

A comparison of hispine beetles (Coleoptera: Chrysomelidae) associated with three orders of monocot host plants in lowland Panama

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Abstract. The feeding traces in fossil ginger leaves and the conserved phylogenetic relationships seen today in certain clades of hispine beetles on their monocot hosts point towards a long and intimate plant–insect evolutionary relationship. Studies in the 1970s and 1980s documented the rich fauna of rolled-leaf hispine beetles and their association with the Neotropical monocot family Heliconiaceae in Central America. In this report, the taxonomic breadth of these early studies is expanded to include species in the families, Marantaceae, Poaceae, Arecaceae and Costaceae, all with species occurring sympatrically with the Heliconiaceae in lowland Panama. Additionally, the analysis is widened to include open-leaf scraping and internal leaf-mining clades of hispoid Cassidinae. The censuses add more than 5080 Cassidinae herbivore occurrence records on both open and unfurled new leaf rolls of 4600 individual plants. Cluster analysis reveals that while many Hispinae species tend to group with plant species in only one of the three monocot orders, 9 of 16 Hispinae species on Zingiberales hosts were recorded in substantial numbers on both the Heliconiaceae and the Marantaceae, indicating an underlying pattern of feeding flexibility at the host plant family level.

Key words: Cassidinae, Arecales, Poales, Zingiberales, hispine beetles, species assemblages

Introduction

Interactions among plants and phytophagous insects play a key role in community function and evolution. Trophic relationships provide useful insights into understanding important community dynamics (Chave, 2004). Relationships with host plants and food specialization that evolved in the past may shape contemporaneous communities and species assemblages (Novotny *et al.*, 2002). Diet and degree of food specialization are poorly

characterized for many tropical phytophagous insects, including many species of hispine beetles (Coleoptera: Chrysomelidae: Cassidinae), a group distributed worldwide containing approximately 3000 species distributed among 24 tribes and 170 genera (Staines, 2002). The relationships between selected groups of host plants and their phytophagous insects can provide insights into the evolution of feeding associations and their phylogenetic constraints (Dobler *et al.*, 1996). Such an approach

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requires sampling a large number of potential host plant species during consecutive seasons for an entire guild of the phytophagous insects. Our study compares the assemblages of hispine beetle species associated with five families of monocotyledonous plants, Heliconiaceae, Marantaceae and Costaceae (Zingiberales), Poaceae (Poales) and Arecaceae (Arecales) in a single geographical area, the lowland Neotropical forests of central Panama.

The diverse New World hispine beetles include 11 tribes, among which Cephaloleiini or 'rolled-leaf hispines' have cryptophagic larvae and are largely associated with three orders of plants, Poales, Arecales and Zingiberales (Jolivet, 1997). Feeding grooves in fossilized ginger leaves of late Cretaceous age suggest that the Cephaloleiini clade of hispine beetles may have had a long and intimate plant-insect evolutionary relationship with their monocot hosts (Wilf *et al.*, 2000). The hispine tribes, Chalepini and Sceloenoplini, have endophagic or 'leaf-mining' larvae and are associated with a diverse set of both monocotyledonous and dicotyledonous host plant families (Hespenheide and Dang, 1999). The Hemisphaerotini, a relatively small tribe of putative hispines, have exophagic larvae and are associated principally with the New World monocotyledonous orders: Pandanales (Cyclanthaceae) (D. Windsor, personal communication), Poales (Bromeliaceae) (D. Windsor, personal communication), Arecales and Zingiberales. Aspects of the natural history and ecology of these three groups are discussed in Jolivet (1997), Strong (1977a,b, 1983), Chaboo and Nguyen (2004) and Chaboo (2007). Thus, with the exception of some genera of leaf-mining hispines, the vast majority of taxa are associates of basal angiosperms in the Monocotyledonae (Soltis *et al.*, 2005). A basal position for several hispine taxa within the Cassidinae subfamily was indicated in an early phylogenetic reconstruction using a fragment of the 12S ribosomal mtDNA gene (Hsiao and Windsor, 1999) and in a subsequent morphological study (Chaboo, 2007). With only minor exceptions, both adult and larval hispines feed on the same host plant species (Jolivet, 1997). Previous studies of hispine beetles examined communities associated with a single monocot genus *Heliconia* (Seifert and Seifert, 1976; Seifert, 1982; Strong, 1977a,b, 1982, 1983; Staines, 2002). The actual host plants of more than 70% of hispine genera are still unidentified or poorly supported (Jolivet, 1997). In this paper, we present additional data on the assemblage of hispine species associated with *Heliconia* species and we compare those to the hispine assemblages occurring on other species of monocots in the same habitat. Because the term 'community' when referring to a collection of hispine species can be misleading or ambiguous (Strong, 1977b, 1982), we

use the term 'species assemblage' (Allaby, 2003) in this paper to refer collectively to all hispine species occurring on a specified set of host plants.

The summary trees of commelinoid phylogeny (Janssen and Bremer, 2004; Soltis *et al.*, 2005) indicate that the Arecaceae is the older plant family in our study, splitting off from the core monocots before the Poaceae and the Zingiberales. The Heliconiaceae branched from Zingiberales stock earlier than the Costaceae and these earlier than the Marantaceae. Janssen and Bremer (2004) showed that these three phylogenetically separated families of Zingiberales often co-occur in the understorey of Neotropical forests. Williams and Harborne (1977) reported substantial differences in flavonoid content between these three Zingiberales families. The Marantaceae and Heliconiaceae are the most abundant Zingiberales in many lowland areas of Central America (Seifert and Seifert, 1976; Seifert, 1982; Strong, 1977a,b, 1982; Flowers and Janzen, 1997). Our study examines the hypothesis that hispine beetles will share structurally similar species of plants occurring in the same or nearby habitats. An alternative hypothesis is that hispine beetles will share plants having similar histories of diversification. That is, flowering plants may have shaped beetle diversification leaving host plant associations at the family level as an enduring record of this process. In particular, older plant families such as the Arecaceae (Soltis *et al.*, 2005) are expected to support more diverse faunas and more basal taxa than are found on younger families such as the Marantaceae. Our study also examines Strong's contentions that broadly distributed Zingiberales food plants will have more associated hispine species than narrowly distributed ones and that a plant species will have a constant number of associated hispines throughout its range but that the particular species may change according to location ('geographical replacement') (Strong, 1977b, 1982).

Materials and methods

Study area, host plants and sampling method

Observations were made within forests along the east bank of the Panama Canal including a site near Gamboa in Parque Natural Soberania (9°7'N, 79°42'W), along the trail system on Barro Colorado Island (9°9'N, 79°51'W) and within the Parque Metropolitano (8°59'N, 79°33'W; near Panama City). Observations were made during 18 total months within both dry and wet seasons between May 2002 and July 2006.

Our study is based on observations recorded from a total of 17 species of monocot host plants (Table 1). Twelve of these species are within the

order Zingiberales and included six species of Heliconiaceae, five species of Marantaceae and one species of Costaceae. Species within these families, all native to the Neotropics (Croat, 1978), are characterized by large and colourful bracts and by young leaves rolled into long tubes opening after the periods from a few days to a few weeks into large mature leaves with transverse venation, long petioles and high-flavonoid content (Berry and Kress, 1991). The scroll-like immature leaves are the typical habitat for 'rolled-leaf' hispid Cassidinae (Strong, 1983; Jolivet, 1997). This study additionally includes four species of the Arecaceae whose young leaflets initially form relatively compact sword-like structures, which subsequently begin to open and provide habitat for hispines as the leaf expands to full size. Finally, the study includes one species of the Poaceae (*Chusquea simpliciflora* Munro) whose new leaves are rolled at the base.

Ecologically, these five families occur in a range of habitats from old forest to human-disturbed areas. We classified each species into one of seven habitat categories according to the information given by Croat (1978) and Kress (1990) (Table 1). Host plant species were chosen because of their abundance in the study sites (Croat, 1978; personal observations) and the presence of a diversified

hispid Cassidinae fauna (Strong, 1977a,b; Staines, 2004; D. Windsor, personal communication). Thus, the study lacks data from rare host plant taxa and those taxa that are rarely or never fed upon by hispines. While the host plant species are all part of the native flora of Panama (Croat, 1978), all of them occur elsewhere in Central and South America (Table 1).

We recorded the number of leaves unrolled or expanded for each individual plant that was sampled and the number of species and developmental stage (egg, larva and pupa) for hispine beetles observed or collected. Plants occurring along trails were surveyed inside young (60- to 120-year-old) and old (400-year-old) forests on BCI and along trails in mainland forests (Parque Soberania and Parque Metropolitano). Hispine species were identified based on the plates in the Biologia Centrali-Americana (Baly, 1885–1894), descriptions and key given by Staines (1996, 2002) and by comparison to identified specimens in the STRI insect collection. Beetles species were normally easily identified in the field but were collected when their identification was in doubt, as with leaf-miner larvae which were reared in the laboratory to obtain and identify the adults. Study sites were visited once a week during the sampling

Table 1. Host plant characteristics and numbers of each plant species sampled

Host plant family and species	Range ¹	Habitat ²	Size ¹	No.
Arecaceae				
<i>Bactris major</i> Jacq.	Panama	Understorey (forest)	8 m	172
<i>Chamaedorea wendlandiana</i> Hemsl.	Nicaragua to Panama	Understorey (forest, near streams)	Up to 5 m	358
<i>Cryosophila warscewiczii</i> Bartl.	Belize to Colombia	Understorey (younger forest)	Up to 10 m	196
<i>Oenocarpus panamanus</i> Bailey	Costa Rica and Panama	Understorey (forest)	8–20 m	289
Costaceae				
<i>Costus pulverulentus</i> Presl	Mexico to Ecuador	Old tree-fall gap	0.5–3.5 m	63
Heliconiaceae				
<i>Heliconia catheta</i> R. R. Smith	Panama	Tree-fall gap	3–5 m	156
<i>Heliconia irrasa</i> R. R. Smith Lane	Mexico to Colombia	Understorey (forest)	1.5–2 m	242
<i>Heliconia latispatha</i> Benth.	Mexico to South America	Disturbed area	1.5–3 m	± 740
<i>Heliconia mariae</i> Hook	Belize to Panama	Understorey (forest)	3–6 m	82
<i>Heliconia vaginalis</i> Benth.	Mexico to Ecuador	Tree-fall gap	1–3 m	125
<i>Heliconia wagneriana</i> O. G. Petersen	Honduras to Panama	Clearings	Up to 6 m	39
Marantaceae				
<i>Calathea inocephala</i> Kenn. & Nic.	Mexico to Peru	Understorey (forest)	2–4 m	831
<i>Calathea insignis</i> O. G. Petersen	Mexico to Peru	Tree-fall gap	2–3 m	227
<i>Calathea latifolia</i> Klotzsch	Venezuela to Panama	Disturbed area	1–2 m	452
<i>Calathea lutea</i> Schult.	Mexico to Peru	Tree-fall gap	2–3 m	51
<i>Ischnosiphon pruinosis</i> O. G. Petersen	Belize to Panama	Tree-fall gap	2–3 m	247
Poaceae				
<i>Chusquea simpliciflora</i> Munro	Guatemala to Panama	Forest	Up to 25 m ± 400	

Range, distribution area; No., number of visited individuals per host plant species in the present study.

¹Croat (1978).

²Croat (1978) and Berry and Kress (1991).

periods. Because leaf rolls and mined leaves had to be destructively removed, each individual plant was surveyed only once a field season to minimize disturbance to beetle assemblages. The scroll-like immature leaves were only included in the study if they were closed at the bottom (watertight) thereby maintaining a high degree of humidity, which appeared to be an important factor for 'rolled-leaf' Cassidinae communities (Strong, 1977a). A set of voucher specimens was deposited in the collections of the Smithsonian Tropical Research Institute in Panama.

Statistical analysis: species richness and community analysis

Hispine species accumulation curves for each host plant species were generated using five different estimators: Chao 1, Chao 2, Jackknife 1, Jackknife 2 and Bootstrap (Species Diversity and Richness, Version 3.0, Pisces Conservation Ltd, 2002b). The sampling of hispine species was considered exhaustive when the species accumulation curve for that plant species approached an asymptote. The faunas of two species, *Costus pulverulentus* Presl and *Heliconia vaginalis* Benth., did not reach an asymptote indicating further sampling is warranted for these species.

To determine whether factors proposed by Strong (1977a,b) had a significant impact on Cassidinae species richness and average abundance per plant, we performed a Kruskal–Wallis one-way ANOVA (Systat, Version 10, SPSS Inc., 2000). This analysis was applied independently to six matrices (three factors: plant size (height: <3 m or >3 m), plant habitat (forest and understorey or disturbed) and plant range (Central America or Central America plus South America); two variables (richness and average abundance). The sampling effort was not constant for each plant species; therefore, we normalized herbivore species occurrence data by dividing the number of hispine individuals observed on a particular host plant species by the number of individuals of that plant species surveyed. To define the hispine species assemblages associated with each host plant species, we applied 'average linkage' method and Bray–Curtis' distance clustering to normalized field data, procedures widely used in ecological studies with species abundance. The cluster analysis allows identifying a set of groups (hispine assemblages), which both minimize within-group variation and maximize between-group variation (Dufrêne and Legendre, 1997; Legendre and Legendre, 1998; Community Analysis Package, Version 2.0, Pisces Conservation Ltd, 2002a).

Results

More than 4670 host plant individuals (17 species) were inspected (Table 1) and 5080 hispid Cassidinae individuals were collected or identified in place (1013 on five species of Marantaceae, 1072 on six species of Heliconiaceae, 11 on one species of Costaceae, 36 on one species of Poaceae and 2948 on four species of Arecaceae). The host plant species were classified into seven habitats and had ranges varying from Mexico to South America (Table 1). The Cassidinae observed in this study belonged to 30 species distributed among 11 genera (Table 2). Two Cassidinae species are apparently undescribed and were assigned provisional morphospecies names (*Cephaloleia* sp61 and *Spaethiella* sp47).

Hispine species richness per plant species ranged from 1 to 12 species (median number of hispine species per host = 5) (Fig. 1). The plant species with the richest assemblages of hispines were *Heliconia latispatha*, *H. catheta* and *Calathea latifolia* with 12, 10 and 9 species, respectively. The smallest assemblages were found on *Cryosophila warszewiczii*, *C. simpliciflora* and *H. vaginalis* with 1 (*Prosopodonta dorsata*), 3 and 3 species, respectively. At the family level, the average richness of hispines was 7.8 for Heliconiaceae, 5.8 for Marantaceae, 4 for Costaceae, 3 for Poaceae and 2.5 for Arecaceae. The numbers of hispine species associated with Heliconiaceae and Marantaceae species were not significantly different (Mann–Whitney $U = 19.0$, $P = 0.457$, $df = 1$).

Neither host plant habitat, host plant range nor the average size of host plants had a significant effect on hispine richness (host plant habitat: Mann–Whitney $U = 29.5$, $P = 0.53$, $df = 1$; plant range: Mann–Whitney $U = 47.5$, $P = 0.26$, $df = 1$; plant size: Mann–Whitney $U = 49.5$, $P = 0.19$, $df = 1$) or on hispine abundance (host plant habitat: Mann–Whitney $U = 42.0$, $P = 0.56$, $df = 1$; plant range: Mann–Whitney $U = 51.0$, $P = 0.15$, $df = 1$; plant size: Mann–Whitney $U = 30.0$, $P = 0.56$, $df = 1$).

Ten well-defined hispine species groups (containing a total of 30 species) were arranged by cluster analysis (average linkage method; Bray–Curtis' distance) (Fig. 2). The first dichotomy in the output separates the hispine species clusters associated with the Zingiberales from those associated with the Poales + Arecales. The next branching within the Zingiberales separates the hispines associated with the three Zingiberales families, Marantaceae, Heliconiaceae and Costaceae. The second branching of the Poales + Arecales separates these two orders. Each hispine species fell within only one cluster group (Table 3). Two hispine species, *C. stevensi* and *D. panamensis*, clustered only with themselves and diverged early in the statistical output (Fig. 2).

Table 2. The list of hispine species observed in the present study ($n = 30$), species abbreviations used in the cluster analysis (Fig. 2), and species feeding habits and larval morphologies (Strong, 1977a,b, 1983; Staines, 1996, 2002, 2004; D. Windsor pers. commun.)

Hispine species	Code	Habitat	Larvae shape	Dorsal shield	Adult habitat
Arescini					
<i>Chelobasis bicolor</i> Gray	Ch.bic	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. perplexa</i> Baly	Ch.per	Rolled leaf	Onisciform	None	Rolled leaf
Cephaloleiini					
<i>Aslamidium semicircularum</i> Olivier	As.sem	Open leaf	Onisciform	None	Open leaf-upper side
<i>Cephaloleia belti</i> Baly	Ce.bel	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. costaricensis</i> Uhmann	Ce.cos	Rolled or open leaf	Onisciform	None	Rolled leaf
<i>C. dilaticollis</i> Baly	Ce.dil	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. dorsalis</i> Baly	Ce.dor	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. erichsonii</i> Baly	Ce.eri	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. instabilis</i> Baly	Ce.ins	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. lata</i> Baly	Ce.lat	Rolled or open leaf	Onisciform	None	Rolled leaf
<i>C. metalescens</i> Baly	Ce.met	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. partita</i> Weise	Ce.par	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. pretiosa</i> Baly	Ce.pre	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. sallei</i> Baly	Ce.sal	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. stevensi</i> Baly	Ce.ste	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. suturalis</i> Baly	Ce.sut	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. trimaculata</i> Baly	Ce.tri	Rolled leaf	Onisciform	None	Rolled leaf
<i>C. sp61</i>	Ce.sp61	Rolled or open leaf	Onisciform	None	Rolled leaf
<i>Imatidium thoracicum</i> Fabr.	Im.tho	Open leaf	Onisciform	None	Open leaf-lower side
<i>Homalispis nevermanni</i> Uhmann	Ho.nev	Rolled leaf	Onisciform	None	Rolled or open leaf
Chalepini					
<i>Platocthispa emorsitans</i> Baly	Pl.emo	Leaf miner	Onisciform	None	Open leaf
Delocraniini					
<i>Delocrania panamensis</i> Champion	De.pan	Open leaf	Onisciform	Detritus	Open leaf-lower side
Hemisphaerotini					
<i>Spaethiella circumdata</i> Boheman	Sp.cir	Open leaf	Eruciform	Exuvial-fecal shield	Open leaf
<i>S. marginata</i> Champion	Sp.mar	Open leaf	Eruciform	Exuvial-fecal shield	Open leaf
<i>S. tristis</i> Boheman	Sp.tri	Open leaf	Eruciform	Exuvial-fecal shield	Open leaf
<i>S. sp47</i>	Sp.47	Open leaf	Eruciform	Exuvial-fecal shield	Open leaf
Imatidiini					
<i>Stilpnaspis panamensis</i> Borowiec	St.pan	Rolled or open leaf	Onisciform	None	Open leaf
Prosopodontini					
<i>Prosopodonta dorsata</i> Baly	Pr.dor	Open leaf	Onisciform	Urogomphus	Open leaf
Spilophorini					
<i>Calyptocephala antennata</i> Spaeth	Ca.ant	Open leaf	Onisciform	Exuvial shield	Open leaf
<i>Calyptocephala brevicornis</i> Boheman	Ca.bre	Open leaf	Onisciform	Exuvial shield	Open leaf

'Onisciform' larvae are flattened with extended segmental margins contacting the substrate, approximately oval in shape when viewed from above, while 'eruciform' larvae are circular in cross section, their mass largely supported by their legs and they are tubular shaped.

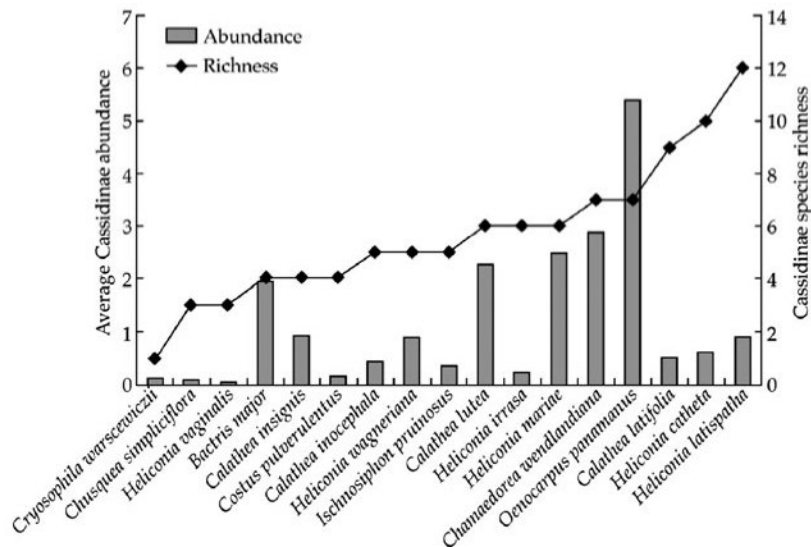


Fig. 1. Hispine species abundance (the average number of hispine individuals observed per plant) and richness (number of hispines species per plant species) on 17 species of monocotyledonous host plants

Discussion

This study represents the first large-scale survey of hispid Cassidinae on the Isthmus of Panama. Thirty species of hispid Cassidinae were encountered on leaves of more than 4670 individual plants in five monocotyledonous families (Table 4). When normalized by number of individuals

per host plant species, the occurrence data suggest that strong preferences exist within the hispid Cassidinae at the host plant ordinal level but less so at the familial and generic levels. Only one hispine species *Spaethiella tristis* was found on more than one plant order (Poales and Arecales), while 12 hispine species were found on more than

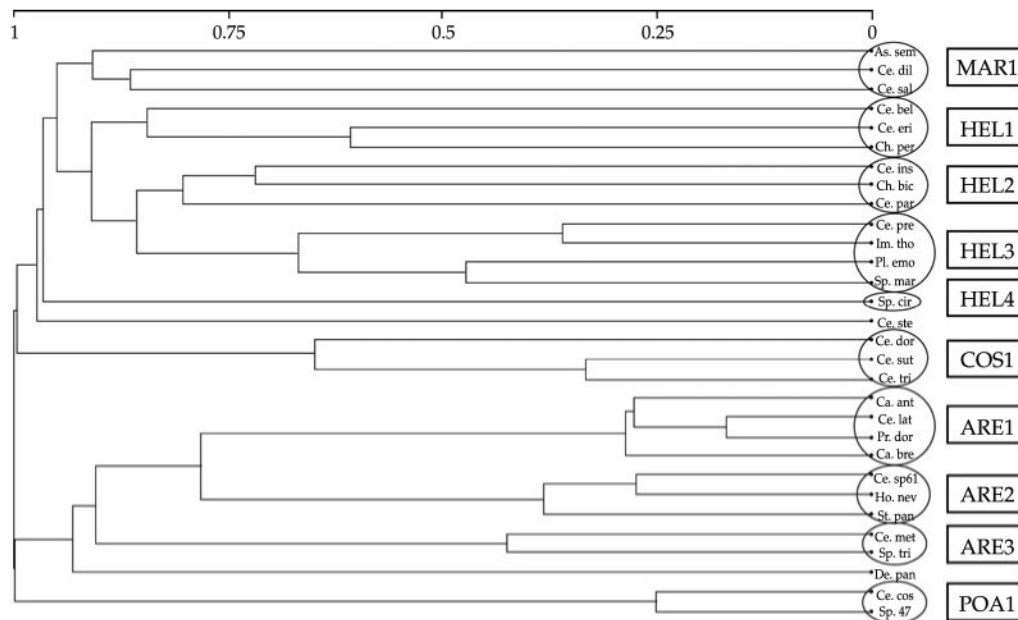


Fig. 2. Cluster analysis: Dendrogram of hispid Cassidinae species (average linkage method; Bray-Curtis' distance). ARE1, first cluster group of Areaceae; ARE2, second cluster group of Areaceae; ARE3, third cluster group of Areaceae; HEL1, first cluster group of Heliconiaceae; HEL2, second cluster group of Heliconiaceae; HEL3, third cluster group of Heliconiaceae; HEL4, fourth cluster group of Heliconiaceae; COS1, cluster group of Costaceae; MAR1, cluster group of Marantaceae; POA1, cluster group of Poaceae

Table 3. Results of cluster group analyses (dendrogram; average linkage method; Bray–Curtis' distance)

Code for cluster groups	Included host plant species	Habitat	Hispine species ⁺
ARE1	<i>Chamaedorea wendlandiana</i>	Understorey (forest, near streams)	Ca.ant, Ce.lat, Pr.dor, Ca.bre
ARE2	<i>Cryosophila warscewiczii</i>	Understorey (younger forest)	Ce.sp61, Ho.nev, St.pan
	<i>Oenocarpus panamanus</i>	Understorey (forest)	
ARE3	<i>Bactris major</i>	Understorey (forest)	Ce.met, Sp.tri
COS1	<i>Costus pulverulentus</i>	Old tree-fall gap	Ce.dor, Ce.sut, Ce.tri
HEL1	<i>Heliconia mariae</i>	Forest	Ce.bel, Ce.eri, Ch.per
HEL2	<i>Heliconia catheta</i>	Tree-fall gap	Ce.ins, Ch.bic, Ce.par
	<i>Heliconia vaginalis</i>	Tree-fall gap	
	<i>Heliconia wagneriana</i>	Clearings	
HEL3	<i>Heliconia latispatha</i>	Highly disturbed area (anthropic)	Ce.pre, Im.tho, Pl.emo, Sp.mar
HEL4	<i>Heliconia irrasa</i>	Understorey (forest)	Sp.cir
MAR1	<i>Calathea inocephala</i>	Understorey (forest)	As.sem, Ce.dil, Ce.sal
	<i>Calathea insignis</i>	Tree-fall gap	
	<i>Calathea latifolia</i>	Disturbed area	
	<i>Calathea lutea</i>	Tree-fall gap	
	<i>Ischnosiphon pruinosus</i>	Tree-fall gap	
POA1	<i>Chusquea simpliciflora</i>	Forest	Ce.cos, Sp.47

Habitat, plant species habitat (Croat, 1978; Berry and Kress, 1991). ARE1, first cluster group of Arecaceae; ARE2, second cluster group of Arecaceae; ARE3, third cluster group of Arecaceae; HEL1, first cluster group of Heliconiaceae; HEL2, second cluster group of Heliconiaceae; HEL3, third cluster group of Heliconiaceae; HEL4, fourth cluster group of Heliconiaceae; COS1, cluster group of Costaceae; MAR1, cluster group of Marantaceae; POA1, cluster group of Poaceae.

⁺ See Table 2 for hispine species abbreviations.

one plant family. No hispine species were found to be randomly distributed among host plant families, genera or host plant species (Fig. 2).

While several factors have been suggested to influence hispid Cassidinae richness on particular host plant species (Strong, 1977a, b), our study does not support either host plant size, range or habitat type as factors significantly affecting either richness or abundance. However, it is possible that some or all of these factors working in combination could explain the high richness of the hispine assemblage associated with *H. latispatha*. This species has 12 hispine species (4 open-leaf scrapers, 1 leaf miner and 7 'rolled-leaf' hispines), and the largest range of the 17 plant species in the study. It is of moderate to large size, frequents forest margins and occurs in dense patches. Moreover, each of the three richest plant species *C. latifolia*, *H. catheta* and *H. latispatha* occurs in disturbed areas and tree-fall gaps (Table 1). Differences among species in major chemical content at the family level as described by Williams and Harborne (1977) could also contribute to the differences in hispine richness we observed among plant species. Seasonality in the production of rolled leaves is another factor that could affect species richness. The Heliconiaceae (Croat, 1978) produce young rolled leaves throughout the year while other plant families may be more seasonal in the production of rolled leaves. The Heliconiaceae thus may offer a more stable habitat than the other understorey species and thereby attract and retain

more 'rolled-leaf' hispine colonists (Fig. 1). The inclusion of only four Arecaceae species in the study, two of which had very low-hispine richness and two of which had moderate-to-high hispine richness, makes any conclusion about the importance of the phylogenetic 'age' of this family premature.

Strong (1977a,b, 1982) recorded 12 'rolled-leaf' hispid Cassidinae species occurring on *Heliconia* spp. in different areas in Central America while we found nine 'rolled-leaf' species in Panama (Table 2). However, only three of these species *Cephaloleia instabilis*, *Chelobasis bicolor* and *C. perplexa* occur on the same host plant species (Table 2). The difference in these faunas (Table 2) does suggest that considerable geographical replacement of hispine species in *Heliconia* communities may occur across Central America.

The results of our study also suggest that host plant data are under reported in the literature, possibly a product of sampling a narrower range of plant families and hispine tribes (Jolivet, 1997; Fernandez and Hilker, 2007). Laboratory feeding trials will be essential to know just how accurately occurrence data as presented in this study reflect actual feeding habits of hispines (Descampe *et al.*, 2007, in press).

Cluster analysis shows that hispine assemblages (Figs 2 and 3) are primarily determined by host plant affiliation (order and family) and secondarily by plant habitat. *H. latispatha* and *C. latifolia*, which occur in similar habitats, present clearly distinct

Table 4. Hispine occurrence data from the present study and that reported by previous studies

Hispine species	Host plant in present study	No.	McKenna ¹	Staines ²	Strong ³
<i>Aslamidium semicircularum</i>	<i>Calathea inocephala</i>	113	–	Zingiberales	–
	<i>Calathea insignis</i>	180			
	<i>Calathea latifolia</i>	32			
	<i>Calathea lutea</i>	102			
	<i>Heliconia latispatha</i>	1			
<i>Calyptocephala antennata</i>	<i>Ischnosiphon pruinosus</i>	79			
	<i>Chamaedorea wendlandiana</i>	286	–	–	–
	<i>Oenocarpus panamanus</i>	10			
<i>Calyptocephala brevicornis</i>	<i>C. wendlandiana</i>	197	–	–	–
<i>Cephaloleia belti</i>	<i>C. latifolia</i>	4	<i>Heliconia</i> sp. – PA	<i>Heliconia imbricata</i> – CR	
	<i>C. lutea</i>	2	<i>H. sp.</i> – CR	<i>H. latispatha</i>	
	<i>I. pruinosus</i>	1		<i>Heliconia pogonantha</i>	
	<i>Heliconia catheta</i>	47		<i>Heliconia mariae</i>	
	<i>Heliconia irrasa</i>	5		<i>Heliconia tortuosa</i>	
	<i>H. latispatha</i>	101			
	<i>H. mariae</i>	85			
	<i>Heliconia wagneriana</i>	13			
<i>C. costaricensis</i>	<i>Chusquea simpliciflora</i>	12	–	–	–
<i>C. dilaticollis</i>	<i>C. inocephala</i>	86	<i>C. lutea</i> – PA	<i>C. insignis</i> – CR	–
	<i>C. insignis</i>	4	<i>Calathea</i> sp. – PA		
	<i>C. lutea</i>	2	<i>Calathea</i> sp. – EC		
	<i>I. pruinosus</i>	1	<i>Renealmia</i> sp. – EC		
	<i>Costus pulverulentus</i>	7	<i>Renealmia</i> sp. – EC	<i>Costus</i> sp. – CR	–
<i>C. dorsalis</i>				<i>C. gymnocarpa</i> – CR	–
<i>C. erichsonii</i>	<i>C. inocephala</i>	186	<i>Calathea gymnocarpa</i> – CR	<i>C. inocephala</i>	
	<i>C. insignis</i>	1	<i>Calathea</i> sp. – CR	<i>Calathea leucostachys</i>	
	<i>C. latifolia</i>	4	<i>Heliconia</i> sp. – CO		
	<i>C. lutea</i>	3	<i>Calathea</i> sp. – PA		
	<i>H. catheta</i>	1			
	<i>H. latispatha</i>	5			
	<i>H. mariae</i>	70			
	<i>Heliconia vaginalis</i>	1			
	<i>C. latifolia</i>	1	<i>H. wagneriana</i> – CR	<i>H. latispatha</i>	<i>Heliconia platystachys</i> – PA
	<i>H. catheta</i>	1		<i>H. imbricata</i>	<i>Heliconia curtispatha</i> – PA
<i>C. insignis</i>	<i>H. latispatha</i>	38		<i>H. wagneriana</i>	<i>H. latispatha</i> – PA
	<i>H. mariae</i>	2			<i>H. latispatha</i> – CR
					<i>H. wagneriana</i> – CR
<i>C. lata</i>	<i>C. wendlandiana</i>	195	<i>Chamaedorea tepejilote</i>	–	–
<i>C. metalescens</i>	<i>Bactris major</i>	215	Unknown – CR	–	–
	<i>C. wendlandiana</i>	4			
<i>C. partita</i>	<i>C. latifolia</i>	2	<i>H. latispatha</i> – PA	<i>Heliconia</i> sp.	–
	<i>H. catheta</i>	2	<i>Heliconia</i> sp. – PA		

Table 4. Continued

Hispine species	Host plant in present study	No.	McKenna ¹	Staines ²	Strong ³
<i>C. pretiosa</i>	<i>H. latispatha</i>	1	<i>Heliconia</i> sp. – CO	–	–
	<i>C. pulverulentus</i>	1			
	<i>H. catheta</i>	12			
	<i>H. latispatha</i>	50			
	<i>H. mariae</i>	12			
<i>Cephaloleia sallei</i>	<i>H. wagneriana</i>	8	<i>H. irrasa</i> – CR	<i>Heliconia</i> sp. – CR <i>Renealmia strobilifera</i>	–
	<i>C. inocephala</i>	1			
	<i>C. latifolia</i>	168			
	<i>C. lutea</i>	4			
	<i>I. pruinus</i>	4			
	<i>H. catheta</i>	1			
	<i>H. latispatha</i>	11			
	<i>H. mariae</i>	1			
<i>C. stevensi</i>	<i>C. inocephala</i>	2	<i>Calathea micans</i> – CR	<i>Heliconia</i> sp. – CR	–
	<i>C. latifolia</i>	1			
	<i>I. pruinus</i>	1			
<i>C. suturalis</i>	<i>C. pulverulentus</i>	1	<i>Costus</i> sp. – CR	–	–
<i>C. trimaculata</i>	<i>C. pulverulentus</i>	2	<i>Renealmia</i> sp. – PA	Costaceae – CR	–
<i>C. sp61</i>	<i>B. major</i>	1	–	–	–
	<i>C. wendlandiana</i>	1			
	<i>O. panamanus</i>	376			
	<i>C. latifolia</i>	1	–	<i>Heliconia</i> sp. – CR	<i>H. curtispatha</i> – CR
<i>Chelobasis bicolor</i>	<i>H. catheta</i>	1			<i>H. imbricata</i> – CR
	<i>H. irrasa</i>	1			<i>H. irrasa</i> – CR
	<i>H. latispatha</i>	10			<i>H. latispatha</i> – PA
	<i>H. vaginalis</i>	2			<i>H. latispatha</i> – CR
	<i>H. wagneriana</i>	1			<i>Heliconia metallica</i> – CR
					<i>H. platystachys</i> – PA
<i>Chelobasis perplexa</i>	<i>H. irrasa</i>	1	<i>H. pogonantha</i> – CR	<i>C. insignis</i> – CR	<i>H. wagneriana</i> – CR
	<i>H. mariae</i>	23		<i>H. imbricata</i> – CR	<i>H. imbricata</i> – CR
<i>Delocrania panamensis</i>	<i>B. major</i>	1	–	–	–
	<i>O. panamanus</i>	20			
<i>Homalispia nevermanni</i>	<i>O. panamanus</i>	659	–	–	–
<i>Imatidium thoracicum</i>	<i>C. lutea</i>	3	–	Arecaceae – CR	–
	<i>H. catheta</i>	11		Zingiberales	
	<i>H. irrasa</i>	1		<i>C. insignis</i> – CR	
	<i>H. latispatha</i>	83			
	<i>H. wagneriana</i>	12			
<i>Platocthispa emorsitans</i>	<i>C. insignis</i>	8	–	<i>Calathea</i> sp.	–

Table 4. Continued

Hispine species	Host plant in present study	No.	McKenna ¹	Staines ²	Strong ³
<i>Prosopodonta dorsata</i>	<i>C. latifolia</i>	17		<i>Costus</i> sp.	
	<i>H. catheta</i>	1			
	<i>H. irrasa</i>	11		Piperaceae	
	<i>H. latispatha</i>	221			
	<i>C. wendlandiana</i>	170	–	Arecaceae	–
	<i>Cryosophila warscewiczii</i>	24			
<i>Spaethiella circumdata</i>	<i>O. panamanus</i>	287			
	<i>H. irrasa</i>	37	–	–	–
	<i>H. latispatha</i>	1			
<i>Spaethiella marginata</i>	<i>H. vaginalis</i>	2			
	<i>C. insignis</i>	19	–	–	–
	<i>H. catheta</i>	16			
	<i>H. latispatha</i>	159			
<i>Spaethiella tristis</i>	<i>H. mariae</i>	9			
	<i>H. wagneriana</i>	1			
	<i>B. major</i>	117	–	–	–
	<i>C. simpliciflora</i>	4			
	<i>O. panamanus</i>	122			
	<i>C. simpliciflora</i>	20	–	–	–
<i>Stilpnaspis panamensis</i>	<i>C. wendlandiana</i>	180	–	–	–
	<i>O. panamanus</i>	83			

Country codes are: PA, Panama; CR, Costa Rica; CO, Colombia; EC, Ecuador; species in bold, species common to both this study and previous reports.

¹McKenna and Farrel (2005, 2006).

²Staines (1996, 2002, 2006).

³Strong (1977a,b, 1983); Strong et al. (1984).

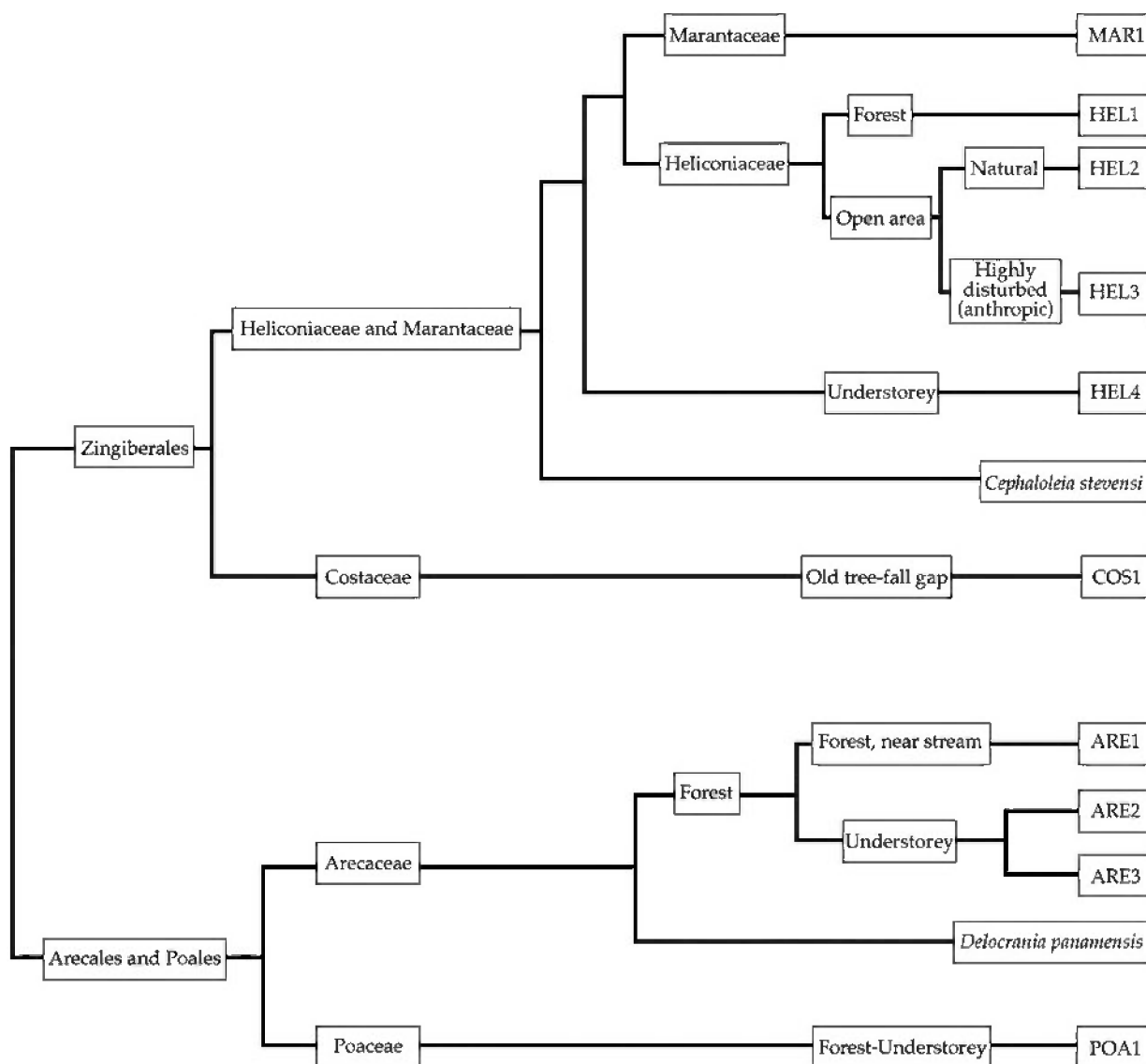


Fig. 3. Synthesis of the relationships between plant family, habitat and hispid Cassidine assemblage composition based on cluster analysis results (average linkage method; Bray–Curtis' distance). ARE1, first cluster group of Arecaceae; ARE2, second cluster group of Arecaceae; ARE3, third cluster group of Arecaceae; HEL1, first cluster group of Heliconiaceae; HEL2, second cluster group of Heliconiaceae; HEL3, third cluster group of Heliconiaceae; HEL4, fourth cluster group of Heliconiaceae; COS1, cluster group of Costaceae; MAR1, cluster group of Marantaceae; POA1, cluster group of Poaceae

beetle assemblages (Tables 3 and 4). Table 3 clarifies the cluster nodes (Fig. 2) according to the host plant species' habitats. However, the fit of some hispine species into these nodes is not clear-cut. For example, *Aslamidium semicircularum*, an 'open-leaf scraper' species that seems to be a 'family specialist' (Barone, 1998; McKenna and Farrell, 2005, 2006), actually occurs on many Marantaceae species (Table 3). This species does not, to our knowledge, feed on other monocot families, including the Arecaceae as indicated by Staines (2002). Conversely, *Cephaloleia sallei* seems to be strongly specialized on

a single plant species: *C. latifolia*, much as *Homalispia nevermanni* occurs on only *Oenocarpus panamanus* among the four palm species we examined. If we follow the feeding concepts of Barone (1998), *C. sallei* and *H. nevermanni* could be termed as 'species specialists'. In our study areas, *C. sallei* is a feeding specialist on primarily one species of Zingiberales (Table 3), contrary to other accounts that suggest it is a generalist (McKenna and Farrell, 2005).

Taxonomy can complicate the comparison and interpretation of results from this study and previous studies. For example, the polymorphic

leaf miner, *Platocthispa emorsitans*, is intimately associated with *H. latispatha*, a common resident in shady anthropically disturbed areas such as roadsides and power transmission line cuts. However, this hispine was also occasionally reared from *Calathea insignis*, *C. latifolia* and *Heliconia irrasa*. Future genetic studies will be needed to test whether individuals reared from *Calathea* and *Heliconia* mines belong to the same or different species. An additional example is provided by *Cephaloleia vicina* Baly and *Cephaloleia belti* whose morphologies are extremely similar and occur on the same species of the Heliconiaceae in different areas. *Cephaloleia belti* occurred in this study primarily on the Heliconiaceae but also in smaller numbers on the Marantaceae. According to a preliminary molecular phylogenetic analysis (Descampe and Meskens unpublished data), the *C. belti* individuals collected in this study appear to belong to a single species.

Another taxonomic problem is presented by the genus *Nympharescus* Weise, which Strong (1982) records from the northern Panama Canal area, but both Blackwelder (1946) and Staines (2002) indicate it is absent from Central America but present from Colombia to Peru. It now appears that this species, which was recorded as a common resident in *Heliconia mariae* scrolls, is a morphological variant of *Chelobasis perplexa* which closely resembles some *Nympharescus* species.

Based on the occurrence data presented in this study and feeding trials (Descampe *et al.*, 2007, in press; Meskens unpublished data), we conclude that most hispines feed on more than one related plant species, often on several genera and families but rarely cross to feed on different orders of monocot plants. The feeding flexibility which we have documented is best exemplified by *C. belti* which occurred in the leaf scrolls of eight species, three genera and two families of Zingiberales.

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