1 Effects of productive management of an agroforestry system on the associated diversity

2 of parasitoids (Hymenoptera: Braconidae)

3 Cecilia Marisol Pech Cutis, Luis Enrique Castillo Sánchez, Jorge Rodolfo Canul Solís,

4 Ermilo López Coba, Nery Maria Ruz Febles, Maria Jose Campos-Navarrete*

5 Tecnológico Nacional de México/ Instituto Tecnológico de Tizimin. Final of Cupul Airport

6 S/N C.P. 97700 Tizimin, Yucatán, Mexico.

7 *maria.campos@ittizimin.edu.mx

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9 Summary

10 Tropical agroecosystems have emerged from the continuous modification of natural 11 environments, as a sustainability alternative for food production and biodiversity 12 conservation. This work explores how the diversity of parasitoids is modified in environments where plant diversity is limited e.g. crops and when these are adjacent to 13 secondary vegetation, i.e. a scenario fragmented continuously in a limited space. It was found 14 that there is no direct effect of plant diversity in the group of parasitoids studied; but the 15 16 number of specialist species is high, which indicates that in diversified agroecosystems these probably function as remnants of natural habitat or as a refuge for parasitoids that disperse 17 18 to different types of management within the agroecosystem. Therefore, it is necessary to consider in future studies the controls exerted by the plant diversity effects bottom-up and 19 consumer top-down. Adding to this the context of the interactions that occur in 20 21 agroecosystems.

22 Key words: agroecosystems, biodiversity, Braconidae, conservation, specialists

23 Introduction

Agroecosystems arise in the tropics derived from the loss of vegetation cover, the conversion 24 and fragmentation of natural areas in ecosystems. These activities modify the species that 25 inhabit and depend on vegetation, with effects that depend on the range of the species and its 26 habitat requirements (Scolozzi & Geneletti, 2012; Campos-Navarrete et al 2015a). On the 27 other hand, the speed in the loss of species has increased the interest in the study of 28 biological diversity in agroecosystems, mainly insects, because they constitute the most 29 30 important fraction of the diversity of a territory since they provide multiple ecological services to ecosystems (Kleinet al 2002). 31

Parasitoids are known for their high number of species and their habits that result in the 32 provision of regulatory services for the populations of their hosts (Lepidoptera, Coleoptera 33 34 and Diptera) (Shaw & Huddleston, 1991; Abdala-Roberts et al 2016a). In agroecosystems, the study of these organisms is interesting from the anthropogenic point of view (integrated 35 36 pest control) and even little known in terms of the mechanisms that regulate it (Nicholls, 37 2008; Schmidt et al 2003; Thies et al 2005). Also in agroecosystems, plant species diversity has effects on secondary productivity comparable to natural systems. In this sense, it has been 38 observed that environments with greater diversity of plant species promote increases in 39 richness and abundance in trophic levels (Abdala-Roberts et al 2016 b; Castillo-Sánchez et 40 al 2019). 41

The effect of land use, in conjunction with the associated plant diversity at higher trophic levels, has been explained primarily because greater diversity, by generating a more complex environment, consequently offers a greater number of shelters and prey (Russell, 1989; Obermaier, et al 2008; Moreira et al 2016),which in turn generates increases in predation

rates, causing a reduction in the abundance of parasitoid prey ("up-down" effects). Evidently, 46 47 this type of effects are highly relevant to be considered in the design of productive systems such as forestcrops, due to their potential in pest control (Russell, 1989; Abdala-Roberts et 48 al 2015: 2016a). This last factor can reduce or increase the abundance of prev (herbivores) 49 depending on characteristics such as the specialization of their diet (generalists vs. 50 specialists) and their interactions (Campos-Navarrete et al 2015b). For example, there is 51 52 evidence that for specialists the effects of an increase in diversity can be negative, due to the low density of their priority resource (Hambäck et al 2014). 53

In contrast, for generalists the effects of increased diversity can vary and, in some cases, can be positive due to their mixed diet and increased availability of places of refuge (Unsicker et al 2008; Castagneyrol et al 2013). In this sense, the presence of herbivores can mediate their interaction with the next trophic level of consumers where parasitoids are included (Abdala-Roberts et al 2016b).

The present work explores the effect of four productive areas in an agroecosystem of multiple production in relation to the diversity of parasitoids (Braconidae) associated. Particularly changes in richness, abundance, similarity in areas and in parasitism strategies represented by the proportion of koinobiont (specialist) and idiobiont (generalist) species. Trying to infer how the diversity of parasitoids is modified in environments where plant diversity is limited e.g. crops and when these are adjacent to secondary vegetation, i.e. a scenario with continuously fragmented in a limited space.

66 Materials and Methods

Study area. The present work was carried out in the east of Yucatán in the municipality of Tizimin, Yucatán, Mexico. This area is characterized by the conversion of land use from native jungle to grasslands for the production of pasture for cattle feed. Livestock represents 30% of the economic activity of this area (INEGI 2015). The agroecosystem of study is located in the area of agricultural and livestock production of the TecNM Campus Tizimin, located at the end of the airport Cupul s/n C.P. 97700 with the coordinates 21°09′29" N 88°10′21"W.

The agroecosystem has three areas of crops: a plantation of *Cocos nucifera* "PC" (monoculture), a plantation of Citrus *lemon* "PL" (monoculture), in the livestock production area is located the grassland area with star grass *Cynodon sp.* named grassland "PT" and the fourth type is the matrix of secondary vegetation with more than 30 years of recovery "VS". This contrast in a limited space provides a frequent scenario today, originated by the processes of fragmentation, originating contrasting sites.

Fieldwork. For the capture of insects were used malaise traps, which capture large numbers 80 of organisms, widely recommended and used to capture parasitic Hymenoptera (Noves, 81 82 1982) (Nieves-Aldrey & Castillo., 1991). The orientation of the trap was from north to south, because it is more effective if the openings are placed in the position where the wind comes 83 from. For the placement of the trap it was considered that it was far from the edges of each 84 area, in order to reduce the edge effect and between traps there was a distance of 500 m, to 85 ensure the independence of the samples. The traps remained active every day during the 86 period from October 2015 to March 2016, were checked every fortnight and the specimens 87 placed in jars with 70% alcohol. 88

Identification of the specimens: The samples were identified in the Laboratory of Parasitology of the TecNM Campus Tizimín, before the identification the braconids were separated from the rest of the insects, later they were counted and labeled with the corresponding data, for their identification the specimens were assembled according to the technique traditionally used for these organisms. Identification was carried out using the taxonomic key for Braconidae (Wharton et al 1998).

The assembly was carried out using entomological pins correctly placing each insect, which were also labeled for a systematic control of these. The reason for mounting is that in this way it is easier to observe insects when they are dry, since when they are wet it is difficult to observe certain characteristics.

99 The material was determined up to the taxonomic level of genus subsequently the concept of morphospecie was used in which (Simpson, 1961; Mayr & Ashlock., 1991; Delfín & Burgos 100 2000) mention that due to the existing difficulty in determining parasitoids at a specific level 101 102 it is advisable to use this concept to identify and separate individuals who present different morphological characteristics. In addition, morphospecies were classified according to their 103 parasitism strategies as koinobiont species (specialists) and idiobionts (generalists), with the 104 105 help of specialized literature (Goulet & Huber, 1993). All the collected material is sheltered in the TecNM Campus Tizimín. 106

Data analysis. We performed a classical diversity analysis in the braconid community in all
 four areas using the Shannon-Wiener diversity index. This index expresses the uniformity of
 the values of importance across all species in the sample; assumes that all individuals are

110 randomly selected, and that all species are represented in the sample (Ludwing and Reynolds,

111 1988; Magurran, 1988).

The similarity between the braconid communities was estimated using the Sorensen index (Krebs, 1989) which is based on the presence/absence of species and the number of species whether common or rare (Spellerberg, 1991). It relates the number of species in common to the arithmetic mean of the species at the sites (Magurran, 1988).

To estimate the representativeness of the species richness of the samples in each type of vegetation, the EstimateS version 9 program was used, which presents specialized estimators for different types and sizes of data. We used ice wealth estimators based on the number of rare species (those observed in less than 10 sampling units) and jacknife first order which is based on the number of unique species (Colwell and Coddington, 1994).

With the data of the morphospecies recorded by area, using relative abundance, as the 121 response variable, the change between the four areas was evaluated. This is through a model 122 123 that explores the quantitative response of braconids to areas. This model used as the main factor the area with four levels (PC, PL, PT, VS). The second model explored the parasitism 124 strategies of braconist species, using as the main factor parasitism strategy with two levels 125 126 (Idiobionts "I" and Koinobiontes, "K"), in relation to abundance. The third model explored the interaction between the two main factors mentioned above in the relative abundance of 127 braconids. We fitted all models using the Penalized Quasi-Likelihood method (Crawley 128 2007). We conducted GLMM analyses using the R statistical package v. 3.01. (R Core Team 129 2020). We used a posteriori contrasts to test for differences among pairs of means for a given 130 factor within each level of the other factor (Crawley 2007). 131

132 results

Collected specimens. A total of 2031 specimens of braconids belonging to 20 subfamilies, 47 genera and 140 morphospecies (Apendice 1) were collected. In general, the subfamily Microgastrinae was the most abundant representing 49% of the total number of individuals captured, other abundant subfamilies were Doryctinae and Hormiinae with 8% and 6% individuals respectively, the least abundant subfamilies were Ichneutinae and Macrocentrinae which are represented by an individual equivalent to 0.05% (Table 1).

Richness and abundance of species by typearea. Wealth in each area of the agroecosystem 139 was distributed as follows from highest to lowest. PL with the largest number of subfamilies, 140 141 followed by PC, VS and PT (Table 1). As for gender from highest to lowest PL, PC, VS and PT (Table 1). The morphospecies were distributed from highest to lowest in the areass 142 in the following order first PL, second PC and with the same number PT and VS (Table 1). 143 Finally, for the number of individuals the largest was concentrated in PL, followed by PC, 144 145 PT and finally VS (Table 1). For wealth in general it was observed that the PL and 146 PC, considereds monocultures concentrated the largest number of individuals and taxonomic wealth, while VS occupied the third site and with the least wealth and abundance 147 148 the PT with the simplest vegetation structure composed solely of grass.

The general pattern for the number of rare and unique morphospecies, as well as species diversity according to the Shannon Index by area is described below. For rare species the order of greatest number was in PT, PL, PC and VS (Table 2). As for unique species from highest to lowest these were found in PC, PT, PL and finally VS (Table 2). In general, it is observed that the VS considered a polyculture and with greater complexity was the one with

the lowest number of rare and unique species present. This contrasts with what was found
for the Shannon-Wienner diversity index, where the highest diversity was observed in PL,
followed by VS, finally PC and PT (Table 2).

Specific wealth estimation: The wealth estimators used were ICE and Jacknife of the first order. According to the ICE, 60% of the species were captured on average for the agroecosystem, while for the jacknife of the first order 70% were captured according to the estimate for the agroecosystem. Table 3 shows in particular the percentage of wealth estimators by area.

Similarity of communities: The Sorensen index indicated that the sites that presented the greatest similarity were PC with VS, followed by PT with VS, in third place, PL with VS and those that presented the least similarity were PC with PT (Table 4). The above highlights the importance of conserving remnants of secondary vegetation in agroecosystems, since it was observed that VS is similar to crop areas.

Effect of the areas on the abundance. As for the quantitative response in the abundance 167 168 relative to the area of cultivation and the functional role of the braconist in this system, it was observed that the Area has a statistically significant effect on abundance (F $_{1,3}$ = 2.8125 p = 169 0.000). Higher values of relative abundance were observed in PC, PT and VS with respect 170 171 to PL (Figure 1). As for the parasitism strategy, a significant effect was found on abundance (F₁₂=8.5947 p =0.000), being greater in koinobionts (specialists) compared to idiobionts 172 (generalists) (Figure 2). However, no interaction effect was found between the Area and 173 parasitism strategies ($F_{1,1}$ = 0.6097 p = 0.65)in this agroecosystem. 174

175 **Proportion of the estrategia of parasitism.** The distribution was as **follows**, in general 176 the Koinobionte (specialist) represented 63% on average, while 36% on average is classified as Idiobiont (generalists) and only 1% that corresponds to the registror a species of the genus 177 Epsilogaster (Mendesellinae) of which its biology is unknown (Table 5). The percentage 178 179 from highest to lowest in the areas of koinobiont species was higher percentage was presented or in PT, followed by PC, VS and finally PL (Table 5). As for Idiobionts with 180 181 the highest and equal percentages PL and VS, followed by PC and finally PT (Table 5). This result suggests that in a diversified agroecosystem, species of braconide specialist 182 183 nests are more abundant, mainly in complex areas such as monocultures; and generalists 184 concentrate on areas of greater complexity such as secondary vegetation.

185 **Discussion**

Braconids are generally considered to be of great importance for their participation in the natural control of other insects and for their use in biological control programs for forest pests, fruit trees, vegetables and extensive crops worldwide (Coronado et al 2010). But the mechanisms of this possible regulation in management systems are little explored, particularly in agroecosystems that expect high levels of plant diversity and maintain high levels ofbiodiversity, with abiotic and abiotic changes that can affect insect communities (Klein et al 2002).

In the study of this diversified agroecosystem, it was found that the areas contain high levels of diversity of bracornests, with the presence of high percentages of specialists in the crops and communities of these organisms very similar to the remaining secondary vegetation. This supports the benefit of agroecosystems as conservation strategies, as some insects (e.g.

eumenid wasp and solitary bees) do not decline significantly to the modification of theenvironment (Klein et al 2002), which apparently happens in the case of bracornests.

According to the data obtained in this work it was observed that Microgastrinae was the most 199 200 abundant subfamily, which is expected. For this subfamily it has been estimated between 4000 to 10 000 species in he world (Joneset al 2009). The high abundance of microgastrinae 201 may be due to the fact that they are braconids with a wide range of hosts, which attack almost 202 203 all families of Lepidoptera, except the family Hepialidae (Shaw, 1994). And it includes 204 species of small size and with dark colors that possibly help them not to be seen by their 205 predators (Rathcke and Price, 1976; Hawkins et al 1992). The subfamilies Ichneutinae and Macrocentrinae were the least abundant, this is in line with the predicted for these subfamilies 206 207 which estimate there are between 100 -300 species worldwide (Jones et al 2009).

The study of braconids in Mexico is focused on three areas and they are, the knowledge of their taxonomic richness, which includes both faunal studies and descriptions of new taxa; research in ecology, mainly using these organisms as indicators of biodiversity; their use as biological control agents of other insects, with potential applications in agriculture and forestry activities (Coronado et al 2010; Coronado, 2011).

Recent studies have reported 318 genera of Braconidae for Mexico, 194 belonging to Yucatán (Coronado and Zaldívar, 2014). In this work, 47 genera were registered, which constitutes 24% of what was recorded for Yucatán. This family is well represented in the peninsula as it is the state with the highest number of genera recorded so far. It should be mentioned that the high number of genera found for this state is due to the fact that it is very close to the Gulf of Mexico and because it is one of the states where national and foreign

specialists work and /or where more collections have been made (Coronado and Zaldívar,
200 2014).

With the data obtained in this agroecosystem it was possible to observe that the areas of monocultures (e.g. PL), presented greater diversity of species, in terms of taxonomic richness and abundance of individuals. However, for parasitoid hymenoptera considered "specialised enemies" are not affected by the manipulation of plant diversity (Koricheva et al 2000; Abdala-Roberts et al 2016a). This was observed in three of the four areas where there were no significant differences in the abundance of bracor nests.

In this agroecosystem it was found that the largest number of individuals were classified as specialists (koinobionts) and that the presence of these is independent of the area of cultivation. One mechanism that could explain the patterns found in this agroecosystem may be More specialization (sensu Obermair et al 2008)which indicates specialization prevents competition, so productive communities support a greater number of species.

232 In this agroecosystem it remains to explore the role of the abundance of herbivores, which 233 has been pointed out in other cases, as a factor that suggests that the presence of herbivores 234 affects the number of parasitoids and the density of their population with respect to landscape, climate and management (Koricheva et al 2000; Schmidt et al 2003). This 235 236 indicates that the presence of herbivores in crops, caused by the concentration of a single 237 resource would affect the incidence of certain pests, which result in an important source for 238 the attraction of certain parasitoids, increased consumer abundance as proposed by the More 239 Individuals Hypothesis (Srivastava & Lawton 1998).

On the other hand, polyculture (VS) can present greater plant diversity than other sites,probably functioning as a remnant of natural habitat or as a refuge for parasitoids that disperse

242	to different	types of	of management	within	the	agroecosystem,	depending	closely	on	the
243	managemen	t (Thies	and Tscharntke,	, 1999) v	whic	h should be expl	ored.			

- 244 The results correspond to what was reported by Askew and Shaw (1986), Hawkins (1994),
- Ruiz-Guerra et al (2014), Rodríguez-Soliset al(2016) where they indicate that the braconid
- community is dominated by koinobiont parasitoids, mentions that the specialty of the host is
- not related to the degree of disturbance and therefore koinobiont parasitoids can occur both
- in the early stage and in the late succession phase of vegetation.
- Finally, the controls exerted by the plant diversity bottom-up and consumer top-down effects
- should be considered in future experimental work. Adding to this the context of the
- interactions that occur in agroecosystems must be studied with respect to the communities
- 252 (e.g. insects, animals) associated with the different trophic levels and the functional role they
- take in these systems.
- 254

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257 Declaration of interest statement

258 The authors declare that there is no conflict of interest

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- 381
- 382 Appendices
- 383 Appendix 1
- 384 List of subfamilies, morphospeies, abundances and biology of braconids in four sites:
- 385 PS=Parasitims strategy, K =Koinobionte, I = Idiobionte, ?= Unknow, PC = Coconut
- 386 Plantation, PL = Lemon Plantation, PT = P astizal, VS = Secondary Vegetation and TL =
- 387 Total *in the* agroecosystem.

subfamily	species	B PC	pl	Pt	VS	T1
Agathidinae	Alabagrus sp1	K 8	7	1	2	18
Agathidinae	Alabagrus sp2	K 0	2	0	0	2
Agathidinae	Bassus sp1	Κ 7	2	1	0	10
Agathidinae	Bassus sp2	K 4	0	0	0	4
Agathidinae	Bassus sp3	K 10	13	2	0	25
Agathidinae	Bassus sp4	K 3	0	1	0	4
Agathidinae	Pharpa sp1	K 0	0	0	2	2
Agathidinae	Plesiocoelus sp1	K 0	0	1	0	1
Alysiinae	Aphaereta sp1	K 3	7	1	3	14
Alysiinae	Asobara sp1	K 0	23	0	0	23
Alysiinae	Aspilota sp1	K 0	0	1	2	3
Alysiinae	Aspilota sp2	K 0	1	0	0	1

Alysiinae	Microcrasis sp1	K	0	2	0	0	2
Aphidiinae	Aphidiinae sp1	K	0	78	0	0	78
Blacinae	Blacus sp1	K	0	0	0	5	5
Blacinae	Blacus sp2	K	0	0	0	2	2
Braconinae	Bracon sp1	Ι	25	5	1	0	31
Braconinae	Bracon sp2	Ι	16	26	2	0	44
Braconinae	Bracon sp3	Ι	1	2	1	0	4
Braconinae	Compsobraconoides sp1	Ι	1	1	0	0	2
Braconinae	Habrobracon spl	Ι	0	3	1	0	4
Braconinae	Lapicida sp1?	Ι	0	2	0	3	5
Braconinae	Vipio sp1	Ι	1	0	2	0	3
Cardiochilinae	Schoenlandella sp1	K	0	17	0	0	17
Cardiochilinae	sp1	K	0	1	1	1	3
Cheloninae	Chelonus sp1	K	1	17	0	0	18
Cheloninae	Chelonus sp2	K	0	10	4	3	17
Cheloninae	Chelonus sp3	K	0	0	0	1	1
Cheloninae	Chelonus sp4	K	2	9	0	0	11
Cheloninae	Chelonus sp5	K	0	4	0	0	4
Cheloninae	Chelonus sp6	K	0	4	0	0	4
Cheloninae	Chelonus sp7	K	1	0	0	0	1
Cheloninae	Chelonus sp8	K	3	0	0	0	3
Cheloninae	Chelonus sp9	K	0	0	1	0	1
Cheloninae	Chelonus sp10	K	0	0	1	0	1
Cheloninae	Chelonus sp11	K	0	2	0	0	2
Cheloninae	Phanerotoma sp1	K	3	12	0	0	15
Cheloninae	Phanerotoma sp2	K	1	3	2	0	6
Cheloninae	Phanerotoma sp3	K	0	0	0	1	1
Cheloninae	Phanerotoma sp4	K	0	0	1	0	1
Cheloninae	Pseudophanerotoma sp1	K	0	2	0	0	2
Doryctinae	Acrophasmus spl	Ι	7	1	1	0	9
Doryctinae	Curtisella sp1	Ι	0	1	0	0	1

Doryctinae	Ecphylus sp1	Ι	0	4	0	0	4
Doryctinae	Ecphylus sp2	Ι	1	3	0	0	4
Doryctinae	Heterospilus sp1	Ι	0	3	0	1	4
Doryctinae	Heterospilus sp2	Ι	0	1	0	1	2
Doryctinae	Heterospilus sp3	Ι	0	17	4	4	25
Doryctinae	Heterospilus sp4	Ι	1	19	6	5	31
Doryctinae	Heterospilus sp5	Ι	1	8	2	1	12
Doryctinae	Heterospilus sp6	Ι	0	1	0	1	2
Doryctinae	Heterospilus sp7	Ι	0	3	0	0	3
Doryctinae	Heterospilus sp8	Ι	1	0	0	0	1
Doryctinae	Heterospilus sp9	Ι	3	0	1	1	5
Doryctinae	Heterospilus sp10	Ι	0	4	0	1	5
Doryctinae	Heterospilus sp11	Ι	0	1	0	0	1
Doryctinae	Heterospilus sp12	Ι	0	0	0	1	1
Doryctinae	Heterospilus sp13	Ι	0	0	0	2	2
Doryctinae	Heterospilus sp14	Ι	1	6	0	0	7
Doryctinae	Heterospilus sp15	Ι	0	6	0	2	8
Doryctinae	Heterospilus sp16	Ι	0	0	2	0	2
Doryctinae	Heterospilus sp17	Ι	0	0	4	1	5
Doryctinae	Heterospilus sp18	Ι	1	0	1	0	2
Doryctinae	Heterospilus sp19	Ι	0	1	0	0	1
Doryctinae	Heterospilus sp20	Ι	2	6	0	2	10
Doryctinae	Heterospilus sp21	Ι	0	1	0	0	1
Doryctinae	Heterospilus sp22	Ι	0	1	0	0	1
Doryctinae	Heterospilus sp23	Ι	0	1	0	0	1
Doryctinae	Odontobracon sp1	Ι	0	3	0	1	4
Doryctinae	Rhaconotus sp1	Ι	1	2	4	0	7
Doryctinae	Rhaconotus sp2	Ι	0	1	0	0	1
Doryctinae	Sp1	Ι	0	1	0	0	1
Euphorinae	Aridelus sp1	K	1	12	2	0	15
Euphorinae	Leiophron sp1	K	0	10	0	0	10

Euphorinae	Leiophron sp2	K 2	2	0	0	4
Euphorinae	Microctonus sp1	K 0	0	0	2	2
Euphorinae	Microctonus sp2	K 1	0	0	0	1
Euphorinae	Peristenus sp1	K 0	1	0	0	1
Homolobinae	Exasticolus sp1	K 0	1	1	0	2
Homolobinae	Homolobus sp1	K 0	5	0	0	5
Homolobinae	Homolobus sp2	K 0	0	1	0	1
Hormiinae	Cantharoctonus sp1	I 0	1	0	0	1
Hormiinae	Hormius sp1	I 1	1	1	0	3
Hormiinae	Hormius sp2	I 5	23	0	0	28
Hormiinae	Hormius sp3	I 0	1	0	1	2
Hormiinae	Hormius sp4	I 3	33	0	0	36
Hormiinae	Oncophanes sp1	I 0	1	0	0	1
Hormiinae	Oncophanes sp2	I 2	2	0	0	4
Hormiinae	Oncophanes sp3	I 1	5	0	1	7
Hormiinae	Oncophanes sp4	I 0	4	0	2	6
Hormiinae	Pambolus sp1	I 0	5	0	2	7
Hormiinae	Parahormius sp1	I 1	1	0	0	2
Hormiinae	Rhysipolis sp1	I 4	4	7	1	16
Hormiinae	Rhysipolis sp2	I 0	1	0	1	2
Hormiinae	Rhysipolis sp3	I 0	1	0	0	1
Hormiinae	Xenarcha sp1	I 0	0	0	3	3
Hormiinae	Xenarcha sp2	I 0	1	0	0	1
Ichneutinae	Oligoneurus sp1	K 0	1	0	0	1
Macrocentrinae	Macrocentrus sp1	K 1	0	0	0	1
Mendesellinae	Epsilogaster sp1	? 2	7	0	0	9
Meteorinae	Meteorus sp1	K 1	0	1	0	2
Microgastrinae	sp1	K 24	139	14	4	181
Microgastrinae	sp2	K 5	0	4	0	9
Microgastrinae	sp3	K 14	62	17	3	96
Microgastrinae	sp4	K 1	0	2	0	3

M	icrogastrinae	sp5	K	18	54	8	8	88
M	icrogastrinae	sp6	K	26	47	8	2	83
M	icrogastrinae	sp7	K	30	55	38	0	123
M	icrogastrinae	sp8	K	11	55	7	1	74
M	icrogastrinae	sp9	K	6	14	0	6	26
M	icrogastrinae	sp10	K	65	37	2	0	104
M	icrogastrinae	sp11	K	2	0	3	0	5
M	icrogastrinae	sp12	K	3	6	3	0	12
M	icrogastrinae	sp13	K	31	9	3	0	43
M	icrogastrinae	sp14	K	1	6	2	0	9
M	icrogastrinae	sp15	K	0	5	0	0	5
M	icrogastrinae	sp16	K	6	43	5	12	66
M	icrogastrinae	sp17	Κ	1	5	1	0	7
M	icrogastrinae	sp18	Κ	0	2	1	2	5
M	icrogastrinae	sp19	K	1	4	1	0	6
M	icrogastrinae	sp20	K	2	12	1	0	15
M	icrogastrinae	sp21	K	0	15	0	7	22
M	icrogastrinae	sp22	K	0	0	0	1	1
M	icrogastrinae	sp23	K	0	2	1	0	3
M	iracinae	Mirax sp1	K	4	17	0	6	27
Op	piinae	Opius sp1	K	30	81	2	27	140
Op	piinae	Opius sp2	K	0	3	0	0	3
Op	piinae	Opius sp3	K	0	2	0	0	2
Op	piinae	Opius sp4	K	4	4	0	0	8
Op	piinae	Opius sp5	K	1	10	1	8	20
Op	piinae	Opius sp6	K	11	37	2	3	53
Op	piinae	<i>Opius</i> sp7	K	1	6	1	1	9
Or	gilinae	Orgilus sp1	K	0	0	0	2	2
Or	gilinae	Orgilus sp2	K	4	0	0	0	4
Or	gilinae	Stantonia sp1	K	1	0	1	0	2
Or	gilinae	Stantonia sp2	K	0	4	3	2	9

Rogadinae	Aleiodes sp1	Κ	1	0	0	1	2
Rogadinae	Aleiodes sp2	Κ	0	4	0	0	4
Rogadinae	Stiropius sp1	Κ	9	12	0	0	21
Rogadinae	Stiropius sp2	Κ	0	7	0	0	7
			447	1232	194	158	2031

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391 Table(s) with caption(s) (on individual pages)

- Table 1. Number of genera (G), morphospecies (S) and abundance (A) by subfamilies in
- each area PC= Coconut Plantation, PL= Lemon Plantation, PT= Pastizal, VS=Secondary
- 394 Vegetation and the Total, in the agroecosystem.

subfamily	PC			pl			Pt			vs				
Number of	447	7		123	1232		194	194		158	3		total	
individuals														
	G	s	to	G	S	to	G	s	to	G	s	to	to	S
Agathidinae	2	5	32	2	4	24	3	5	6	2	2	4	66	8
Alysiinae	1	1	3	4	4	33	2	2	2	2	2	5	43	5
Aphidiinae	-	-	-	-	1	78	-	-	-	-	-	-	78	1
Blacinae	-	-	-	-	-	-	-	-	-	1	2	7	7	2
Braconinae	3	5	44	4	6	39	3	5	7	1	1	3	93	7
Cardiochilina	-	-	-	1	2	18	-	1	1	-	1	1	20	2
e														
Cheloninae	2	6	11	3	9	63	2	5	9	2	3	5	88	16

Doryctinae	4	1	19	6	24	95	3	9	25	2	1	24	163	31
		0									4			
Euphorinae	3	3	4	3	4	25	1	1	2	1	1	2	33	6
Homolobinae	-	-	-	2	2	6	2	2	2	-	-	-	8	3
Hormiinae	4	7	17	7	15	84	2	2	8	5	7	11	120	16
Ichneutinae	-	-	-	1	1	1	-	-	-	-	-	-	1	1
Macrocentrin	1	1	1	-	-	-	-	-	-	-	-	-	1	1
ae														
Mendesellinae	1	1	2	1	1	7	-	-	-	-	-	-	9	1
Meteorinae	1	1	1	-	-	-	1	1	1	-	-	-	2	1
Microgastrina	-	1	24	-	19	572	-	1	12	-	1	46	986	23
e		8	7					9	1		0			
Miracinae	1	1	4	1	1	17	-	-		1	1	6	27	1
Opiinae	1	5	47	1	7	143	1	4	6	1	4	39	35	7
Orgilinae	2	2	5	1	1	4	1	2	4	2	2	4	17	4
Rogadinae	2	2	10	2	3	23	-	-	-	1	1	1	34	4
total	2	6	44	3	10	123	2	5	19	2	5	15	203	14
	8	8	7	9	4	2	1	8	4	1	1	8	1	0

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399 Table 2. Shannon-Wiener diversity index, Total number of morphospecies, Singletons,

400 Doubletons, Unique species and rare species in PC coconut plantation areas; PL lemon

401 plantation; PT grassland and VS secondary vegetation in the agroecosystem.

		PC	pl	Pt	vs
	Shannon-Wiener index	3.43	3.77	3.41	3.45
	Total number of species	68	104	58	51
	Singletons	28	25	28	20
	Doubletons	7	13	12	14
	Unique species	35	31	32	30
	Rare species	35	38	40	34
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- 415 Table 3. Ice and Jacknife wealth estimators of the first order (Jacknife 1), with respect to the
- 416 morphospecies captured, calculated in the areas of PC coconut plantation; PL lemon
- 417 plantation; PT grassland and VS secondary vegetation in the agroecosystem.

	PC	PL	РТ		
MORPHOSPECIES	68	104	58	51	
CAPTURED					
ICE	120	129	103	94	
% CAPTURED FROM	57%	81%	56%	54%	
ICE					
JACKNIFE 1	97	130	85	76	
% CAPTURED FROM	70%	80%	68%	67%	
JACKNIFE 1					

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431 Table 4. Similarity of the braconid community in the areas of PC coconut plantation; PL

432 lemon plantation; PT grassland and VS secondary vegetation in the agroecosystem.

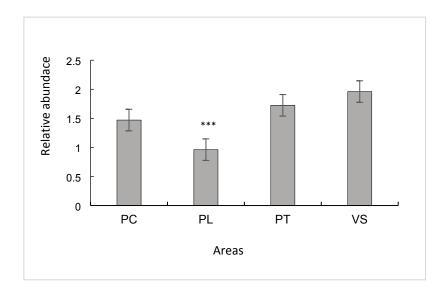
		PC	PL	РТ	VS
	РС	-	0.41	0.33	0.65
	PL	-	-	0.49	0.54
	РТ	-	-	-	0.58
	VS	-	-	-	-
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Table 5. Strategies of parasitism and morphospecies in which it is not known (?=Unknow)
expressed in total and percentage for PC coconut plantation; PL lemon plantation; PT
grassland and VS secondary vegetation in the agroecosystem.

		PC	%	PL	%	РТ	%	VS	%
	KOINOBIO	45	66	58	56	42	72	29	57
	NTS								
	IDIOBIONT	22	32	45	43	16	28	22	43
	S								
	?	1	2	1	1	-	-	-	-
	TOTAL	68	100	104	100	58	100	51	100
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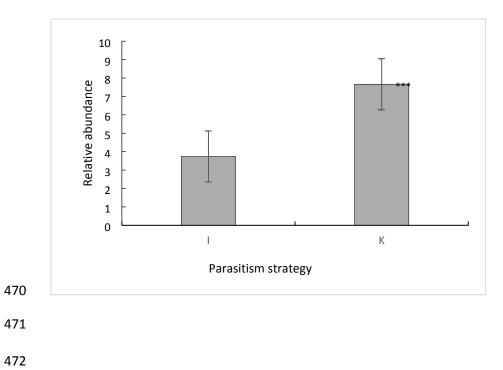


- 465
- 466 Figures
- 467 Figure 1





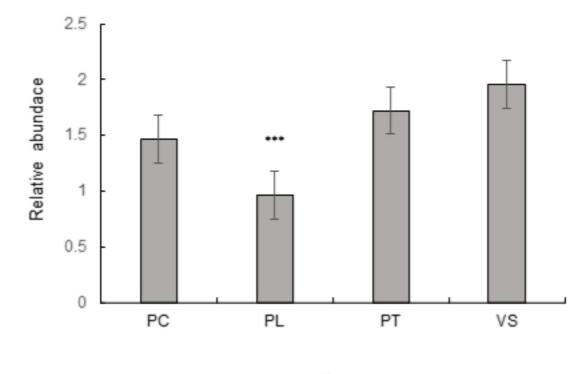






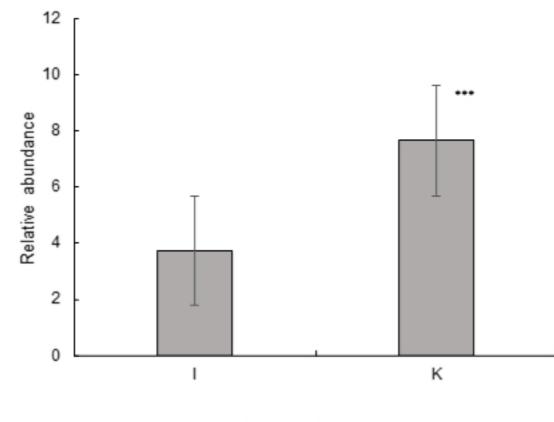
474 Figure captions (as a list).

- 475 Figure 1. Average relative abundance (EE±) of braconid morphospecies per pc coconut
- 476 plantation area; PL lemon plantation; PT grassland and VS secondary vegetation in the
- 477 agroecosystem (***p<0.05)
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- 479 Figure 2. Relative average abundance by parasitism strategy ($EE \pm$) of braconid
- 480 morphospecies per area PC coconut plantation; PL lemon plantation; PT grassland and VS
- 481 secondary vegetation in the agroecosystem (***p < 0.05)
- 482



Areas

Figure 1



Parasitism strategy

Figure 2