

REVIEW

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Oak canopy arthropod communities: which factors shape its structure?

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Abstract

Canopy of forest ecosystems has been recognized as a habitat that supports a wide variety of plants, vertebrates, invertebrates, and microbes. Within the invertebrate group, arthropods are characterized by their great abundance, diversity, and functional importance. Particularly in temperate forests, species of the genus *Quercus* (oaks) are one of the most important tree canopy groups, for its diversity and dominance. Different studies have shown that the oak canopy contains a high diversity of arthropods suggesting their importance as habitat for this group of organisms. In this review, we investigated the factors that determine the establishment, organization, and maintenance of arthropod communities in the oak canopy. In general, it was found that there is a lack of literature that addresses the study of oak canopy arthropod communities. Also, the following patterns were found: (a) the research has covered a wide variety of topics; however, there are differences in the depth to which each topic has been analyzed, (b) there are ambiguous criteria to define the structure of the canopy, (c) groups with different habitat preferences belonging to different guilds and uneven development stages have been studied, avoiding generalizations about patterns found, (d) the standardization in sampling techniques and collection has been difficult, (e) bias exists towards the study of phytophagous insects belonging to the Coleoptera, Hymenoptera, and Lepidoptera orders, and (f) there are few studies in other groups of arthropods, for example, acorn borers, whose activity has an impact on the fitness and dispersion of the host plants. Finally, we propose that the detection and study of patterns in oak canopy communities can be of great value to propose management and conservation strategies in these forests.

Keywords: *Quercus*; Temperate forests; Species diversity; Conservation

Introduction

The study of forest canopies has evolved from a new line of research into a fascinating discipline that has changed the way we perceive forest ecosystems (Nadkarni 2001; Nadkarni et al. 2004). Historically, studies in forest canopies were limited by access techniques and collection methods (Nadkarni 1994; Sutton 2001). The standardization of new access and collection methods started in the 1970s; from studies of only descriptive interest, researchers started to investigate the distribution, abundance and biodiversity of organisms, structure and functioning of the associated communities, and the dynamics of the ecosystems (Nadkarni and Parker 1994; Mitchell 2001; Stork et al. 2008). As access (i.e., balloons, cranes, platforms, bridges) and sampling methods (i.e.,

sprays, light traps, hand collecting) have evolved, canopy research became a viable option for researchers in many scientific disciplines, such as botany, zoology, landscape and ecosystem ecology, meteorology, conservation biology, etc. (Sutton 2001; Nadkarni et al. 2004; Stork 2007). This development promoted the diversification of research topics in the field, such as the study of community structure, biodiversity, choice of hosts, succession, and climate change (Lowman and Nadkarni 1995; Adis et al. 1998; Ozanne et al. 2003; Stork and Hammond 2013). Currently, canopy studies have matured, which is reflected in an increase in the number publications on cutting-edge topics. For example, the impact of global climate change and habitat disturbance on biodiversity and ecosystem dynamics is currently being assessed in the canopy of forests (Ozanne et al. 2003; Rinker and Lowman 2004; Lowman 2009) using a predictive approach.

The term “canopy” has been defined by different authors (Moffett 2000; Ozanne et al. 2003; Nadkarni et al. 2004);

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however, in this review, we use the definitions associated with forests. In general, the proposals made by different authors are integrated in the following definition: “it is the aerial part where all the foliage, buds, fine branches, associated flora and fauna, the suspended soil and the interstices (air) of a forest are combined.” Depending on the type of vegetation, the canopy can be defined vertically according to the different layers of vegetation between the ground and the canopy, which gradually modulates biotic (floristic composition, leaf area, biomass density, species diversity) and abiotic parameters (temperature, humidity, sunshine, wind velocity) (Basset et al. 2003a). This stratification creates a mosaic of resources and conditions that may be occupied by different organisms.

Several studies have documented that the canopy contains most of the photosynthetically active biomass in forest ecosystems (Wright and Van Schaik 1994; Ozanne et al. 2003) and is therefore the main energy absorption site in the biosphere (Basset et al. 2003b), as well as a site in which an intensive exchange of oxygen, water vapor, and carbon dioxide takes place (Hurtado et al. 2003). Furthermore, the canopy is a habitat that supports a wide variety of organisms, including plants, arboreal vertebrates, invertebrates, and microorganisms (Nadkarni et al. 2004), a discovery that has had a significant impact on the perception of biodiversity patterns at local, regional, and global levels (Erwin 1982; Novotny et al. 2002; Basset et al. 2003b). Since the early 1980s, the arthropod community associated with canopies is the one that has received the most attention. In particular, studies have focused on taxonomically describing the species associated with canopies, as well as the ecological processes that regulate populations, communities, and ecosystems (Southwood et al. 1982; Erwin 1995; Palacios-Vargas and Castaño-Meneses 2003; Tovar-Sánchez et al. 2013). In addition, the functional importance of arthropods in terrestrial ecosystems (pollinators, prey, parasites, parasitoids, herbivores, and detritivores) has stimulated the interest of researchers in the study of their communities (McIntyre et al. 2001). In general, the parameters that have been used to describe the structure of the canopy arthropod communities are the following: abundance, biomass, composition, richness, and diversity of species.

The results obtained by the works mentioned above indicate that the arthropod fauna associated with the canopy of temperate and tropical forests constitutes a very significant proportion of the global biodiversity (Erwin 1982; Novotny et al. 2002; Novotny and Basset 2005; Basset et al. 2007); the most recent estimate suggests 6.1 million species (Hamilton et al. 2013). In general, the diversity of arthropod species associated with the canopy is influenced by a latitudinal gradient (i.e., mites, collembola, beetles, and ants.), for example, there is greater diversity in tropical forests compared to

temperate forests (Gaston 2000; Nigel and Hughes 2004; Rinker and Lowman 2004; Novotny et al. 2006), a condition that could be a response to the increasing diversity of plants in the same direction (Novotny et al. 2006). The great diversity of plants and canopy arthropods that has been reported in tropical forests, coupled with their great exuberance, could explain the fact that, historically, the study of arthropod communities associated with the canopy has focused on these forests.

In particular, studies around the world have emphasized the importance of studying the arthropod fauna associated with the canopy of temperate forests, as well as the factors that determine the structure of their communities, since it has been shown that the canopy of these forests contains unique communities with a high diversity of species (Summerville et al. 2003a,b; Sobek et al. 2008; Tovar-Sánchez et al. 2013).

Oaks (Fagaceae: *Quercus*) are one of most important tree and shrub groups in temperate forests; they are widely distributed worldwide (Nixon 1993), have a great diversity of species (531, Govaerts and Frodin 1998), and are dominant components of the canopy (Challenger 1998). According to the latest taxonomic classification, the genus *Quercus* is divided into two subgenera, *Cyclobalanopsis* and *Quercus* (Nixon 1993; Manos and Stanford 2001). In particular, the subgenus *Quercus* includes four sections: *Cerris*, *Lobatae* (red oaks), *Protobalanus* (intermediate oaks), and *Quercus* (white oaks).

Most works that have addressed the study of oak canopy arthropod communities have included species belonging to the commonly called “red oaks” (*Lobatae*, i.e., Yarnes et al. 2008; Tovar-Sánchez 2009, Tovar-Sánchez et al. 2013; Tovar-Sánchez et al. 2015a,b) and “white oaks” (*Quercus*, i.e., Southwood et al. 2004, 2005; Yarnes et al. 2008; Tack et al. 2010). There are few studies in oaks belonging to other groups (Southwood et al. 2004, 2005; Nazemi et al. 2008). The works mentioned above included species belonging to different geographic regions (i.e., Africa (Moran and Southwood 1982; Southwood et al. 1982), America (Preszler and Boecklen 1994; Forkner et al. 2004; Yarnes et al. 2008; Tovar-Sánchez et al. 2013), Asia (Ishida et al. 2003; Ito and Ozaki 2005; Nakamura et al. 2008), and Europe (Southwood et al. 2004, 2005; Sobek et al. 2008)]. In general, these studies have revealed a great diversity of canopy arthropods (including epiphytes and suspended soil), a pattern that suggests that oaks are important habitats for these organisms.

Therefore, this paper aims to review the different factors that contribute to the establishment, organization, and maintenance of arthropod communities associated with the canopy of oaks. The following paragraphs describe and discuss the factors affecting this structure,

the patterns found, and the implications on the diversity and conservation of oak forests. The depth with which each topic is covered depends on the number of publications that support it.

Review

Genetic diversity of the host plant

Genetic diversity can be defined as any measurement that quantifies the magnitude of genetic variability at the individual, population, or species level (Nason 2002; Hughes et al. 2008). Genetic diversity has been considered the raw material for evolution by natural selection (Fisher 1930) and a fundamental source of biodiversity (Hughes et al. 2008). In general, its ecological consequences at the population level have been well studied (i.e., greater fitness of the population and lower risk of extinction, Vellend and Geber 2005). However, the role of genetic diversity in the organization and dynamics of communities is not yet clear (Johnson and Stinchcombe 2007; Hersch-Green et al. 2011; Wymore et al. 2011). It has been proposed that for genetic diversity to have a significant impact on a community, the latter must be dominated by one or a few foundation species (Bangert and Whitham 2007; Hughes et al. 2008). Foundation species are those that define and structure communities by creating locally stable conditions and providing resources for other species, besides participating in the modulation and stabilization of ecosystem processes (Ellison et al. 2005; Whitham et al. 2006). The trees of forest ecosystems are excellent candidates for foundation species because their architectural, functional, and physiological characteristics define the structure of the forests and can influence the microclimate, and their biomass and chemical constitution contribute significantly to ecosystem processes (Ellison et al. 2005). However, not only the trees of temperate zones have attributes of foundation species; tropical trees, seagrass, alpine herbs, coastal scrub, terrestrial grasses, and ferns have also been considered foundation species within their ecosystems (Whitham et al. 2012).

To date, most studies that have shown a genetic basis of variation in the phenotype of a community have been conducted in hybrid plants (i.e., *Eucalyptus*, Dungey et al. 2000; *Salix*, Hochwender and Fritz 2004; *Quercus*, Tovar-Sánchez and Oyama 2006b) and in specific genotypes within a species (i.e., *Oenothera*, Johnson and Agrawal 2005; *Populus*, Shuster et al. 2006, Schweitzer et al. 2008; *Solidago*, Crutsinger et al. 2008). In contrast, few studies have assessed a parameter of genetic diversity with respect to a parameter of community structure (i.e., *Populus*, Wimp et al. 2004; *Quercus*, Tovar-Sánchez and Oyama 2006 b, Tovar-Sánchez et al. 2013). One of the groups of herbivorous insects most commonly used in these studies are endophagous insects, probably due to their high

degree of specialization and their close relationship with the host plants (Tovar-Sánchez and Oyama 2006b). However, it is important to mention that communities of endophytic fungi, mycorrhizae, epiphytic and terrestrial plants, and soil microorganisms have also responded significantly to plant genetics (Whitham et al. 2012).

In particular, the effect of the genetic diversity of host species on arthropod communities associated with the canopy has been analyzed in terms of composition (Dungey et al. 2000; Bangert et al. 2005; Wimp et al. 2005; Bailey et al. 2006), richness (Dungey et al. 2000; Bangert et al. 2005, 2006, 2008), relative abundance (McIntyre and Whitham 2003; Wimp et al. 2005), and diversity of species (Wimp et al. 2004; Tovar-Sánchez and Oyama 2006b; Tovar-Sánchez et al. 2013). In general, the following patterns have been detected: (1) the diversity of arthropod communities is positively and significantly related to the genetic diversity of host species (Wimp et al. 2004; Tovar-Sánchez and Oyama 2006a; Tovar-Sánchez et al. 2013); (2) genetically more similar hosts support more similar associated arthropod communities (Bangert et al. 2006; Bangert and Whitham 2007; Kiers et al. 2010). In the first case, it has been proposed that increasing the genetic diversity of the host plant can generate changes in its morphological (Lambert et al. 1995; González-Rodríguez et al. 2004; Tovar-Sánchez and Oyama 2004), phenological (Hunter et al. 1997), architectural (Martinsen and Whitham 1994; Whitham et al. 1999; Bangert et al. 2005), and chemical characteristics (Fritz 1999; Cheng et al. 2011). These changes can be translated into a broader mosaic of resources and conditions that can be used by canopy arthropods. In the second case, it has been suggested that more genetically similar populations have a greater similarity in their physical, chemical, and phenological characteristics, favoring more similar arthropod communities associated with the canopy (Bangert and Whitham 2007).

Oaks are ideal species to study the effect of the genetic diversity of host plants on associated communities (Tovar-Sánchez et al. 2013; Tovar-Sánchez et al. 2015 a, b). This is because many oak species can be considered foundation species due to their wide geographic distribution (Valencia 2004) and dominance in the canopy of forests (Tovar-Sánchez et al. 2013) and because they are habitat to different species. They also show a high frequency of natural hybridization (Curtu et al. 2007; Peñaloza-Ramírez et al. 2010; Valencia-Cuevas et al. 2015), a condition that can promote an increase in genetic diversity (Tovar-Sánchez et al. 2008; Valencia-Cuevas et al. 2014) and the appearance of new features in host plants (Tovar-Sánchez and Oyama 2004) which can be exploited by arthropod communities. However, to date, there are few studies that have evaluated the structure of arthropod communities associated with the canopy of oaks from a genetic perspective.

Most studies that have evaluated the effect of the genetics of the host plant were conducted in hybrid systems (Preszler and Boecklen 1994; Yarnes et al. 2008). These studies have emphasized the response of arthropods, particularly phytophagous, to the variation found in these zones due to the generation of unique combinations of genetically based features of host plants, which could be associated with the oviposition preferences of insects and with the resistance responses of the plants (Boecklen and Spellenberg 1990, Aguilar and Boecklen 1992). However, most of these studies do not include an analysis to quantify the genetic diversity of the host plant and are based only on theoretical predictions that assume that the presence of two species, rather than one, increases genetic diversity of the host and that genetic diversity increases in hybrid individuals, especially when they are fertile, due to the huge number of genetic combinations that may occur (Whitham et al. 1999).

In general, there are few studies that have assessed a parameter of genetic diversity with respect to a parameter of the oak canopy arthropod community structure. Furthermore, these studies have yielded conflicting results. On one hand, we can mention the case of the complex of Mexican oaks formed by *Quercus crassipes* and *Q. crassifolia*; it was found that the composition, abundance, and diversity of the community of endophagous insects (gall-forming wasps and leaf miner moths) were affected by the genetic status of the host tree. Moreover, a significant positive relationship was found between the diversity of gall-forming insects and the genetic diversity of the host plant at the population level (Tovar-Sánchez and Oyama 2006b). Similarly, Tovar-Sánchez et al. (2013) found that the diversity and density of canopy arthropods associated to *Q. castanea* and *Q. crassipes* responded positively and significantly to the genetic diversity of the host tree at the individual level in central Mexico.

In contrast, Tack et al. (2010) found that genetic diversity had little influence on the community structure of endophagous insects (gall-forming, leaf miner and leaf roller insects) associated with the canopy of *Q. robur* in Finland. Similar results were reported by Castagneyrol et al. (2012), who found that the genetic attributes of the host plant (genetic diversity, kinship, and genetic identity) had no significant effect on the community of phytophagous insects (endophagous and ectophagous) associated with the canopy of *Q. robur* in France. The absence of a response by the canopy communities associated with *Q. robur* in Europe is probably due to the lower levels of genetic diversity of European species (whose populations were affected by glaciations) compared to the diversity reported for Mexican oak species (i.e., Magri et al. 2007; Marsico et al. 2009). Vakkari et al. (2006) reported very low levels of genetic diversity ($He = 0.162$) of *Q. robur* in Finland. The authors

suggest that these low values of genetic diversity may be explained by several factors: the high degree of fragmentation of the populations that this country represents the limits of the distribution of *Q. robur* and that *Q. petraea*, a species with which frequently exchange genetic material in central Europe, is not distributed in this country. Under this scenario, other factors may exert a greater influence on the structure of the associated communities. Bangert et al. (2006, 2008) proposed that if genetic variability does not increase on par with environmental variability, the latter will start to become a more important factor for the organization of the communities.

Because there are few studies that have evaluated the ecological consequences of the genetic diversity of oaks on their canopy arthropod communities and also because the results have been contradictory, it is essential to conduct further research to determine whether the effect of the genetic diversity of the host oak is the rule or the exception. Undoubtedly, the study of the genetic diversity of oaks and of its effect on the arthropod fauna associated with the canopy is a line of research with potentially important implications for conservation; previous studies in different species of poplar, eucalyptus, and willows (with attributes of foundation species, like oaks) suggest not only that genetic diversity influences the fitness of unique species but also that its benefits can extend to the associated arthropod communities.

Host plant hybridization

Interspecific hybridization is a common phenomenon in plants (Whittemore and Schaal 1991; Avise 1994; Rieseberg 1995; López-Caamal and Tovar-Sánchez 2014), especially common among species of the genus *Quercus* (Curtu et al. 2007; Peñaloza-Ramírez et al. 2010; Valencia-Cuevas et al. 2015). Particularly, for 20 years, the hybrid zones have been attractive as unique scenarios to study the effects of natural hybridization on plant-insect interactions (Fritz et al. 1994, 1998; Whitham et al. 1999; Wimp et al. 2004; Yarnes et al. 2008). Different studies have documented that the interspecific hybridization of plants can influence the distribution of herbivores between hybrids and putative parentals in the process of evolution of herbivores and in the coevolution process of insects and plants (Fritz et al. 1994; Fritz 1999; Dungey et al. 2000; Hochwender and Fritz 2004; Bangert et al. 2005). Community-level studies have found that arthropods show five patterns of response to the hybridization of their host plants: (1) *Susceptibility*: more species of insects in hybrid hosts than in parental species (Fritz et al. 1994; Whitham et al. 1994; Fritz 1999), (2) *Dominance*: hybrids support as many species of herbivores as one of the parental species (Fritz et al. 1994, 1996; Fritz 1999), (3) *Resistance*: hybrids support less herbivores than

both parental species (Boecklen and Spellenberg 1990; Fritz et al. 1994, 1996; Fritz 1999), (4) *Additivity*: hybrids support an intermediate number of insects compared to the parental species (Boecklen and Spellenberg 1990; Fritz et al. 1994, 1996; Fritz 1999) and, finally, (5) *No differences*: hybrids have an equal number of herbivorous insects than both parental species (Fritz 1999). The existence of these different patterns of response of arthropods in hybrid zones has been attributed to the age and extent of the geographical distribution of the hybrid zone, to environmental gradients, to the genetic status of hybrids, to the backcross direction, to the morphological and chemical similarity of the parental species, and to the genetic mechanisms that determine the inheritance of resistance mechanisms in hybrids (Boecklen and Spellenberg 1990; Fritz et al. 1994; Strauss 1994).

In the particular case of oak canopy arthropods, the previously described responses to the hybridization of the host plants have all been recorded. For example, the hypothesis of susceptibility has been reported in the case of the hybrids of the *Q. crassipes* × *Q. crassifolia* complex in Mexico, which supported the greatest richness and the highest number of rare species of ectophagous insects compared to parental individuals (Tovar-Sánchez and Oyama 2006a). Also, there are studies that have supported the hypothesis of dominance. For example, the *Q. dumosa* × *Q. engelmannii* complex in which hybrids showed a similar density of *Nepticula* spp., *Tischeria* spp., and *Coptodisca* spp. compared to pure individuals of *Q. gambelii* (Yarnes et al. 2008). Other studies have supported the hypothesis of resistance reporting a lower density and diversity of leaf mining insects (Lepidoptera: Nepticulidae and Gracillariidae) and gall-forming wasps (Hymenoptera: Cynipidae) in hybrid plants of the *Quercus depressipes* × *Q. rugosa* and *Q. emoryi* × *Q. coccolobifolia* complexes in northern Mexico (Boecklen and Spellenberg 1990), and a lower density of leaf miner moths (*Phyllonorycter*) in hybrid plants of the *Q. grisea* × *Q. gambelii* complex in New Mexico (Preszler and Boecklen 1994). The additive hypothesis has been supported by Aguilar and Boecklen (1992), who analyzed the density patterns of herbivores (leaf miner moths: Lepidoptera: Nepticulidae and Gracillariidae; gall-forming wasps: Hymenoptera: Cynipidae) in the *Q. grisea* × *Q. gambelii* complex in New Mexico and found that the hybrid hosts maintained intermediate densities compared to the parental hosts. Similar responses have been reported by Ishida et al. (2003), Tovar-Sánchez and Oyama (2006a, b) and Yarnes et al. (2008) in the *Q. crispula* × *Q. dentata* complex in Japan, *Q. crassipes* × *Q. crassifolia* in Mexico and *Q. grisea* × *Q. gambelii* in New Mexico, respectively. Finally, the hypothesis of no differences was demonstrated by the abundance of phytophagous

insects in the hybrids of the *Q. crispula* × *Q. dentata* complex in Japan (Ishida et al. 2003).

It has also been detected that the response of arthropods can be different in the same system. In the *Q. crispula* × *Q. dentata* complex in Japan, it was found that the abundance of phytophagous insects (leaf miner and leaf-chewing insects) responded differently to the hybridization of the host plant; the hybrids showed intermediate abundance (additivity), similar to one of the parental species (dominance) or similar to both parental species (no differences) (Ishida et al. 2003). The authors suggest that this differential response of herbivores resulted from differences in the expression of foliar chemistry in different genotypes of the host plant (F1, backcrosses) generated as a result of different patterns of introgression (unidirectional or bidirectional) in this system.

Hybrid bridges and change of host species

The mechanisms by which arthropod species can switch to a new host plant species are crucial to understanding the coevolution of plant-insect interactions (Floate and Whitham 1993). Currently, the change of host plant species has been explained by the hypothesis of the pre-adaptation or mutation required by arthropod species to “jump” and switch from one host to another (Floate and Whitham 1993). The pre-adaptation hypothesis suggests that herbivores are pre-adapted to switch to a new host species, but they do not do it because the new host is not present (Thomas et al. 1987). In contrast, when herbivores are not pre-adapted to the host, one or more key mutations must occur for the herbivore to be able to recognize a new and better host (Jermy 1984). Floate and Whitham (1993) proposed the hybrid bridge hypothesis, which predicts that hybrid plants facilitate the change of arthropods from one host species to another. For example, Keim et al. (1989) found that when a plant species presents allopatric distribution (with respect to a second potential host), this scenario acts as a barrier to the switching of hosts by the herbivore. The pre-adaptation hypothesis suggests that arthropods will not switch host species unless the hosts have a sympatric distribution (Thomas et al. 1987). But if two species hybridize so that the space between their distributions constitute “space bridges” formed by hybrid intermediaries, arthropods may switch to the new host species through the hybrids even though the parental plant species maintain an allopatric distribution.

This change of host can be affected by the hybridization patterns of plants, which determine the hybrid genotype that opens the genetic space between the parental species. Floate and Whitham (1993) proposed four hypothetical scenarios: *first*, no hybridization between the parental species; *second*, if parental species hybridize and produce

sterile F1 offspring, all hybrid individuals will have 50 % of the genome of each parental species; *third*, if the parental species show hybridization and produce sterile F1 offspring, the hybrids will show unidirectional introgression, i.e., the hybrids will cross with only one of the parental species to produce a genotype continuum of hybrids towards that parent. Under this scenario, the change of one host species for another is facilitated, but the space that remains still represents a barrier to the gradual change of host. The *fourth* scenario proposes that hybridization between parental species will produce fertile F1 hybrid individuals, which will backcross with the two parental species, generating a pattern of bidirectional introgression that will result in a genetic continuum between the hybrid individuals and the parental species, facilitating the change of host (Floate and Whitham 1993).

Considering the high frequency of hybridization between species of oaks (Spellenberg 1995; Howard et al. 1997; Tovar-Sánchez and Oyama 2004; Curtu et al. 2007; Peñaloza-Ramírez et al. 2010; Valencia-Cuevas et al. 2015) and that hybrid host plants facilitate the establishment of herbivore species (Whitham 1989; Boecklen and Spellenberg 1990; Floate and Whitham 1993; Tovar-Sánchez and Oyama 2006a), one would expect that the change of host by herbivores between species of the genus *Quercus* would be well documented. However, actually, this subject has been little studied. In a recent study, Tovar-Sánchez and Oyama (2006b) found that *Q. × dysophylla* (hybrid resulting from a cross between *Q. crassipes* and *Q. crassifolia*) has 25.7 % ($n = 9$) of the insects that are specific to the two parental species supporting the hypothesis of the hybrid bridge. A proposal associated with this hypothesis is that the hybridization patterns of plants (described above) can influence the structure of their communities. For example, it has been reported that species richness can be higher in hybrid zones with bidirectional introgression, where a genetic continuum can facilitate the accumulation in hybrids of herbivores of the two parental species (Whitham et al. 1999); intermediate species richness has been found in hybrid zones formed by unidirectional introgression where hybrids can accumulate herbivore species associated with only one of the parental species; the lowest species richness was found in hybrid zones formed by sterile F1 hybrids due to the lack of backcrossing with the parental species, which prevented the formation of a continuum of hybrid genotypes that could act as bridges for host switching (Whitham et al. 1999). In this regard, the study by Tovar-Sánchez and Oyama (2006b) in seven hybrid zones of the *Q. crassipes* × *Q. crassifolia* complex in Mexico showed that in the hybrid zone where bidirectional introgression was detected, the hybrids supported the highest richness of insects compared to

hybrids in the other six hybrid zones where there was unidirectional introgression (asymmetric).

Considering the high frequency of hybridization between species of the genus *Quercus*, the studies that have evaluated the effect of hybridization suggest that hybrid zones are centers of biodiversity of endophagous and ectophagous insects (Tovar-Sánchez and Oyama 2006a, b) and areas of great potential for exploring ecological and evolutionary processes at multiple levels (Strauss 1994). Therefore, the conservation of these zones may be crucial for the maintenance of arthropod communities associated with the canopy of the oak species involved in hybridization events.

Origin of the species (native or introduced)

The geographical distribution of plants can vary naturally due to geological or climatic changes or to anthropogenic causes (Strong et al. 1984), a situation that has allowed to compare the structure of arthropod communities in native and exotic habitats. For example, community-level studies have revealed that introduced plants support a lower total diversity of phytophagous arthropods and a change of herbivores from specialists to generalists compared to native plants (Colautti et al. 2004; Cripps et al. 2006; Hill and Kotanen 2010). This has been explained on the assumption that native herbivores do not find introduced plants to be suitable hosts because they prefer to feed on plants with which they share a common evolutionary history (Bernays and Graham 1988). Furthermore, the differences in the structure of guilds (generalists vs. specialists) between native and introduced plants are explained because specialists are more closely linked to the host plant; thus, they are expected to need more adaptations in order to be able to use a new host species (Gaston et al. 1992). Alternatively, other studies have shown that the abundance, richness, and foliar damage caused by arthropods is not different between native and introduced plants (Agrawal and Kotanen 2003; Frenzel and Brandl 2003; Agrawal et al. 2005), suggesting that some specialist herbivorous insects and native generalists can use the resources provided by introduced plants (Keane and Crawley 2002; Parker et al. 2006), especially when exotic plants coexist with closely related native plants, with whom they share similar mechanisms of defense (Tallamy 2004). In this case, natural enemies may not be associated in this new relationship, which may facilitate their establishment and an increase in herbivory levels. It has even been found that the abundance of herbivores may be greater in introduced plants than in native plants (Cripps et al. 2006), suggesting that native plants may be better adapted to the local herbivore fauna (Shea and Chesson 2002) or that introduced plants represent predator-free spaces, facilitating the

switch to a new host species (Keane and Crawley 2002; Murphy 2004).

There are few studies on the arthropod fauna associated with the canopy of oaks. However, the studies found reported a decrease in the abundance of arthropods in introduced host plants. For example, Southwood et al. (1982) found that the richness, diversity, and abundance of phytophagous insects associated with the canopy of *Q. robur* was lower in South Africa, where it is an introduced species, compared to Britain, where it is a native species. Similarly, Auerbach and Simberloff (1988) found a lower density of leaf miner specialists in *Q. acutissima* in Florida, where this species is an introduced one. More recently, Southwood et al. (2004) studied the fauna of phytophagous insects of two species of native oaks (*Q. robur* and *Q. petraea*) and two species introduced to Britain (*Q. cerris* and *Q. ilex*); they found that the two introduced species supported a lower density and richness of gall-forming insects and leaf miners compared to native species. With this same system, Southwood et al. (2005) found that, as in the case of the guild of herbivores, there was lower species richness and abundance of predators and parasitoids in introduced host species compared to native ones.

In general, there are very few studies that have evaluated the effect of the origin of the host oak species on the arthropod fauna associated with the canopy. However, assessing the evolution process of organisms that lack a history of coexistence or the response to the origin of the host species among arthropod specialists and generalists can provide very important information in terms of the conservation of native diversity and for designing management strategies for biological control.

Chemistry of the host plant

The chemical compounds of plants that can influence arthropod communities can be grouped into two categories: food and defense (Strong et al. 1984). In the first case, leaf nitrogen is a critical component for phytophagous insects (Strong et al. 1984), as there are records of a positive relationship between leaf nitrogen concentration and the rate of growth, reproduction, and survival of herbivorous insects (Mattson Jr. 1980). In addition, leaf nitrogen content can vary between species of plants and through foliar ontogeny (Jeffries et al. 2006; Ricklefs 2008), influencing the selection of the host plant by herbivores (Coley and Barone 1996). For example, a low nitrogen content has been associated with a reduced preference and performance of insects, as the palatability of the plant for herbivores depends on the ratio of carbon/nitrogen in the leaves (Schädler et al. 2003).

In the case of arthropod associated to oak species, several studies have supported the hypothesis proposed by Mattson Jr. (1980); these studies found a positive relationship between leaf nitrogen concentration in *Q. alba* (Wold and Marquis 1997), *Q. prinus*, *Q. rubra* (Forkner and Hunter 2000), *Q. dentata* (Nakamura et al. 2008), *Q. geminata*, *Q. laevis* (Cornelissen and Stiling 2006, 2008), *Q. alba*, *Q. coccinea*, and *Q. velutina* (Marquis and Lill 2010) and the density of herbivorous insects (i.e., leaf miners, leaf-chewing, gall-forming, and leaf rollers). Similarly, it has been reported that higher leaf nitrogen content in *Q. crispula* favors greater species richness of leaf-chewing insects (Lepidoptera) (Murakami et al. 2005, 2007, 2008).

In general, there are records of a significant effect of the individual plant species on the concentration of nitrogen and secondary metabolites (Laitinen et al. 2000; Osier et al. 2000; Cornelissen and Stiling 2008; Marquis and Lill 2010), suggesting that such variability affects the foraging activity and spatial distribution of arthropods. It has been suggested that the variation in the concentration of nitrogen and secondary metabolites depends on the following: (1) the genotype of the host plant (Glynn et al. 2004), (2) environmental conditions (Larsson et al. 1986; Henriksson et al. 2003; Niinemets and Kull 2003), and (3) the resources of the host plant (Ricklefs 2008). In addition, the chemical composition of the leaves of host plants may change as a result of damage by herbivores (plant's induced response), which has an effect on subsequent attacks (Karban and Myers 1989). This induced response has also been documented in oaks. For example, Wold and Marquis (1997) found a decreased nitrogen content in seedlings of *Q. alba* that had previously been subjected to herbivory in an experimental field in Missouri. This change in foliar chemical composition favored a reduction in the subsequent damage caused by herbivore insect compared to the seedlings that had initially shown lower herbivory.

Furthermore, tannins have been the most studied chemical components of oaks regarding their effect on the structure of arthropod communities (Feeny 1970; Abrahamson et al. 2003; Forkner et al. 2004; Murakami et al. 2005, 2007, 2008; Yarnes et al. 2008). In a pioneering work, Feeny (1970) documented a significant and negative relationship between the concentration of leaf tannins in *Q. robur* and the abundance of phytophagous insects (*Operophtera brumata*), suggesting a defensive role for these compounds. Specifically, tannins have been reported to reduce the growth and survival of phytophagous insects (Kause et al. 1999; Kopper et al. 2002), to produce lethal deformities (Barbenhenn and Martin 1994) and to increase rates of parasitism (Faeth and Bultman 1986). It has also been documented that there is a negative relationship

between the concentration of tannins and the biomass of Lepidoptera in the pupal stage (Lill and Marquis 2001).

Other studies in oaks found the same results as Feeny (1970); they reported a negative relationship between the concentration of tannins in the plant and the abundance and richness of herbivores (leaf-chewing insects, leaf miners, and gall-forming) that inhabit the canopy of *Q. alba*, *Q. velutina* (Le Corff and Marquis 1999; Forkner et al. 2004), *Q. crispula* (Murakami et al. 2005, 2007, 2008), *Q. geminata*, *Q. laevis* (Cornelissen and Stiling 2006, 2008), and *Q. gambelii* × *Q. grisea* (Yarnes et al. 2008).

It has also been well documented that plants produce other chemical substances (i.e., oxalic acid, alkaloids, phenolic content, toxic lipids, flavonoids, and lignins) that act as defenses or insect attractants (Becerra et al. 2001; Salminen et al. 2004). In response to these signals or chemical defenses, arthropod communities can have different structures (Inoue 2003). In particular, it has been documented that gall-forming insects (Cynipidae) are a group that responds sensitively to differences in foliar chemistry between oak host species. For example, Abrahamson et al. (1998, 2003) found that the community composition of gall-forming wasps in six species of oak (*Q. laevis*, *Q. myrtifolia*, *Q. inopina*, *Q. chapmanii*, *Q. geminata*, and *Q. minima*) in the east of the USA was different and unique for each species. Similar results were reported for the *Q. crassipes* × *Q. crassifolia* complex in Mexico (Tovar-Sánchez and Oyama 2006b), for *Q. infectoria* and *Q. brantii* (Nazemi et al. 2008) in Iran and for *Q. castanea* and *Q. crassipes* (Tovar-Sánchez et al. 2013) in Mexico. The authors suggest that this close relationship is a result of the high degree of specialization of insects to host plant chemicals.

Moreover, oaks have been found to show seasonal variations in foliar chemistry. Some examples have been documented in *Q. robur* (Feeny 1970; Salminen et al. 2004), *Q. alba*, *Q. velutina* (Le Corff and Marquis 1999), *Q. alba* (Lill and Marquis 2001), *Q. crispula* (Murakami et al. 2005, 2007, 2008), *Q. geminata*, and *Q. laevis* (Cornelissen and Stiling 2006, 2008). These studies found a temporal variation in the nutritional quality of the leaves, as the ontogenetic development of leaves progressed, the content of tannins and lignins increased and the content of water and nitrogen decreased (Feeny 1970; Lill and Marquis 2001; Forkner et al. 2004; Murakami et al. 2005). Several studies have shown that this seasonal variation in foliar chemistry is reflected in the structure of the arthropod community associated with the canopy of oaks in terms of composition, species richness, diversity, biomass, and abundance (Southwood et al. 2004, 2005; Murakami et al. 2005, 2007; Tovar-Sánchez 2009; Tovar-Sánchez et al. 2013). In general, the pattern that has been observed is that the richness, diversity,

abundance, and biomass of canopy arthropods decrease as the season progresses, while the composition varies in response to changes in the chemistry of the host plant.

Considering the great diversity of oak species worldwide, and the great diversity of canopy arthropods, the study of the chemistry of the host and of its influence on plant-insect interactions provides an ideal scenario to study the oviposition preferences, food, ontogenetic performance, host switching, and community structure of arthropods in response to variations in the chemical defenses of the host plant.

Spatial variation

One of the most important patterns of the communities of plants and animals is that organisms are not spread evenly in nature (Agrawal et al. 2006). A consequence of this phenomenon is that the identity and intensity of the interactions between plant and insect species shows a high spatial variability (Agrawal et al. 2006). For example, if the composition and diversity of plant communities vary predictably across habitats and biogeographical zones (Gurevitch et al. 2002), it is reasonable to assume that the strength of plant-herbivore interactions can also vary. For example, a latitudinal gradient in the intensity of herbivory is predicted at the global scale, which has led to a greater diversity of unpalatable plants in the tropics compared to temperate regions (Pennings and Silliman 2005). In addition, plant communities show large spatial differences in richness, genetic diversity, abundance, and total biomass, helping create heterogeneous habitats that promote differences in arthropod communities. In forests, horizontal (between localities, Gering and Crist 2000; Summerville et al. 2003a, b) and vertical spatial variation (between layers) can have an influence on canopy insect communities (Le Corff and Marquis 1999; Simmon and Linsenmair 2001; Forkner et al. 2004; Barber and Marquis 2009). The horizontal differences can be generated by changes in biotic (i.e., dominant canopy species, composition, abundance, and diversity of host species, forest age) and abiotic factors (i.e., geological age, topography, soil type, altitude) between localities (Gering et al. 2003; Price et al. 2004). On the other hand, vertical variation results from the presence of different layers of vegetation in the forest, which gradually modulates biotic (i.e., floristic composition, leaf area, biomass density, species diversity) and abiotic parameters (i.e., temperature, wind velocity, sunshine, humidity) along a vertical gradient running from the forest floor to the canopy (Basset et al. 2003a). This stratification of vegetation results in a great diversity of microhabitats, which can support different communities of arthropods (Parker 1995; Forkner et al. 2004; Barber and Marquis 2009). However, there has been no consensus among researchers to define the canopy vertically in

temperate forests, as there are some who recognize two layers, overstory (canopy and aerial part), and understory (shrubs and herbaceous plants, Le Corff and Marquis 1999; Lewinsohn et al. 2005; Murakami et al. 2005) and some who divide the canopy into an upper and a lower layer (Wagner et al. 1995; Simmon and Linsenmair 2001). In general, the spatial variation of resources and conditions between locations and layers generates changes in the composition, richness, abundance, and diversity of arthropod communities associated with canopies (Strong et al. 1984; Gering et al. 2003). On the other hand, the studies that have been conducted on the vertical structure of arthropod communities have not found consistent patterns of stratification. In some cases, species are more abundant in the lower canopy (Basset 1991); in other cases, they are more abundant in the upper canopy (Meagher and Hull 1987), while in other cases, the species are evenly distributed (Brown et al. 1997).

In the particular case of the arthropod fauna associated with the canopy of oaks, it has been found that parameters such as composition, richness, and diversity respond to the spatial variation between localities and layers. In the first case, Gering et al. (2003) found that the communities of coleoptera associated with the canopy of *Q. alba* and *Q. rubra* were different in terms of composition, richness, and diversity between sites and ecoregions, which they attributed to the heterogeneity of resources and conditions in the different scales studied. Similarly, Tovar-Sánchez et al. (2003) found that the composition, richness, and diversity of the arthropod community associated with the canopy of *Q. castanea*, *Q. crassipes*, *Q. crassifolia*, *Q. greggii*, *Q. laeta*, and *Q. rugosa* in three locations in central Mexico showed significant differences in response to different degrees of disturbance and fragmentation between locations. Similarly, Price et al. (2004) and Barber and Marquis (2009) reported differences between locations in the species richness of gall-forming insects (Cynipidae) associated with the canopy of *Q. myrtifolia* and in the abundance of herbivorous and predatory canopy arthropods associated to *Q. alba*.

In particular, the spatial variation in the physical and chemical characteristics of the leaves within the canopy of oaks was documented (Roslin et al. 2006), and its effect on the arthropod communities associated was evaluated. However, the studies that have examined the stratification of arthropod communities in the canopy of oaks have shown contrasting results. On one hand, some studies have reported that the upper canopy supports the highest density and abundance of arthropods; such is the case of the study by Simmon and Linsenmair (2001), who found that the density of herbivorous insects associated with the canopy of *Q. subsericea* in a rainforest in

Borneo, Malaysia, was greater in the upper canopy compared to the lower canopy, suggesting that in these microenvironments, differences in temperature or in foliar chemistry, and palatability of leaves between layers could be responsible for differences in the density and abundance of the associated arthropod species. In addition, this study found differences in the composition and abundance of the arthropod community between layers; in the lower layer, homoptera and ants were the dominant groups, while at in the upper layer, the dominant groups were diptera and hymenoptera (except ants). On the other hand, there are studies that have found that the richness and abundance of oak canopy arthropods are lower in the canopy than in the understory. For example, Le Corff and Marquis (1999) found that the richness of herbivorous insect species (larvae of Lepidoptera) in *Q. alba* and *Q. velutina* was higher in the understory than in the canopy, suggesting that environmental factors such as the amount of light can be important in explaining the results. In addition, Nakamura et al. (2008) found that *Q. crispula* showed a greater abundance of herbivorous leaf miners in the canopy compared to the understory, suggesting that the results were determined by the higher leaf nutritional quality in the understory compared to the canopy.

A higher species richness was also been found in the inside of the canopy of oaks. For example, Sobek et al. (2008) found that the species richness of mites (Acari: Oribatida) associated with the canopy of *Q. robur* was higher in the inside of the canopy, suggesting that mites are able to live in the thick branches near the trunk, while they become rare in the outer branches, which are constantly exposed to rain, wind, and solar radiation, making these sites unfavorable habitats for many arthropods.

In addition, there are studies that have found that vertical stratification in arthropod communities associated with the canopy of oaks changes seasonally in response to changes in the nutritional quality of the leaves between layers. Examples of this are the patterns of vertical stratification in the community of leaf-chewing insects associated with the canopy and understory of *Q. alba* and *Q. velutina* (Forkner et al. 2004) and in the community of lepidopteran larvae in the canopy and understory of *Q. crispula* (Murakami et al. 2005).

The influence of the spatial variation of the host plant on the arthropod fauna associated with oak species is probably due to the lack of consensus about the vertical structure of the canopy in temperate forests, which leads to comparing arthropod groups with different habitat preferences belonging to different guilds and in unequal stages of development. A control of the above variables could identify clear response patterns of canopy arthropods to the vertical spatial variation in oak forests.

Seasonal variation

Temperate deciduous forests show predictable changes in highly synchronized phenological events, as most species of trees show sprouts of leaves in spring, followed by senescence and eventual leaf fall in autumn (Strong et al. 1984; Murakami et al. 2007). Consequently, these seasonal cyclical changes change the structure of the arthropod community associated with the canopy. The change in the composition, as well as the decrease in relative abundance and richness of canopy arthropod species in temperate forests as seasonality advances is well documented (Gering et al. 2003; Southwood et al. 2004, 2005; Murakami et al. 2005, 2007). In this regard, it has been suggested that in temperate forests, the nutritional quality of the leaves decreases as the season progresses (Feeny 1970); the leaves become harder, its water and nitrogen content decreases, and the content of tannins and fibers increases (Feeny 1970; Le Corff and Marquis 1999; Lill and Marquis 2001; Salminen et al. 2004; Cornelissen and Stiling 2008; Murakami et al. 2007, 2008). These foliar changes have also been used to explain the differences in the composition of arthropod communities; as the season progresses, species with different feeding preferences appear (i.e., leaf-chewing insects at the start of the season and sap-sucking insects at the end, Strong et al. 1984; Southwood et al. 2004; Stork and Hammond 2013).

In addition, temperate forests exhibit a seasonal rainfall pattern in which the rains are distributed throughout a season that lasts 6–7 months and is interrupted by a dry season that can last from 5–6 months (Rzedowski 1978). This seasonal variation has several effects on the phenology of the plants and on their associated arthropod communities; in the rainy season, there is an increase in the formation of branches, leaves and fruits, as well as in the development of epiphytic plants, and this favors the production of a range of resources and conditions that can be used by canopy arthropods. This can promote changes on a microclimate level (Basset and Novotny 1999; Peeters et al. 2001), as well as an increase in environmental heterogeneity, which can be used by arthropods (Yarnes and Boecklen 2005). In addition, during the rainy season, the young leaves are the most abundant and nutritious, they are less hard and have less chemical defenses (Kursar and Coley 2003; Boege 2004; Coley et al. 2006). Finally, the increase of vegetation in the forests (i.e., annual vegetation) during the rainy season can serve as a bridge so that new species of arthropods have a greater chance of colonizing new trees (Basset et al. 1992; Campos et al. 2006).

Studies in oaks have documented the effect of host plant phenology and seasonality on the canopy arthropod fauna. In the first case, a decrease in the density,

richness, diversity, and biomass of insects associated with the canopy of oaks has been detected as the season progresses. Examples include the community of leaf-chewing insects associated with the canopy of *Q. alba* and *Q. velutina* in Missouri (Forkner et al. 2004), the community of phytophagous insects associated with the canopy of *Q. cerris*, *Q. ilex*, *Q. petraea*, and *Q. robur* in a forest in France (Southwood et al. 2004, 2005), and the community of beetles associated with the canopy of *Quercus* spp. in two forests in Turkey (Şen and Gök 2009).

Furthermore, it has been found that during the rainy season the canopy of oaks supports greater species diversity, richness, density, and biomass compared to the dry season. Examples of this have been reported in the community of collembola inhabiting *Tillandsia* spp., an epiphytic plant of the canopy of *Quercus* spp. in a temperate forest in central Mexico and in the community of ectophagous insects associated with the canopy of *Q. laurina* and *Q. rugosa* (Tovar-Sánchez 2009) and of *Q. castanea* and *Q. crassipes* (Tovar-Sánchez et al. 2013) in a temperate forest in Central Mexico.

The high sensitivity of arthropod species to changing environmental parameters can be a tool to predict the effect of climate change on biodiversity in oak forests, using canopy arthropods as bioindicators.

Habitat heterogeneity

The hypothesis of habitat heterogeneity is one of the cornerstones of ecological studies; it predicts that structurally more complex habitats provide a wider range of resources and conditions, which favors the establishment of a greater diversity of species (Halaj et al. 2000, Tews et al. 2004; Affeld 2008). Because plant communities determine the physical structure of the environment, they have great influence on the structure of animal communities (Strong et al. 1984). For example, Martinsen and Whitham (1994) found that the hybrid plants of the *Populus angustifolia* × *P. fremontii* complex had a more complex architecture compared to the putative parental species. This favored the presence of a greater number of bird nests in hybrid plants and, therefore, an increase in the richness of associated bird species.

In the case of arthropods inhabiting the canopy of trees, the structure and complexity of their habitat is determined by the abundance and architecture of the plants (i.e., shape and size of leaves, buds, twigs, and epiphytes as well as the bark texture, Halaj et al. 2000; Affeld 2008; Ulyshen and Hanula 2009).

In addition, important resources for herbivores such as food, shelter, foraging sites, oviposition, and sexual deployment are directly related to habitat heterogeneity (Halaj et al. 2000). For example, it has been documented

that the structural complexity of the canopy is increased by the presence of epiphytic plants (i.e., orchids, bromeliads, ferns, mosses, and lichens), since they differ significantly in structure, growth habit, and function, providing a great diversity of microhabitats and resources (Ishii et al. 2004; Affeld 2008) that can be used by arthropods.

Moreover, the forest canopy can be structurally more complex when it is formed by more than one species (Ishii et al. 2004; Novotny et al. 2006; Sobek et al. 2009), resulting in a more complex environment due to the differences in plant architecture and the increased availability of resources and conditions (Novotny et al. 2006). This hypothesis is supported by studies that have shown that some groups of birds (Martinsen and Whitham 1994; Poulsen 2002), mammals (Southwell et al. 1999; Williams et al. 2002), and amphibians (Atauri and Lucio 2001) respond positively to the physiognomic complexity of host plants and, furthermore, by studies that have shown a positive effect of taxonomic diversity of host plant species on the associated communities (Siemann et al. 1998; Beals 2006).

In the case of arthropods, it has also been reported that more complex habitats offer a wider range of niches that can support a larger number of species (Humphrey et al. 1999; Hansen 2000; Hamer et al. 2003; Lassau and Hochuli 2005). Specifically, a positive relationship between species richness of arthropods and plant diversity has been reported (Gaston 1991; Siemann et al. 1998; Novotny et al. 2006; Sobek et al. 2009).

In addition, several authors have suggested that the structural complexity of the forest increases with age, i.e., mature forests are structurally more complex compared to young forests or plantations (Schowalter 1995; Evans and Jukes 2000; Jeffries et al. 2006), as the former have larger trees in terms of height and biomass. Moreover, mature forests contain trees of different ages, which favor greater structural complexity (Ishii et al. 2004).

Several studies in oaks have documented that structurally more complex habitats offer more resources and conditions to canopy arthropods, which has a positive effect on the richness, diversity, and abundance of their communities. For example, Marquis et al. (2002) demonstrated the effect of the architecture of *Q. alba* on the abundance of shelter-building caterpillars. Their results showed a positive relationship between the structural complexity of the canopy of the mentioned species (percentage of overlapping leaves) and abundance of caterpillars (Lepidoptera), suggesting the importance of plant architecture for these herbivores. Subsequently, Lill and Marquis (2003) found, in the same system, that the building of shelters by these caterpillars favors a greater structural complexity in *Q. alba*, resulting in

an increase in the species richness of leaf-chewing insects.

Other studies have shown the positive effect of structural habitat complexity related to forest age on the richness of the arthropod fauna associated with the canopy. An example of this is the community of leaf-chewing insects associated with the canopy of *Q. alba* and *Q. velutina* (Marquis and Le Corff 1997; Marquis et al. 2000) and the lepidopteran community associated with the canopy of *Quercus* spp. (Summerville and Crist 2002, 2003).

Recognizing the positive effect of habitat heterogeneity on the arthropod communities associated with the canopy could be useful to propose management strategies that promote the structural complexity of the canopy of oak forests and thereby benefit the associated arthropod communities.

Biotic interactions

A community is a group of species that share the same habitat, in which, at least some of the component species populations will interact with each other. These interactions and the resulting population dynamics underlie broad patterns in the structure of ecological communities (Strong et al. 1984). For example, in arthropod communities the interactions can broadly affect the local abundance and distribution of specialist and generalist predators and parasitoids, as well as herbivores in several feeding guilds (Styrsky and Eubanks 2007; Tack et al. 2012). In general, the interactions between different species inside the communities can be established “horizontally,” when the species are potential or actual competitors or “vertically,” when the interactions involving natural enemies (Morin 1999).

In particular, interspecific competition is any mutually negative interaction between two or more species within the same guild or trophic level (Polis and Holt 1992). Competitive interactions manifest themselves as reduced abundance, decreased fitness, or a decrease in some fitness component, such as body size, growth rate, fecundity, or survivorship (Begon et al. 2005). The assumption is that decreases in fitness components would eventually cause the reduction in the abundance of species involved, a condition that could change the community structure (Hooper et al. 2000). Competition between species for limited resources has been regarded as a process that structures ecological communities, because it limits the number of coexisting species, molding and constraining what those species do, where and how they feed, their body sizes, seasonal distribution, etc. (Strong et al. 1984; Begon et al. 2005).

For insects, the role of intra- and interspecific competition in ecological theory has changed throughout the years from the notion that competition was weak and

infrequent on phytophagous insect communities in the early 1980s (Lawton 1982; Strong et al. 1984), to the resurrection of the importance of competition among phytophagous insects in the 1990s (Denno et al. 1995; Reitz and Trumble 2002). Most notably, Kaplan and Denno (2007) reviewed the evidence for competition in phytophagous insects from 145 published studies and found signal for interspecific competition in 73 % of the individual cases compared, although the magnitude of effects varied among the variables analyzed.

On the other hand, the ecology theory considers that phytophagous insects are kept rare, relative to the availability of potentially limiting resources, by the impact of natural enemies as insect parasitoids, insect predators, birds, pathogens, etc. (Strong et al. 1984). Hence, it has been suggested that the major processes acting in many phytophagous insect communities work vertically through the food chain, not horizontally with others species in the same trophic level (Morin 1999).

Predation is defined as the interaction between an individual predator and a prey, where the predator benefits from the interaction, while the consumed prey does not (Rosenheim et al. 1993). Predator-prey interactions involve species that reside on many different trophic levels, including the impacts of herbivores on plants, carnivores on herbivores, carnivores on other carnivores, and parasites and parasitoids on hosts. Predators affect community composition in diverse ways. Some predators feed selectively on competitively superior species that would otherwise exclude weaker competitors. This fact enhances the number of prey species that are able to coexist, since predators reduce the interspecific competition among surviving prey species (Murakami and Nakano 2000). For the above mentioned, several studies have proposed predation as a potent agent of natural selection, population regulation, and community structure (Marquis and Whelan 1994; Murakami and Nakano 2000), being considered as one of the most important balancer mechanism of the natural ecosystems.

In contrast, minor importance has been given to positive interactions among species (Morin 1999). The tendency to overlook a positive effect of one species on another neglects the potential importance of some of the more fascinating interspecific interactions that can occur in communities (Morin 1999). This oversight is unfortunate given that mutualisms, while often inconspicuous, are common and potentially important forces that influence the structure and function of communities (Bronstein 1994; Connor 1995; Kiers et al. 2010). Mutualism, defined as a reciprocally beneficial interaction between individuals of two species, is increasingly recognized as a common and important ecological interaction (Bronstein 1994; Stachowicz 2001). Janzen (1985) identified four key types of mutualisms: dispersal,

pollination, nutritional, and protective. For example, community-level effects of nutritional mutualisms, such as mycorrhizal association formed between fungi and the roots of many higher plants can influence seedling establishment and the outcome of competition. Likewise, many higher plants are involved in a facultative mutualism with arthropods and vertebrates that pollinate their flowers and disperse their seeds (Bertness and Callaway 1994). While these positive interactions are often emphasized by ecologists who study various forms of plant—animal interactions, their impact on community organization remains little explored (Bertness and Callaway 1994; Morin 1999; Styrsky and Eubanks 2007).

Another example of positive interactions among species is the existence of ecosystem engineers. Despite controversies about the appropriate use of the term “ecosystem engineering” (Jones and Gutiérrez 2007), it is currently a well-recognized type of positive ecological interaction (Hastings et al. 2007; Burchsted et al. 2010). Ecosystem engineers are organisms that directly or indirectly control the availability of resources for other organisms by causing physical state changes in biotic or abiotic materials (Jones et al. 1997) influencing local patterns of biological diversity. In this sense, Romero et al. (2014) conducted a meta-analysis of 122 studies which explored effects of animal ecosystem engineers on species richness of other organisms in the community. The analysis revealed that the overall effect of ecosystem engineers on diversity is positive and corresponds to a 25 % increase in species richness, indicating that ecosystem engineering is a facilitative process globally.

Specifically, this review revealed that the interaction among species is a factor that has impact on the abundance, distribution, and diversity of arthropods associated to oak canopies. In particular, we found that parameters such as composition, abundance, and species richness respond to the interactions between species as depredation, competition, and mutualism. For example, it has been reported that arthropod communities associated to oak canopies respond to predation by natural enemies as birds (Gunnarsson and Hake 1999; Murakami and Nakano 2000; Barber and Marquis 2009; Böhm et al. 2011; Schönrogge et al. 2013; Ceia and Ramos 2014), bats (Böhm et al. 2011), parasites, and parasitoids (Sanchez et al. 2013), in terms of abundance. However, most the studies have evaluated the influence of birds as regulators of species populations that integrate the arthropod communities. In this context, the study of Murakami and Nakano (2000) found that predation by two bird species (*Parus major* and *Sitta europaea*) reduced the population sizes of herbivorous insects associated to *Q. crispula* in Japan. These results highlight the function of birds as abundance regulators of the herbivores insects in canopy forest

communities and its role in the community structure. Similarly, Ceia and Ramos (2014) revealed the importance of several bird species as potential regulators of the abundance and distribution of defoliator insects associated to *Q. robur* and *Q. rotundifolia* canopy in the Mediterranean. In contrast, Lichtenberg and Lichtenberg (2002) studying the interactions among insectivorous birds, arthropods, and white oak saplings (*Quercus alba*) in a temperate deciduous forest did not find a detectable influence of birds on leaf-chewer densities in either year.

On the other hand, Nakamura et al. (2008) analyzed the population dynamics of several leaf miners on a deciduous oak *Q. dentata* for 9 years in northern Japan. The authors found that density-dependent effects were not explicit in the population dynamics of the species studied. The authors suggested that interspecific competition is not a factor that modeled the community structure in this system, because leaf miners species are often associated with diverse parasitoid species and they are sometimes subjected to the top-down population regulation by these enemies. In order to understand whether parasitoids mediate interactions of the two host leaf miner species (*Phyllonorycter permilis* and *P. leucocorona*), Nakamura and Kimura (2009) studied the parasitism rate in this system. The authors reported a high parasitism rate (24.1–92.6 % in *P. persimilis* and 58.9–81.7 % in *P. leucocorona*) suggesting that parasitism was a major mortality factor in these *Phyllonorycter* species and a promotor of both species coexistence. Though the parasitoid composition was different between the two hosts species, most parasitoids were able to parasitize both leaf miner species, a condition that suggests that the present parasitoids could mediate interactions between the present leaf miner species.

In addition, the influence of positive interactions on arthropod community structure associated to oak canopy, as mutualism has been reported in the literature. For example, Fernandes et al. (1999) reported a mutualistic relationship between three ant species (*Formica neurofibarbis*, *Liometopium apiculatum*, and *Monomorium cyaneum*) and the gall-forming wasp *Disholcaspis edura* that coexist in *Q. turbinella* canopy. The authors found that in the presence of ants, galls with the largest diameter suffered a lower mortality rate due to *Platigaster* sp. parasitoid attack. Thus, presence of ants reduced the selective pressure imposed by the parasitoid on wasp galls. Also, the occurrence of ecosystem engineers has been reported in oak canopy community structure literature. An example is the study of Lill and Marquis (2003) who examined the effect of shelter-building caterpillars *Pseudotelphusa* sp. (Gelechiidae) on the species richness and guild structure of leaf-chewing herbivores occupying individual white oak (*Quercus alba*) samplings in Missouri. The authors

found that the availability of leaf shelters created by *Pseudotelphusa* sp. within a tree's canopy is an important organizing factor because the species richness of the associated insect herbivore community was determinate. Similarly, Wang et al. (2012) using two species of leaf-tying caterpillars *Pseudotelphusa quercinigracella* and *Psilocorsis cryptolechiella* found that the presence of leaf ties increased species density and abundance of herbivores, predators, and scavengers, depending on the oak host species (*Q. alba*, *Q. imbricaria*, *Q. macrocarpa*, *Q. marilandica*, *Q. muehlenbergii*, *Q. velutina*, *Q. rubra*, and *Q. stellata*). Also, arthropod species composition differed between untied leaves and between ties made by the two leaf-tier species.

These associations among organisms across multiple trophic levels (Hooper et al. 2000; Sutherland 2006) may be mediated through “bottom-up” processes whereby diversity at higher trophic levels is governed by diversity at lower trophic levels or “top-down” control, via consumption by higher trophic levels (Hunter and Price 1992). Now, it is generally accepted that bottom-up and top-down forces act in concert to influence populations of most phytophagous insects (Hartvigsen et al. 1995; Hunter et al. 1997; Denno et al. 2002; Schönrogge et al. 2013). For example, temporal variation in the abundance of winter moth on oak trees in England depends largely upon the delayed density-dependent interaction of the herbivore with its natural enemies, a top-down effect (Hunter et al. 1997). In contrast, spatial variation in winter moth abundance among individual trees depends largely upon the density-independent effect of host-plant phenology, a bottom-up force. More recently, Schönrogge et al. (2013) quantified spatial patterns in native bird predation of invading gall inducing *Andricus* wasps associated with introduced Turkey oak (*Quercus cerris*) at eight sites across the UK. They found variation in gall density among trees driven by bottom-up influences of host oak (genotype) and simultaneously, a top-down regulation by birds of gall wasp populations.

In particular, researchers now readily acknowledge that interactions between plants, herbivores and natural enemies are commonplace (Forkner and Hunter 2000; Denno et al. 2002). Thus, host plants can impact herbivores directly by influencing their performance and survival (Lill and Marquis 2001; Sobek et al. 2009) and indirectly by mediating the effects of natural enemies (Lawton and McNeill 1979; Price et al. 1980). On the other hand, predators frequently have important influences on ecosystems through direct effects on the regulation of herbivorous insects densities and indirect effects propagated to primary producers (i.e., Marquis and Whelan 1994; Van Bael et al. 2003; Mäntylä et al. 2011). Finally, these relationships between species across different trophic levels may have effects on the

community structure and ecosystem functioning (Kagata and Ohgushi 2006; van Dam and Heil 2011).

Interactions involving plants, herbivores, and predators have been reported in oaks and the consequences for arthropod communities have been variables. For example, Marquis and Welan (1994) revealed that the presence of three species of birds (*Poliophtila caerulea*, *Regulus calendula*, and *R. satrapa*) enhances the growth of juveniles oaks of *Q. alba* via birds consumption of leaf-chewing insects (Lepidoptera: *Catocala amica*) in Missouri, USA. On the other side, Fernandes et al. (1999) reported that the interaction between ants (*Formica neorofibarbis*, *Liometopium apiculatum*, and *Monomorium cyaneum*) and gall wasps (*Disholcaspis* sp.) in *Quercus turbinella* canopy in Arizona, USA, reduced nearly half of the parasitism rate exercised by the parasitoid *Platygaster* sp. on galls of *D. edura*. In both studies, it is suggested that the third trophic level influences the population density of herbivorous insects but in an opposite direction. On the other hand, control bottom-up regulated by the heterogeneity in plant productivity and quality (i.e., nutrient availability, plant chemistry) has been recognized on the arthropod community structure associated to oaks. For example, Forkner and Hunter (2000) evaluated the importance of interactions bottom-up vs. top-down on *Q. rubra* and *Q. prinus* canopy insect communities. In this study, it was found that herbivorous insects were more abundant in plants of better quality (nitrogen, gallotannins, proanthocyanidins), and in response, predaceous insects were also more abundant. In contrast, exclusion of predators (birds) had not detectable effect on the presence or absence of herbivores or predaceous insects. The authors suggested that in these insect communities associated to oak canopies, bottom-up forces appear to regulate to influence the impact of top-down forces.

Finally, phytophagous insects frequently interact with pathogens within local communities (Hatcher 1995; Simon and Hilker 2003; Stout et al. 2006). Despite this fact, Stout et al. (2006) concluded in a review that pathogen-plant-insect interactions still receive limited attention. Most notably, few studies have examined interactions between plant-feeding insects and plant pathogens in the field or placed such interactions in a wider community context (Stout et al. 2006), oaks are no exception. In the literature, there are few studies that have addressed interactions between plant pathogens and herbivores. For example, Tack et al. (2012) evidence the interactions between three plant-feeding guilds: leaf miners, free-feeding insects, and the oak powdery mildew (*Erysiphe alphitoides*) in *Q. robur* canopy in Finland. The authors reported that individual species of insects responded differently to the intensity of pathogen infection and that the presence of the leaf miner *T. ekebladella* on oak leaves infected by powdery mildew decreased

the abundance of parasitoids emerging from these leaves in the following year. Both results suggest that mildew may drive the relative abundances of different species in different directions, thereby molding the structure of the full community.

We concluded that interactions between species are forces that influence the distribution, diversity, and abundances of arthropods associated to oak canopies and that these may act through different trophic levels in bottom-up and top-down directions. However, the outcome of individual interactions can range from positive or negative from the perspective of each involved species.

Disturbances

A disturbance has been defined as a discrete event that alters the structure of populations, communities, and ecosystems changing the availability of resources and conditions (White and Pickett 1985). Because disturbances are ecological processes that promote succession, producing vegetation mosaics with different degrees of structural complexity (White and Pickett 1985), they may alter the composition of the communities and the spatial patterns of diversity (Fagan et al. 1999; Cantrell et al. 2001). Specifically, the intensity, frequency, duration, and area of disturbance can determine the abundance and species richness in the communities (Huston 1994; Townsend et al. 1997). In general, temperate and tropical forests are subject to the effect of natural disturbances such as fire, storms, hurricanes, and floods (Dziöck et al. 2006; Martikainen et al. 2006), and of anthropogenic disturbances such as deforestation, agriculture, and urbanization (Hirao et al. 2007). In particular, this review addresses the effect of anthropogenic disturbances on the structure of communities associated with the canopy of oaks.

To date, there are several studies in tropical and temperate zones that have evaluated how human disturbances, in various forms and with different intensities is affecting the arthropod communities associated with the canopy (Hill et al. 1995; Ozanne et al. 1997, 2000; Floren and Linsenmair 2001, 2003; Tovar-Sánchez et al. 2003; Floren and Deeleman-Reinhold 2005; Forkner et al. 2006; Müller and Goßner 2007). However, these studies have shown inconsistent results, reporting that a disturbance can have a negative, positive, or no effect on the structure of arthropod communities. It has been suggested that these results can in part be explained by differences in dispersal abilities, habitat requirements, distribution patterns of different species of arthropods (Cooke and Roland 2000), as well as by differences in the historical factors and heterogeneity of the site (Hamer et al. 2003) and the scale and degree of the disturbance (Lewis 2001).

In particular, temperate forests are characterized by their diversity, richness, and longevity (Challenger 1998). Much of this diversity is found in the canopy, and arthropods constitute a major fraction of this species pool (Stork and Hammond 1997). Unfortunately, temperate forests are being converted into monoculture plantations and agricultural land, increasing the dominance of human activity on the landscape; the resulting fragmentation of forests has caused a marked decrease in biodiversity (Kimmins 1997; Maleque et al. 2006). Particularly, it has been suggested that canopy arthropods can be considered indicators of forest condition because they respond quickly to changes in environmental conditions (Schowalter 1995; Ozanne et al. 1997; Maleque et al. 2006, 2009).

Studies on the effect of disturbances on the canopy arthropod fauna associated to oaks have also shown conflicting results. On the one hand, there are studies like that of Tovar-Sánchez et al. (2003), who compared three forests with different degrees of disturbance and found a significant decrease in the abundance and diversity of ectophagous canopy arthropods associated to *Q. castanea*, *Q. crassipes*, *Q. crassifolia*, *Q. greggii*, *Q. laeta*, and *Q. rugosa* in the Valley of Mexico, in response to the disturbance gradient across forests. Similar responses were reported for the richness of herbivores associated with the canopy of *Q. alba* and for the richness and abundance of leaf-chewing insects associated with the canopy of *Q. alba* and *Q. velutina* in Missouri (Forkner et al. 2006, 2008).

Moreover, other studies reported that oak canopy arthropods did not respond to the disturbances. Such is the case of the beetles associated with the canopy of *Quercus* spp. in forest fragments with different degrees of urbanization (rural-suburban-urban); in this case, the composition and species richness did not differ in response to a gradient of disturbance in Bulgaria (Niemelä et al. 2002). The authors suggest that the beetles were not affected because the magnitude of the habitat disturbance in this gradient was moderate or because local factors such as temperature, moisture, and soil conditions determined the results.

As mentioned above, disturbances are environmental events that trigger the succession process of forests (White and Pickett 1985); thus, different studies have focused on the study of the succession process of plants following a disturbance (Jeffries et al. 2006); however, relatively little is known about the responses of arthropod communities to the succession process of their host plants (Morin 1999). It has been proposed that changes in the characteristics of plant communities as a result of a disturbance can influence arthropod communities, particularly if these specialize on certain species of host plants or on microhabitats generated by a particular

form of plant growth (Jeffries et al. 2006). In general, forests in late successional stages have greater structural complexity compared to young forests or plantations (Ishii et al. 2004); this is because the former have larger trees in terms of height, biomass, and structural complexity providing a more complex architecture of young forests (Schowalter 1995; Evans and Jukes 2000; Jeffries et al. 2006).

Studies in oaks on this topic are scarce; however, the few studies that have been conducted showed that the age of the forest does have an effect on the community structure of canopy arthropods in terms of richness, diversity, and density. For example, Summerville and Crist (2002, 2003) found a decrease in species richness and changes in the composition of the community of Lepidoptera associated with the canopy of *Quercus* spp. among forests in different successional stages (recently cleared vs. uncleared). Similarly, Jeffries et al. (2006) found that the diversity and density of leaf-chewing herbivorous insects (larvae of microlepidoptera, orthoptera, coleoptera, hymenoptera, and phasmids) were higher in the canopy of *Q. alba* in forests of 313 years of age, compared to sites recently affected by disturbances (2 years old). In both cases, the authors suggest that changes in the structure of these communities were the result of a shift of the forests to early successional stages.

Implications for conservation

Natural communities have been considered complex ecological systems whose structure and functioning are determined by the interaction of different factors that vary spatially and temporally (Bailey and Whitham 2006). In the previous sections, we discussed how oak canopy arthropods can respond to changes in genetic, architectural, chemical, phenological, successional factors, etc. and the influence of such factors in the establishment, organization, and maintenance of their communities. Under a conservation perspective, the sensitivity of this group of organisms is important because it suggests the role of arthropods as indicators of changes in the coverage, structure, and composition of forests. This condition may be the result of two particular attributes of arthropods: a short life cycle and little resilience (Brown 1997). Both features make these animals highly sensitive to changes in the parameters of the ecosystems in which they inhabit; thus, it has been suggested that in addition to their functional importance and contributions to biodiversity, arthropods can be very useful as an efficient early warning system for changes in habitat characteristics (Kremen et al. 1993; Schowalter 1995; Ozanne et al. 1997; Maleque et al. 2006, 2009). In this way, canopy arthropods could be useful to predict the consequences of climate change, disturbances,

or the loss of genetic diversity in oak forests. The use of canopy arthropods to examine the dynamics of the forest as well as alterations in vegetation cover by humans can be an efficient tool in time and cost compared to a longer-term perspective that implies information about palaeo-ecological record, pollen-based vegetation modeling, and biome reconstruction and global climate simulations (i.e., Feurdean et al. 2015).

Unfortunately, animal conservation strategies have focused on the preservation of species of mammals, birds, and terrestrial or aquatic vertebrates, groups of charismatic animals for which there are many ecological studies documenting the hazards faced by their habitats. This information has been helpful in implementing conservation and management programs (Leather et al. 2008). In contrast, efforts towards the conservation of arthropods have been limited, probably because many species have not been described or because their life history or their role within their communities (Redak 2000) is unknown. However, recent estimates suggest that this group contains the greatest diversity of species on earth (Hamilton et al. 2013), in addition to representing a group of vital organisms for the maintenance of a wide variety of processes at the level of communities and ecosystems, acting as pollinators, herbivores, or predators and participating in decomposition, nutrient cycling, energy transfer, or soil formation (McIntyre et al. 2001). This suggests that the loss of biodiversity of arthropods could have catastrophic consequences in ecological terms. Fortunately, in recent years, it has been proposed that the conservation of biodiversity must be carried out under an integrated approach that includes not only the conservation of unique species but also the conservation of their habitat and ecosystem processes (Bangert et al. 2005). This new approach may be essential to improve our ability to generate efficient conservation strategies (Wimp et al. 2004); in this sense, conserving the habitats and ecosystem processes of oak forests could benefit the arthropod communities associated with their canopy.

Conclusions

The canopy has been recognized for its important role in the functioning of terrestrial ecosystems and the maintenance of biodiversity. Arthropods constitute a major fraction of the species that inhabit this environment and oaks are one of the plant groups most representative of temperate forests worldwide. Despite this, we conclude that there is a lack of literature that addresses the study of arthropod communities associated with the oak canopies. There are still many aspects that haven't been addressed, which prevents us from making generalizations about some of the patterns found.

In general, the existing studies have revealed that the canopy of oaks supports arthropod communities with

high levels of diversity, that this diversity is the result of the great genetic, morphological, chemical, temporal, and spatial variation presented by this group of trees and shrubs, and that this heterogeneity can be observed at different levels: among individuals, species, localities, or regions. Most of these studies have focused on the response of the phytophagous insects of the Coleoptera, Hymenoptera, and Lepidoptera orders to changes in their host oaks. Research topics have been varied; however, most studies have focused on the effect of seasonal, spatial, and chemical variations in the host plant on the structure of arthropod communities, as well as on the effect of the hybridization of host plants on the resistance patterns of arthropods. The least studied aspects have been the effect of genetic diversity and of the origin (native vs. introduced) of the host oak. This review allowed us to find a lack of standardization in sampling and collection techniques, as well as inconsistencies in the criteria for defining the structure of the canopy. We also found that the existing studies did not consider variables such as habitat preference, type of feeding (guilds), or stage of development of the arthropods studied, which prevents generalizations about the patterns found. Furthermore, we could detect a bias towards the study of phytophagous insects and a lack of studies in other important groups such as acorn weevils, whose activity has an impact on the fitness and dispersion of host plants.

As a result of this review, we can conclude that the study of the arthropod fauna associated to oak canopy is a huge field of research to be explored, considering the large number of oak species that exist worldwide and the enormous diversity of arthropods that inhabit their canopy. As the knowledge of how the canopy arthropod communities respond to changes in their surrounding habitat moves forward, these animals may be recognized as bioindicators of oak forests health.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

Both authors participated in the review topic design, in the data analyses, and in the manuscript writing. Both authors read and approved the final manuscript.

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