Degradative succession of the insects in small rodents in subtropical systems

Sucessão degradativa de insetos em carcaças de pequenos roedores em sistemas subtropicais Sucesión degradante de insectos en pequeñas canales de roedores en sistemas subtropicales

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Abstract

We evaluated the insect succession in small rodent carcasses on the soil extracts (surface vs. buried) in two areas (riparian vs. agricultural) and two seasonal periods (spring vs. summer). Daily, the rodent carcasses were weighed and the insects present were collected in them for counting and family level identification. We also measured the air and soil temperature daily, in addition to the physical and chemical characteristics of the soil. The total of 11,059 individuals from 28 *taxa* was collected. The most abundant *taxa* were Calliphoridae (70%), Formicidae (20%), Muscidae (2%), and Sarcophagidae (2%). Insect richness was higher in spring in riparian vegetation areas; however, decomposition and insect abundance were greater in spring in agricultural areas by the high temperature measured. Vespidae, Sarcophagidae, Muscidae, and Calliphoridae decreased over time with abundance peaks respectively at 3, 7, 7, and 8 days in riparian vegetation areas. Calliphoridae decreased abundance over time with a peak at five days, but Armadillidiidae increased with peak at 6-9 days in agricultural areas. Decay and insect abundance was high in agricultural areas, mainly on the soil surface. Insect richness was high in riparian vegetation areas, especially on the soil surface. Only the superficial layers of soil presented *taxon* indicators and all decreased over time with abundance peaks of Calliphoridae, Sarcophagidae, and Muscidae at 5-8, 7, and 7 days, respectively. This abundance peak of insect families help to understand the degradative succession of the insect community in small rodents in subtropical systems of the neotropical region.

Keywords: Forensic entomology; Carcass decomposition; Insect families; Mus musculus.

Resumo

Avaliamos a sucessão de insetos em carcaças de pequenos roedores no solo (superfície vs. enterrado) em duas áreas (riparia vs. agrícola) e dois períodos (primavera vs. verão). Diariamente, as carcaças foram pesadas e os insetos coletados para contagem e identificação em nível familiar. Também medimos diariamente a temperatura do ar e solo, e características físico-químicas do solo. Foram coletados 11.059 indivíduos de 28 *táxons*. Os *táxons* mais abundantes foram Calliphoridae (70%), Formicidae (20%), Muscidae (2%) e Sarcophagidae (2%). A riqueza de insetos foi maior na primavera das áreas de vegetação riparia, no entanto, a decomposição e a abundância de insetos foi maior na primavera em áreas agrícolas. Vespidae, Sarcophagidae, Muscidae e Calliphoridae diminuiu com o tempo, com picos de abundância respectivamente em 3, 7, 7 e 8 dias nas áreas de vegetação riparia. Caliphoridae diminuiu a abundância ao longo do tempo com pico em 5 dias, mas Armadillidiidae aumentou, com pico em 6-9 dias em áreas agrícolas. A decomposição e abundância de insetos foram maiores nas áreas agrícolas, principalmente na superfície do solo. A riqueza de insetos foi maior na súperficiais do solo apresentaram indicadores de *táxons* e todas diminuíram ao longo do tempo com picos de abundância de Calliphoridae, Sarcophagidae e Muscidae respectivamente em 5-8, 7 e 7 dias. Pico de abundância de famílias de insetos podem servir como base de dados para investigação e ajudar a determinar os intervalos *Post Mortem* em sistemas subtropicais do Neotropical.

Palavras-chave: Entomologia forense; Decomposição de carcaça; Famílias de insetos; Mus musculus.

Resumen

Evaluamos la sucesión de insectos en pequeños cadáveres de roedores en el suelo (superficie vs. enterrados) en dos áreas (ribereña vs. agrícola) y dos períodos (primavera vs. verano). Diariamente, se pesaron los cadáveres y se

recolectaron los insectos para el recuento e identificación a nivel del hogar. También medimos diariamente la temperatura del aire y del suelo, y las características físico-químicas del suelo. Se recolectaron 11.059 individuos de 28 taxones. Los taxones más abundantes fueron Calliphoridae (70%), Formicidae (20%), Muscidae (2%) y Sarcophagidae (2%). La riqueza de insectos fue mayor en primavera en áreas de vegetación ribereña, sin embargo, la descomposición y abundancia de insectos fue mayor en primavera en áreas agrícolas. Vespidae, Sarcophagidae, Muscidae y Calliphoridae disminuyeron con el tiempo, con picos de abundancia respectivamente a los 3, 7, 7 y 8 días en áreas de vegetación ribereña. Caliphoridae disminuyó en abundancia con el tiempo con un pico a los 5 días, pero Armadillidiidae aumentó, con un pico a los 6-9 días en áreas agrícolas. La descomposición y la abundancia de insectos fue mayor en las áreas agrícolas, especialmente en la superficie del suelo. La riqueza de insectos fue mayor en las áreas de vegetación ribereña, especialmente en la superficie del suelo. Solo las capas superiores del suelo mostraron indicadores de taxones y todos disminuyeron con el tiempo con picos de abundancia de Calliphoridae, Sarcophagidae y Muscidae respectivamente en 5-8, 7 y 7 días. Las abundancias máximas de familias de insectos pueden servir como base de datos para la investigación y ayudar a determinar los intervalos post mortem en sistemas neotropicales.

Palabras clave: Entomología forense; Descomposición de la canal; Familias de insectos; Mus musculus.

1. Introduction

Decomposition is a continuous degradation process that releases CO₂ and energy (Bornemissza, 1957; Higley & Huntington, 2009; Medeiros et al., 2015; Rezende et al., 2021) that starts at the death of the organism and end when the carcass is reduced to the skeleton (Probst et al., 2020; von Hoermann et al., 2021; Wolff et al., 2001). In addition, another important process is pedogenesis, which depends on animals that mix organic matter with mineral components in the soil (Moskowitz et al., 2015; Oliveira-Costa, 2013). Both processes, decomposition and pedogenesis, are essential for the ecosystem maintenance (Bornemissza, 1957; Moskowitz et al., 2015) as they allow nutrients and minerals to be reused in the food chain (Alvim et al., 2015; Rezende et al., 2019; Rezende et al., 2021). Also, animal carcass is important for the reproduction of some insects, as *Nicrophorus concolor* (Coleoptera: Silphidae) (Ito, 2021). Therefore, insect colonization in carcasses decomposing with abiotic factors influence (e.g. temperature and moisture) and carcass conditions (e.g., completeness, toxicology, and pathology) may determine the decomposition rates in an ecosystem (Andrade-Herrera et al., 2020; Jales et al., 2020; Moretti et al., 2008; Von Hoermann et al., 2021).

Insects constitute the most abundant and diverse group, and they are found in almost all terrestrial habitats and aquatic environment, except in salt water (Haskell et al., 2008). Insects may travel to distances of three to five kilometers for locating a suitable habitat for laying their eggs (Higley & Huntington, 2009). In this sense, insects are usually the first to find a decomposing body. Therefore, through the insect community structure, it is possible to determine the *Post Mortem* Interval (PMI) and the place of death, as the succession of species on carcasses is also considered a serial replacement class (Haskell et al., 2008; Probst et al., 2020). The degradative succession process may occur on a relatively short time scale; however, succession patterns may vary widely and according to the distinctiveness of the fauna in each local region (Oliveira-Costa, 2013). In this way, the study of the degradation succession process must take into account the local patterns of visiting cadaveric insect fauna (Na & Pt, 2013; Wang et al., 2021). Thereby, this studies are important to understand the degrading succession process and provide subsidies to nutrient cycling from these carcasses (Haskell et al., 2008).

Small rodents can be abundant in subtropical systems and have important ecological functions, such as invertebrate control and seed consumption (Rabinowitz & Nottingham, 1989; Terborgh et al., 1993). Small rodents may be a frequent source to dead organic matter of animal origin in systems where they are abundant (Maestri et al., 2014; Terborgh et al., 1993), due to the fast life cycle compared to other mammals (Feng & Himsworth, 2014). In the subtropical region of the Neotropics, native genre such as *Akodon, Bibimys*, and *Brucepattersonius* are abundant (Maestri et al., 2014). On the other hand, non-native rodents are common and abundant as the genus *Mus*. The genus *Mus* was introduced in Brazil in the European colonization, and was the main representative the species *Mus musculus* that can inhabit both urban and rural areas (Feng &

Himsworth, 2014). However, despite many studies in the tropical region (Cruz & Vasconcelos, 2006; Jales et al., 2020; Oliveira-Costa, 2013; Pujol-Luz et al., 2006), few studies were carried out in the Neotropical subtropical region on the survey of cadaveric fauna (Trivia & Carvalho, 2018).

Studies shows a high insect community variation, mainly by habitat differences (Parry et al., 2016), food availability (T. D. C. Moretti et al., 2008), insect community structure (Tembe & Mukaratirwa, 2021), and temporal changes (Richards et al., 2009). Studies in other regions have shown that Diptera and Coleoptera account for 60% of the fauna associated with decaying carcasses, and *Chrysomya* sp. are an important gender of scavenger fauna (Haskell et al., 2008; Oliveira-Costa, 2013; Tembe & Mukaratirwa, 2021). According to Moretti et al. (2008), a higher diversity of arthropods has been found in cadaveric fauna, and *Chrysomya megacephala* (Fabricius, 1974) may be an important specie of scavenger fauna in southern Brazil (Carvalho & Mello-Patiu, 2008; Trivia & Carvalho, 2018). Also Moretti et al. (2008) show the presence of other necrophagous families that are of forensic interest, such as Sarcophagidae and Drosophilidae. In this way, our objectives were the following: i) compare the richness and abundance of insects throughout the rodent carcasses decomposition, as well as the decomposition process itself by repeated-measures ANOVA, using time in days as a random factor; and ii) evaluate which insects were temporal indicators, positively or negatively, along the degradative colonization process through Threshold Indicator Taxa Analysis between areas (riparian vs. agricultural), periods (spring vs. summer), and soil depths (depth vs. surface).

2. Methodology

2.1 Study area

The study was carried out in a riparian vegetation area (26°54'51"S and 52°41'03"W) and an adjacent agricultural area (26°54'46"S and 52°41'08"W) in spring (October 2020) and summer (January 2021) at both locations in the West (Coronel Freitas City) of Santa Catarina State, Brazil. The agricultural area shows culture rotation throughout the year by pasture, oat, and corn in the Atlantic Forest biome. The climate is of Cfa type with an average annual temperature range of 17 to 20°C and precipitation of 1900 to 2200 mm. The predominant soil is Red Latosol, with a soft undulated relief, with a very clayey texture, which may vary with LitholicNeosol and Haplic Cambisol.

2.2 Environmental characteristics

The chemical and physical properties of the soil were collected at the same points where litter bags were incubated. The samples were collected at depths from 0.0 to 10 cm for the chemical and physical characteristics of the soil. Air and soil temperatures were measured daily using a thermometer and the soil moisture by drying at 80°C to constant weight ([initial weight – final weight]/initial weight). Rainfall was measured by a portable rain gauge during the experiment period. The chemical characteristics of the soil were analyzed by the Soil Analysis Laboratory of the Agricultural Research and Rural Extension Company of Santa Catarina - EPAGRI in Chapecó/SC. Clay was determined by densiometry, pH water and SMP by potentiometry, P by Mehlich-1/colorometry, K by Mehlich-1/flame photometer, MO by spectroscopy, and Al, Ca, and Mg by KCl/atomic absorption spectrophotometry. The characterization of the soils was performed according to the methodology proposed by Tedesco et al. (1995).

2.3 Sampling unit

The insect community was collected from rodent carcasses of the *Mus musculus* species (Schwartz; Schwartz, 1943), weighing between 27.33g and 35.70g (mean of 31.11g). All rodents were donated by the Unochapecó Bioterism Center (Ethics Committee - CEUA no. 010/20) and submitted to euthanasia by cervical dislocation. At each area, rodent carcasses were then placed in soil covered by cages (37 cm x 27 cm; 3 cm mesh) and some weight was providing above each cage to keep cages

constant and protect carcasses from the attack of large animals 10m away from each other. Insects attracted to rodent carcasses were collected daily from 13:00 to 14:00 hour. The collected insects were preserved in 70% ethanol after washing them in tap water, and then transferred in labeled tubes to the laboratory.

2.4 Morphological identification

The collected insects were examined under stereomicroscope (Zeiss Stemi 305 binocular) and identified following the morphological identification key of Byrd and Castner (2001) and Carvalho and Mello-Patiu (2008). A part of the collected insects was preserved and deposited in the ecological entomology laboratory of Community University of the Chapecó Region - Unochapecó.

2.5 Experiments

In the experiment I, we tested the degradative succession between periods (spring vs. summer) and areas (riparian vs. agricultural). The rodent carcasses were used at six points on the surface of riparian vegetation areas and in six points on the surface of agricultural areas over two periods (spring vs. summer). The sampling effort totaled 24 rodent carcasses (6 points x 2 areas x 2 periods = 24 carcasses – half male and half female).

In the experiment II, we tested the degradative succession between soil layers – surface vs. buried only in riparian vegetation areas. Rodent carcasses were used at three points, with three carcasses placed on the soil surface, three carcasses buried in the soil of riparian vegetation areas, three carcasses placed on the soil surface, and three carcasses buried in the soil of agricultural areas in two periods (spring vs. summer). The sampling effort totaled 24 rodent carcasses (3 points x 2 areas x 2 periods x 2 depths = 24 carcasses – half male and half female).

2.6 Data analysis

The normality of the data (function "ks.test" from the "dgof" package for R) was assessed with the Kolmogorov-Smirnov test while the homogeneity of variances was assessed with Levene's test and the data were transformed when necessary $(\log_10(x+1))$. In the experiment I, seven Repeated-Measure ANOVA models were created (RM-ANOVA; "aov" + Error (repeated measure factor = days) function from "stats" package for R) to test i) the remaining mass of rodent carcasses by ([initial weight – final weight] * 100 / initial weight), ii) insect richness, iii) insect abundance, temperature of (iv) air and v) soil), vi) rainfall and vii) soil moisture (response variables) between areas (riparian vs. agricultural), periods (spring vs. summer), and interaction between variables (explanatory variable) (Crawley, 2007). Other seven RM-ANOVA models from experiment II, with the same response variables to test areas (riparian vs. agricultural), soil depth (surface vs. buried), and interaction between variables (explanatory variable) also used days as a random repeat factor (Crawley, 2007). Contrast analysis ("coms" function from the "coms.R" package for R) was used to discriminate among explanatory variables (Crawley, 2007). In this contrast analysis (orthogonal), the explanatory variables were increasingly ordered and tested pairwise with the closest values, sequentially adding values to the model month with no differences and testing with the following, in a stepwise simplification of the model (Crawley, 2007).

Threshold Indicator Taxa Analysis ("titan" function from the "TITAN2" package for R) was used to track the relationship (positive or negative) and check in which period in days the insect community on rodent carcasses are associated across four individual models (riparian vs. agricultural and surface vs. buried) (Baker & King, 2013). In experiment I, the studied areas (riparian and agricultural) were tested over the days, using the seasonal periods as replicates. In experiment II, the studied soil depths (surface and buried) were tested over the days, using the seasonal periods as replicates. The Threshold Indicator Taxa Analysis is based on Indicator Species Analysis (Dufrêne & Legendre, 1997) and the Change Point Analysis

(King & Richardson, 2003). Based on the frequency of occurrence and relative abundance of the insect community in rodent carcasses, changes were detected in the distribution of species in environmental gradients by Threshold Indicator Taxa Analysis (King et al., 2011). In this way, Threshold Indicator Taxa Analysis highlights the change point based on alterations in mean and variance of response variable, with z - (negative responders, or decliners) and z + (positive responders, or increasers) of the species being associated with the lowest and highest values of the environmental gradient analyzed, respectively (Baker & King, 2010). These changes in the distribution of species from Threshold Indicator Taxa Analysis were detected in space or over time, and associate individual species with different environmental and temporal scores (King et al., 2011). In the Threshold Indicator Taxa Analysis, only species with a frequency of at least 3 were considered. Also, for the Threshold Indicator Taxa Analysis, the significance was set at p (IndVal) ≤ 0.05 , and purity and reliability set at ≥ 0.70 were considered (Baker & King, 2010).

3. Results

3.1 Environmental characteristics

In experiment I, the air temperature was significantly high in riparian vegetation areas (Repeated Measure ANOVA; $F_{(1,204)} = 8.77$; p = 0.003), while the temperature of soil and rainfall were significantly high in the agricultural area (RM ANOVA; $F_{(1,204)} = 50.21$; p < 0.001). Both temperatures (air and soil) were significantly high in spring (RM ANOVA; $F_{(1,204)} = 62.39$; p < 0.001; $F_{(1,204)} = 20.72$; p < 0.001, respectively). On the other hand, rainfall was significantly high in summer (RM ANOVA; $F_{(1,204)} = 5.51E+29$; p < 0.001). Soil moisture changed significantly only in the interaction between areas and seasonal periods (RM ANOVA; $F_{(1,204)} = 3.99$; p = 0.047), with the high values in the summer of the agricultural areas compared to the spring of the agricultural areas.

In experiment II, air temperature was significantly high in riparian vegetation areas (RM ANOVA; $F_{(1,214)} = 11.57$; p = 0.001), while the temperature of soil and rainfall were significantly high in the agricultural area (RM ANOVA; $F_{(1,214)} = 56.22$; p < 0.001). The temperature of the soil was significantly high at surface compared to deep layers (RM ANOVA; $F_{(1,214)} = 7.52$; p = 0.007). Soil moisture was significantly high in riparian vegetation areas compared to agricultural areas (RM ANOVA; $F_{(1,214)} = 7.52$; p = 0.007). Soil moisture was significantly high in riparian vegetation areas compared to agricultural areas (RM ANOVA; $F_{(1,214)} = 5.10$; p = 0.025). Also, soil moisture significantly changed in the interaction between areas and depth, with high values in deep layers of riparian vegetation areas compared to the superficial soil layers of the agricultural areas (RM ANOVA; $F_{(1,214)} = 25.95$; p < 0.001).

Comparing the soil surface characteristics on the two areas, we observe a high values of clay densiometry (41 ± 3.0 vs. 40 ± 2.6 %; mean and standard deviation), pH (5.5 ± 0.2 vs. 5.0 ± 0.1), organic matter (4.1 ± 0.6 vs. 3.4 ± 0.1 %), Ca (14.6 ± 2.2 vs. 13.0 ± 1.1 cmolc/dm³), and cation exchange capacity (22.4 ± 2.1 vs. 20.8 ± 1.5 cmolc/dm³) in riparian vegetation areas compared to agricultural areas, respectively. On the other hand, P (24.7 ± 3.7 vs. 22.3 ± 4.1 mg.dm³), K (345.6 ± 22.6 vs. 292.4 ± 33.6 mg.dm³), Al (0.23 ± 0.1 vs. 0.20 ± 0.1 cmolc/dm³), and Mg (3.8 ± 0.1 vs. 3.6 ± 0.3 cmolc/dm³) show the high values in agricultural areas compared to riparian vegetation areas, respectively. We also found low values of clay densiometry (31.6 ± 6.5 %; mean and standard deviation), organic matter (3.4 ± 1.2 %), Ca (12.6 ± 1.4 cmolc/dm³) and cation exchange capacity (20.6 ± 2.3 cmolc/dm³) in deep layers of riparian vegetation areas compared to soil surface. On the other hand, the high values of pH layers of riparian vegetation areas compared to soil surface.

3.2 Insect community

The total of 11,059 individuals from 28 *taxa* were collected (Table 1). The most abundant *taxa* were Calliphoridae (70%), Formicidae (20%), Muscidae (2%), and Sarcophagidae (2%). In total, we found 6,605 individuals (36.6±108 ind. mean

and standard deviation) in the riparian vegetation area and 4,454 (24.7 ± 62 ind.) individuals in the agricultural area. Moreover, evaluating the year seasons, we found 6,376 individuals (35.4 ± 97 ind. mean and standard deviation) in summer and 4,683 (26.1 ± 79 ind.) individuals in spring.

Table 1. Occurrence of insects collected on rodent carcasses in riparian vegetation areas and agricultural areas in two periods
(spring - spr and summer - sum) of buried (bur) and soil surface (sur) rodent carcasses.

· · · ·	riporion ogr			ultural	arian agricultural			
	riparian bur sur		agricultural bur sur		spr	sum	spr sum	
Insecta	oui	541	oui	501	5.01	Sum	spi	Sum
Dermaptera								
Forficulidae	2	0	0	2	0	2	0	2
Orthoptera	2	0	0	2	0	2	0	2
Gryllidae	1	0	0	0	1	0	0	0
Isopoda	1	0	0	0	1	0	0	0
Armadillidiidae	2	1	11	12	0	3	0	23
Hemiptera	2	1	11	12	0	5	0	23
Ectrichodiinae	0	1	0	0	0	1	0	0
Hymenoptera	0	1	0	0	0	1	0	0
Pompilidae	2	5	0	0	2	5	0	0
Vespidae	7	14	0	0	9	12	0	0
Sphecidae	2	3	0	0	5	0	0	0
Formicidae	101	1672	4	1138	163	1610	702	440
Coleoptera	101	1072	4	1156	105	1010	702	440
Cerambycidae	13	11	0	0	24	0	0	0
Cleridae	5	3	0	6	24 8	0	0	6
Coccinellidae	0	5	0	0	0 1	0	0	0
Curculionidae	1	0	0	0	1	0	0	0
Chrysomelidae	0	6	0	4	5	1	0	4
Dermestidae	1	0	0	4	0	1	0	4
Glaresidae	0	1	0	0	1	0	0	0
Hydrophilidae	1	0	0	0	1	0	0	0
Leiodidae	2	5	0	0	2	5	0	0
Neriidae		1	0	0	2 1	0	0	0
Nitidulidae	25	3	0	0	1 26	2	0	0
Noteridae	23 13	5 5	0	1	20 16	$\frac{2}{2}$	0	1
	4	5 8	2	1	2	2 10	0	9
Scarabaeidae	4	8 3	2	12	0	3	0	
Staphylinidae	0	3	I	12	0	3	0	13
Lepidoptera	1	0	0	0	1	0	0	0
Lycaenidae	1	0	0	0	1	0	0	0
Diptera	222	01.40	010	5100	004	1 402	0777	0.000
Calliphoridae	229	2148	212	5193	884	1493	2777	2628
Drosophilidae	2	2	0	0	4	0	0	0
Muscidae	28	50	0	0	20	58	0	0
Sarcophagidae	14	52	0	0	24	42	0	0
Tephritidae	2	1	0	0	3	0	0	0

Source: Authors.

In experiment I, the remaining masses were high in the spring of riparian vegetation areas compared to the spring of the agricultural area (Table 2a; Figure 1a). The insect richness was high in riparian vegetation areas (Table 2b; Figure 1c), mainly in spring (Table 2b; Figure 1c). On the other hand, the insect abundance was high in agricultural areas (Table 2c; Figure 1e), mainly in summer (Table 2c; Figure 1e). Threshold values over time ranged between 2.5 to 10 (sumz +) and 1.5 to 7 (sumz -). According to the significance criteria adopted here ($p \le 0.05$, reliability and purity ≥ 0.70), 5 *taxa* (Vespidae, Sarcophagidae, Muscidae, Calliphoridae, and Armadillidiidae) had a systematic relationship with the time in days. In the riparian vegetation areas, the families Vespidae, Sarcophagidae, Muscidae, and Calliphoridae showed a decrease in abundance to time with respective peaks in 3, 7, 7, and 8 days (Figure 2a and 2b). In agricultural areas, the family Calliphoridae also showed a decrease in abundance to time with peak in 5 days (Figure 2c e 2d), butthe family Armadillidiidae showed an

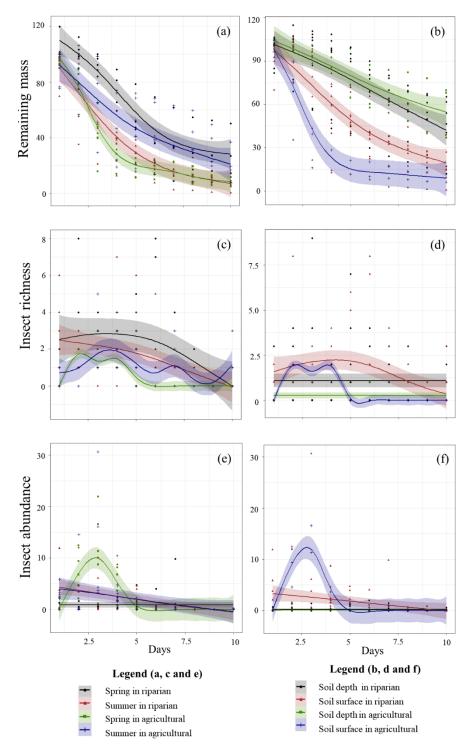
increase in abundance to time with peak in 6-9 days (Figure 2c e 2d).

Table 2. Result of Repeated Measure ANOVA and Contrast Analysis (Post Hoc) of the remaining mass (a), insect richness (b), and insect abundance (c) in rodent carcasses between areas (riparian vs. agricultural), periods (spring vs. summer), and interactions between factors in Experiment I.

		1				
	Df	Sum Sq	Mean Sq	F value	Pr(>F)	Contrast analysis
A. remaining n	nass					
Error x days	1	135183	135183			
area	1	516	516	1.65	0.210	
period	1	602	602	1.92	0.167	
area x period	1	20927	20927	66.82	< 0.001	agricultural in spring < riparian in spring
Residual	204	63887	313			
B. insect richn	ess					
Error x days	1	61.1	61.1			
area	1	40.8	40.9	20.63	< 0.001	agricultural <riparian< td=""></riparian<>
period	1	0.0	0.0	0.01	0.927	с .
area x period	1	15.5	15.5	7.84	0.006	agricultural in spring <riparian in="" spring<="" td=""></riparian>
Residual	204	403.9	2.0			
C. insect abune	dance					
Error x days	1	61.1	61.1			
area	1	40.8	40.9	20.63	< 0.001	riparian <agricultural< td=""></agricultural<>
period	1	0.0	0.0	0.10	0.927	
area x period	1	15.5	15.5	7.84	0.006	agricultural in spring <agricultural in="" summer<="" td=""></agricultural>
Residual	204	403.9	2.0			

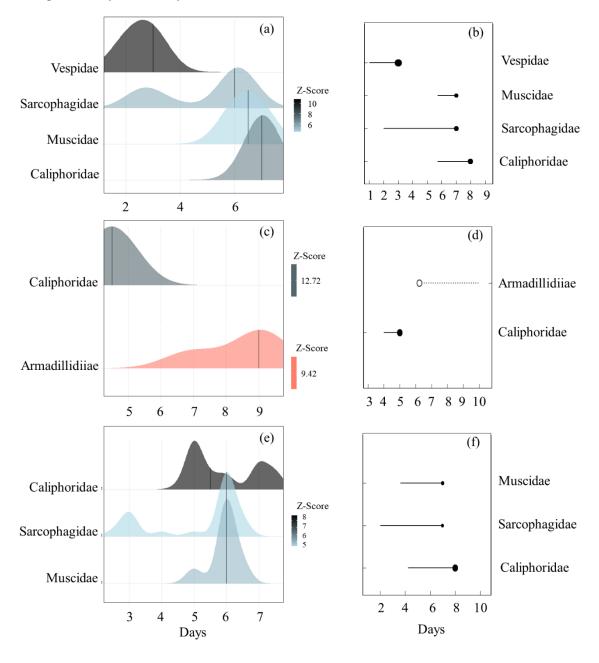
Source: Authors.

Figure 1. Remaining mass of rodent carcasses ("a" and "b"), insect richness in rodent carcasses ("c" and "d"), and insect abundance in rodent carcasses ("e" and "f") between areas (riparian vs. agricultural) periods (spring vs. summer; experiment I), and soil depths (depth vs. surface; experiment II). The solid lines are the representations of the linear model, while the remnants of the same color are the 95% confidence intervals for each model.



Source: Authors.

Figure 2. Values of the largest indicator z value in insect abundance in rodent carcasses change points by probability abundance function of the nBoot bootstrapped replicates (n=500) along days from riparian vegetation areas (a), agricultural areas (b), and surface in both areas (c). *Taxa* names are arranged discretely along the y-axis based on the rank order of the central tendency of the probability abundance function for z- (dark colors negative, decliners) and z+ (red colors positive, increasers), respectively (a, b, e c). The black symbols represent associated species and sumz-, represented by the black cycle, and sumz +, represented by the white cycle (b, d and f).





In experiment II, the remaining masses were high in agricultural areas compared to the riparian vegetation areas (Table 3a; Figure 1b), mainly in soil surface compared to soil deep layers (Table 3a; Figure 1b). Insect richness was high in the riparian vegetation areas (Table 3b; Figure 1d) and in soil surface compared to soil deep layers (Table 3b; Figure 1d). On the

other hand, the insect abundance was high in agricultural areas (Table 3c; Figure 1f), also mainly in soil surface compared to soil deep layers (Table 3c; Figure 1f). Threshold values over time ranged between 3 and 10 (sumz +) and between 2 and 7.3 (sumz -). We observe 3 *taxa* (Sarcophagidae, Muscidae, and Calliphoridae) with a systematic relationship with the time in days. Only the superficial layers of soil presented *taxon* indicatorsand all decreased with time. The abundance peaks of Calliphoridae, Sarcophagidae, and Muscidae were respectively in 5-8, 7, and 7 days.

Table 3. Result of Repeated Measure ANOVA and Contrast Analysis (Post Hoc) of the remaining mass (a), insect richness (b), and insect abundance (c) in rodent carcasses between areas (riparian vs. agricultural), soil depths (depth vs. surface), and interactions between factors in Experiment II.

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	Contrast analysis
A. remaining mass						
Error x days	1	112786	112786			
area	1	8435	8435	23.83	< 0.001	riparian <agricultural< td=""></agricultural<>
soil depths	1	35829	35829	101.21	< 0.001	surface < buried
Local x soil depths	1	1294	1294	3.66	0.057	surface in riparian < buried in agricultural
Residual	214	75754	354			
B. insect richness						
Error x days	1	15.2	15.2			
area	1	54.5	54.5	24.07	< 0.001	agricultural <riparian< td=""></riparian<>
soil depths	1	9.8	9.8	4.35	0.038	buried <surface< td=""></surface<>
area x soil depths	1	0.6	0.6	0.25	0.619	buried in agricultural <surface in="" riparian<="" td=""></surface>
Residual	214	484.8	2.3			
C. insect abundance	•					
Error x days	1	15.2	15.2			
area	1	54.5	54.5	24.07	< 0.001	riparian <agricultural< td=""></agricultural<>
soil depths	1	9.8	9.8	4.35	0.038	buried <surface< td=""></surface<>
area x soil depths	1	0.6	0.6	0.25	0.619	
Residual	214	484.8	2.3			

Source: Authors.

4. Discussion

4.1 Degradative succession between periods

In experiment I, high decomposition occurred in the seasonal period (spring) and in the area (agricultural) with higher soil temperatures. The higher number of cloudy days and constant rains in summer can reduce the direct sunlight incidence on the soil, decreasing soil temperature compared to spring (Ota et al., 2019). The high soil temperatures in the agricultural area are perhaps due to the lack of coverage in tall plant extracts (De Frenne et al., 2019; Rezende et al., 2017). On the other hand, riparian vegetation areas keep a constant mild microclimate (Quintão et al., 2013; Rezende et al., 2017) compared to agricultural areas. High temperatures may accelerate the ecosystem metabolism by stimulating the decomposing community activity in rodents' carcasses (Navarro et al., 2013; Al-Mekhlafi et al., 2020; Probst et al., 2020). As an example, lower and constant soil temperatures may slow down the metabolic activity and oviposition of detritivorous insects, such as Diptera and Coleoptera, and as calliphorids (Souza & Linhares, 1997; Pujol-Luz et al., 2006). Insects are pecilothermic, that is, their temperature varies according to the environment. High soil temperatures may also accelerate the immature development (Kutcherov et al., 2019) and, consequently, the food resources consumption (Kotzé et al., 2016), explaining the high decomposition in the warmer area and period. Furthermore, the high abundances observed in the agricultural area may demand greater food resource consumption (Rodríguez & Liria, 2017; Zeariya & Kabadaia, 2019), helping to explain the faster decomposition. Finally, there is the increase in invertebrate richness in the riparian vegetation area, possibly because carcasses take a longer time to decay (Moretti et al., 2008) and shows that soil temperatures and the abundance of detritivorous insects are more important in accelerating the decomposition process.

In experiment I, the families Vespidae, Sarcophagidae, Muscidae, and Calliphoridae in riparian vegetation areas and Calliphoridae in the agricultural area decreased their abundance over time. The families Calliphoridae, Sarcophagidae, and Muscidae have a preference for colonizing soft tissues during the initial stages of decomposition (Cruise et al., 2018; Tembe & Mukaratirwa, 2021). Furthermore, small carcass decreases the resource available, concentrating the greatest insects' abundances in early stages of decomposition (Al-Mekhlafi et al., 2020; Zeariya & Kabadaia, 2019). Also, the presence of hymenoptera in rodent carcasses is common, especially species of Vespidae families (Haskell et al., 2008). Vespidae may feed on bodily fluids and dipteran larvae (Moretti & Ribeiro, 2006), justifying their high abundance in early decomposition stages. Some Vespidae species may chew the tissues of carcass natural cavities (nose, mouth, and anus), looking for liquids or small pieces of meat (Herdina et al., 2016). On the other side, Armadillidiidae showed an increase in abundance over time in agricultural areas, probably to use the carcasses as a refuge against dissection (Haskell et al., 2008; Oliveira-Costa, 2013).

4.2 Degradative succession between soil layers

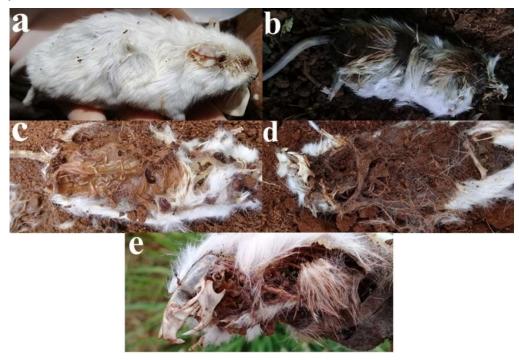
In experiment II, high decomposition occurs in riparian vegetation area, mainly in the superficial layers of the soil, due to differences in moisture. Less vegetation cover increasestemperature variations and decreases soil moisture (De Frenne et al., 2019; Rezende et al., 2017). Lower humidity may make the environment more hostile to specialist organisms, such as detritivorous insects subject to dissection (Sfenthourakis and Hornung, 2018). High moisture in topsoil may accelerate insect colonization compared to lower soil layers. Moreover, richness was high in riparian vegetation areas, especially in superficial soil layers. Soil can act as a barrier to the detritivorous insects, delaying colonization and decreasing richness by reducing decomposition of the buried soil extract eight times compared to the soil surface. On the other hand, insect abundance was high in agricultural areas. Insect abundance may have been influenced by the lower presence of predators. Fewer ants in the carcasses, female of Diptera were able to colonize the available resource more intensively, indicating that the specific insect composition may be important for the decomposition process.

In experiment II, only the superficial layers of the soil showed indicator *taxa*, all (Calliphoridae, Sarcophagidae, and Muscidae) decreasing abundance with time. Succession studies of cadaveric fauna highlight that species of families Calliphoridae, Sarcophagidae, and Muscidaecan stay longer on carcass, but decreasein abundance with time (Cruise et al., 2018; Haskell et al., 2008; Parry et al., 2016; Tembe & Mukaratirwa, 2021). They are attracted by the odors that are produced in the early stages of decomposition (Oliveira-Costa, 2013). These peaks also explain the influence of the carcass condition, due to the high humidity (Tembe & Mukaratirwa, 2021), where rodents rot faster.

4.3 Drying process of rodent carcasses

We observed five decomposition stages of *Mus musculus*: initial decay, putrefaction, black putrefaction, fermentation, and dry decay (Figure 3), as also observed by Moretti et al. (2008). Also, some rodent carcasses in decomposition process may undergo skeletonization or mummification (Figure 4). Mummification is a drying process, dependent on extreme factors of temperature and humidity, which inhibits bacterial activity (Al-Mekhlafi et al., 2020; Tembe & Mukaratirwa, 2021). As an example, high temperatures and low humidity accelerate the decomposition of rodent carcasses, but cause the mummification of some carcasses (Al-Mekhlafi et al., 2020; Haskell et al., 2008). In the present study, the drying process of some carcasses was observed from the 8th day of summer in riparian vegetation areas by high temperatures. In addition, the attractiveness of flies is not hindered, since they were found in the process of drying the carcasses in this study.

Figure 3. Decomposition stages of *Mus musculus*: (a) initial decay, (b) putrefaction, (c) black putrefaction, (d) fermentation and (e) dry decay.



Source: author.

In this way, the drying process of carcasses may then favor the Coleoptera and Formicidae, which colonize the carcasses in the final stages, as they have a preference for cartilage (Haskell et al., 2008). In addition, species of the Coleoptera order families are able to respond to soil changes (Parry et al., 2016). In studies carried out with carcasses, the Coleoptera and Formicidae families were found in advanced stages of decomposition (Cruise et al., 2018; Tembe & Mukaratirwa, 2021; Zeariya & Kabadaia, 2019). In this experiment, the presence of beetle families such as Cerambycidae, Chrysomelidae, Nitidulidae, Noteridae, and Dermestidae was mainly observed in the final stages.

Figure 4. Drying of rodent carcasses.



Source: author.

5. Conclusion

Finally, we can conclude that the temperature and moisture of the soil were the main drivers of carcass decomposition over areas and periods. We also found that insect abundance is more important than insect richness to accelerating carcass decomposition. We also observed that buried carcasses in soil decompose more slowly due to the physical barrier caused by the soil. Most *taxon* indicators (Vespidae, Sarcophagidae, Muscidae, and Calliphoridae) declined over time, except Armadillidiidae, probably to use the carcasses as a refuge against dissection. Lastly, Coleoptera and Formicidae colonize the carcasses in the final stages of decomposition due to the facility of cartilage use. This abundance peak of insect families can serve as the baseline data for investigation and help to determine *Post Mortem* intervals on Neotropical Subtropical systems.

During the decomposition process, it is also important to take into account the influence of dipteran larvae on the carcasses, as they produce metabolic heat. This heat can be due to the increase in temperature caused by the aggregation and movement of larvae. Therefore, it would be interesting to measure the temperature of the larvae in the carcasses, using a thermometer, to better understand how the decomposition process takes place inside the carcass.

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