stricto sensu* "Cerrado" and "Cerradão", compositions of bacterial, archaeal, and fungal communities in the different vegetation physiognomies were associated with temporal variations of soil moisture.

According to Oliveira-Filho & Fontes (2000), there is greater floristic similarity at the species level between Atlantic rain and semi-deciduous forests than between any of these and either Amazonian rain forests or Cerrado. Nevertheless, the Cerrado flora was much more closely related to Atlantic semi-deciduous forests than to the Atlantic rain forests or with Amazonian forests. The results of the similarity analysis and ordination showed that Agaricomycetes diversity of the Cerrado is also much more closely related to semi-deciduous forests.

In Brazil, the only study that evaluated the proportion of common, occasional (or subordinate)



and rare species of Agaricomycetes was conducted by Borba-Silva *et al.* (2015) in the Atlantic Forest in Santa Catarina State. Out of 152 specimens, rare polypore species (less than five records) represented 86% of the total number of species and 50% of total abundance, and only three species were considered common or dominant (5.4% out of all and 26% of total abundance). In our study, most of the species (59.78%) were also considered rare, representing only 12.31% of total abundance, while eleven species were considered common (11.96% out of all species and 57.54% of total abundance). Twenty-six species (28.26%) were considered occasional, representing 30.15% of total abundance. The differences found between our findings and those obtained by those authors are probably the different biomes assessed and the sampling size.

When comparing the most abundant species, *Fuscoporia gilva*, *Tyromyces duracinus*, and *Polyporus guianensis* were more frequent in area B, while *Pycnoporus sanguineus* occurred more often in area A. According to Ryvardeen (2004), *Fuscoporia gilva* is one of the most collected species in tropical regions. According to the Flora Species List of Brazil (Maia *et al.* 2015), *F. gilva* is widespread in Brazil and has been recorded in the Cerrado, the Atlantic Forest, the Amazon Forest, ‘Caatinga’ and ‘Pantanal’.

*Pycnoporus sanguineus* is frequently found in sites exposed to sunlight and with reduced levels of humidity and higher temperatures, being regarded as an indicator of the level of conservation of the areas (Guzmán-Dávalos & Guzmán 1979, Gibertoni *et al.* 2007).

The abundance of some species, such as *Fuscoporia gilva*, *Lentinus crinitus* and *Polyporus guianensis*, can be related to the fact they are species complexes, morphologically quite similar species and, as such, it is difficult to identify them with traditional methods. Those complexes are not yet well known and in the future, they can impact on the results of this paper.

Núñez (1996) studied species showing ecological adaptations to different levels of humidity, some species are perennial and produce basidiomata capable to survive water shortage, forming new tubes and fertile layers instead of new basidiomata after every rainy season, such as some species of the genera *Daedalea*, *Fomes* and *Phellinus s.l.* Some annual species, such as *Coriolopsis* spp., *Fuscoporia gilva*, and annual species of *Hexagonia* develop an annual basidioma that remains sterile under low humidity,

especially on trails, in canopy gaps and dry habitats, but they sporulate under high humidity in closed and humid canopies or during the rainy season in dry forests. On the other hand, species of *Polyporus* and *Tyromyces* are usually dependent on abundant water and are numerous in shaded and humid areas, they produce fleshy basidiomata under suitable conditions, which collapse after sporulating. In our study, *Daedalea aethalodes*, *Fomes fasciatus*, *Fomitiporia maxonii* and *Phellinus punctatiformis* were more abundant in areas of the Cerrado *sensu stricto* (drier), *Fuscoporia gilva* and *Coriolopsis floccosa* occurred in all sampling groups and *Polyporus* (exclusive to area ‘‘B’’) and *Tyromyces* species were much more frequent in area B, where vegetation is more compact and humid.

The riparian forest has higher vegetal diversity than cerrado, it shows Cerrado-preferred species, and common species to riparian, ‘‘cerrado’’ and semi-deciduous forest (Pereira *et al.* 2011). The same was found with the Agaricomycetes diversity studied. Under the same macroclimate, the Agaricomycetes diversity is influenced by factors such as the vegetation physiognomies which is related to the soil type, humidity and preservation level of the area.

### Acknowledgments

The authors are grateful to FAPESP for providing a research grant (number 2009/01403-6). This manuscript is a part of the first author’s Ph.D. thesis, developed in the ‘‘Programa de Pós Graduação em Biodiversidade Vegetal e Meio Ambiente do Instituto de Botânica’’. We would like to thank the staff of the Mogi-Guaçu Biological Reserve for allowing us to use their facilities.

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