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FACULDADE DE FILOSOFIA, CIÊNCIAS E LETRAS DE RIBEIRÃO PRETO  
PROGRAMA DE PÓS-GRADUAÇÃO EM ENTOMOLOGIA

**Small sparkling flies: systematics of *Pholeomyia* Bilimek, 1867  
(Diptera: Milichiidae: Milichiinae)**

**Pequenas moscas brilhantes: sistemática de *Pholeomyia* Bilimek, 1867 (Diptera:  
Milichiidae: Milichiinae)**

**Heloísa Fernandes Flores**

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## Resumo

Milichiidae (Diptera: Schizophora) é uma família de moscas acaliptradas conhecidas pelo fascinante hábito cleptoparasita da maioria das espécies. Dentro da família, Milichiinae é a subfamília com o maior número de espécies descritas. Os machos de muitas espécies da subfamília apresentam pilosidade prateada dorsalmente no abdômen que os faz “relampejar”, refletindo pequenos raios de luz quando enxameiam no início da manhã. Vários aspectos da sistemática do grupo precisam ser devidamente compreendidos, incluindo questões envolvendo as relações filogenéticas entre os gêneros. Dentro da subfamília, *Pholeomyia* Bilimek, 1867 possui 39 espécies descritas, das quais 27 ocorrem na região Neotropical. O gênero não foi completamente revisto na literatura até o momento, e não há estudos que estabeleçam as relações filogenéticas entre as espécies ou mesmo hipóteses de monofilia do gênero baseadas em sinapomorfias não homoplásticas. Além disso, questões relacionadas a sinonímia de *Pseudomilichia* com *Pholeomyia* ainda permanecem. Aqui, a evolução de Milichiinae é discutida e uma hipótese filogenética de *Pholeomyia* baseada na morfologia de machos é apresentada. A análise filogenética incluiu 72 táxons terminais (57 espécies no grupo interno e 15 do grupo externo) e usou 67 caracteres morfológicos, resultando em uma única árvore mais parcimoniosa sob pesagem implícita. A monofilia de *Pholeomyia* é recuperada e o novo cladograma para a subfamília destaca as relações entre os gêneros de Milichiinae. Nossos resultados também corroboram que *Pseudomilichia* é sinônimo de *Pholeomyia*. O cladograma obtido com 57 espécies do gênero mostra quatro clados principais, cuja posição e composição de espécies podem encontrar algum nível de ajuste com a adição de informações do abdômen masculino na matriz de dados para 12 das espécies incluídas como terminais na análise.

# 1. INTRODUCTION

## 1.1. The family Milichiidae

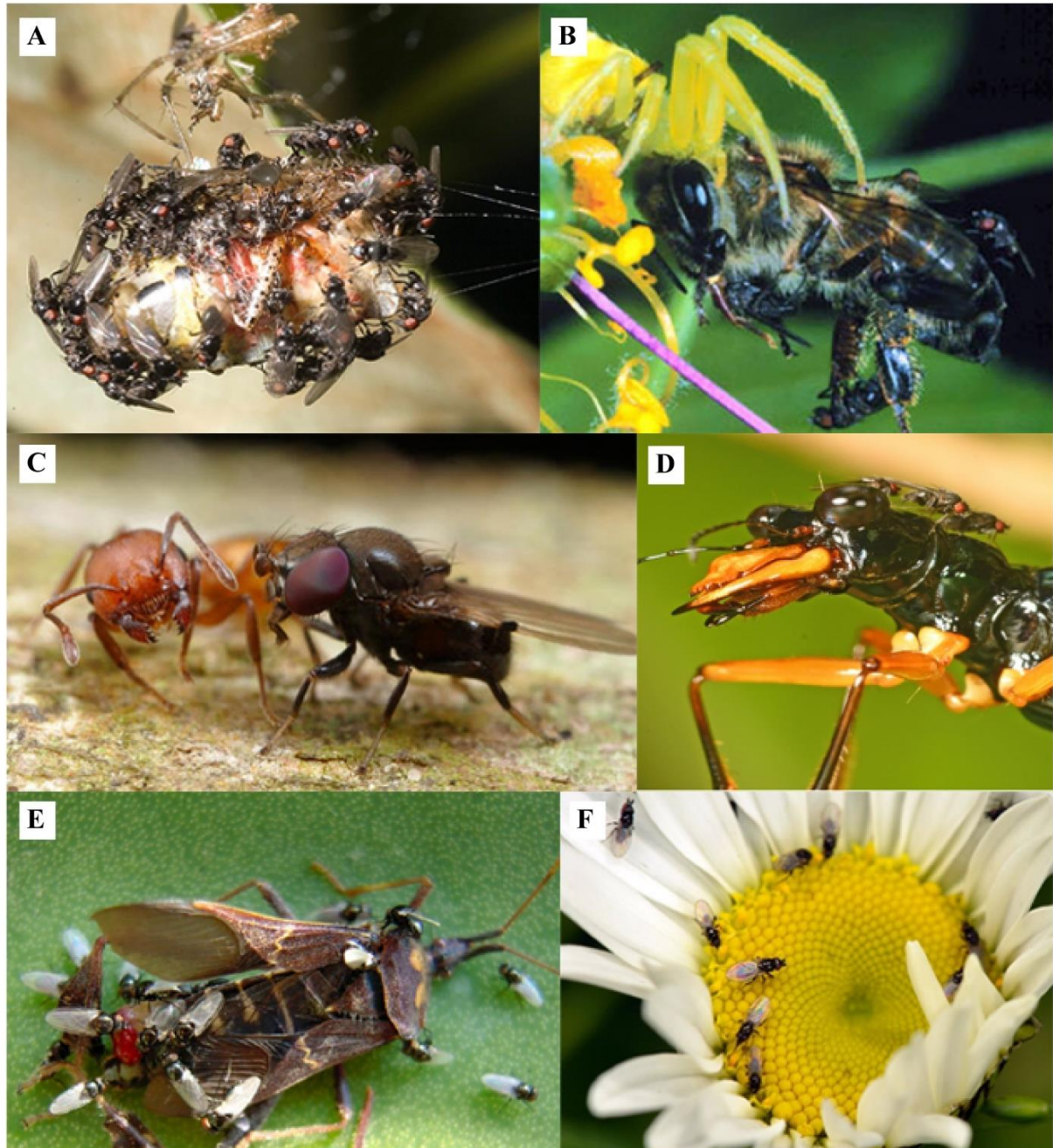
Milichiidae (Diptera: Schizophora), popularly known as ‘jackal flies’ due to the kleptoparasitism habit of most of the species, includes approximately 417 extant and ten fossil species described in 20 genera in the world (Brake, 2000; Swann, 2016). Compared to other fly families, the jackal flies are not particularly species-rich. Nevertheless, considering the undescribed specimens in collections around the world, the number of species to be described may move the diversity of the group to over 1,000.

Milichiids are small flies, ranging in length from 1 to 6 mm. The coloration of most species varies from light brown to black. Males of some genera, as *Pholeomyia* Bilimek, *Milichiella* Giglio-Tos and some others, may have silvery abdominal tergites. The family has a rather broad spectrum of morphological variation, and some species may resemble in some extent other families, as carnids, agromyzids, chloropids and even tachinids (Brake, 2000). This can make it difficult for non-trained entomologists to identify milichiids in collections around the world. Some morphological characters may help in separating the jackal flies from these families, such as the presence of a geniculate proboscis, the presence of both humeral and subcostal breaks, the closed cup cell, and the absence of the postgonites (Brake, 2000; Swann, 2010).

Descriptions of the natural history of the jackal flies demonstrate that they are also ecologically diverse (Brake, 2000) (Figure 1). There are reports of saprophagous, necrophagous and coprophagous larvae (Sabrosky, 1977; Ferrar, 1987; Papp & Wheeler, 1998; Brake, 2000). Coprophagy is a habit present in adults of some species too (Sabrosky, 1959). At least seven genera have been documented to feed on nectar of *Aristolochia* flowers (Brantjes, 1980; Wolda & Sabrosky, 1986).

Within the spectrum of sources of food for adults, one of them stands out: the kleptoparasitism. Kleptoparasitic interactions occur at least in eight genera. Adults feed on the preys of a variety of species of spiders or insects (Reduviidae, Asilidae, Mantidae, among others) (Robinson & Robinson, 1977; Sivinski & Stowe, 1980; Eisner, Eisner & Deyrup, 1991; Sivinski, Marshall & Petersson, 1999; Brake, 2000; Swann, 2008). In almost all cases, only females are kleptoparasites (Sivinski & Stowe, 1980; Eisner, Eisner & Deyrup, 1991) and it is hypothesized that kleptoparasitism provides an extra source of protein for egg maturation (Robinson & Robinson, 1977; Brake, 2000). Kleptoparasitism may be related to the morphological diversity and to the geographical distribution of the family (Brake, 2000),

providing an interesting model for understanding the transition of different life histories across time and space.



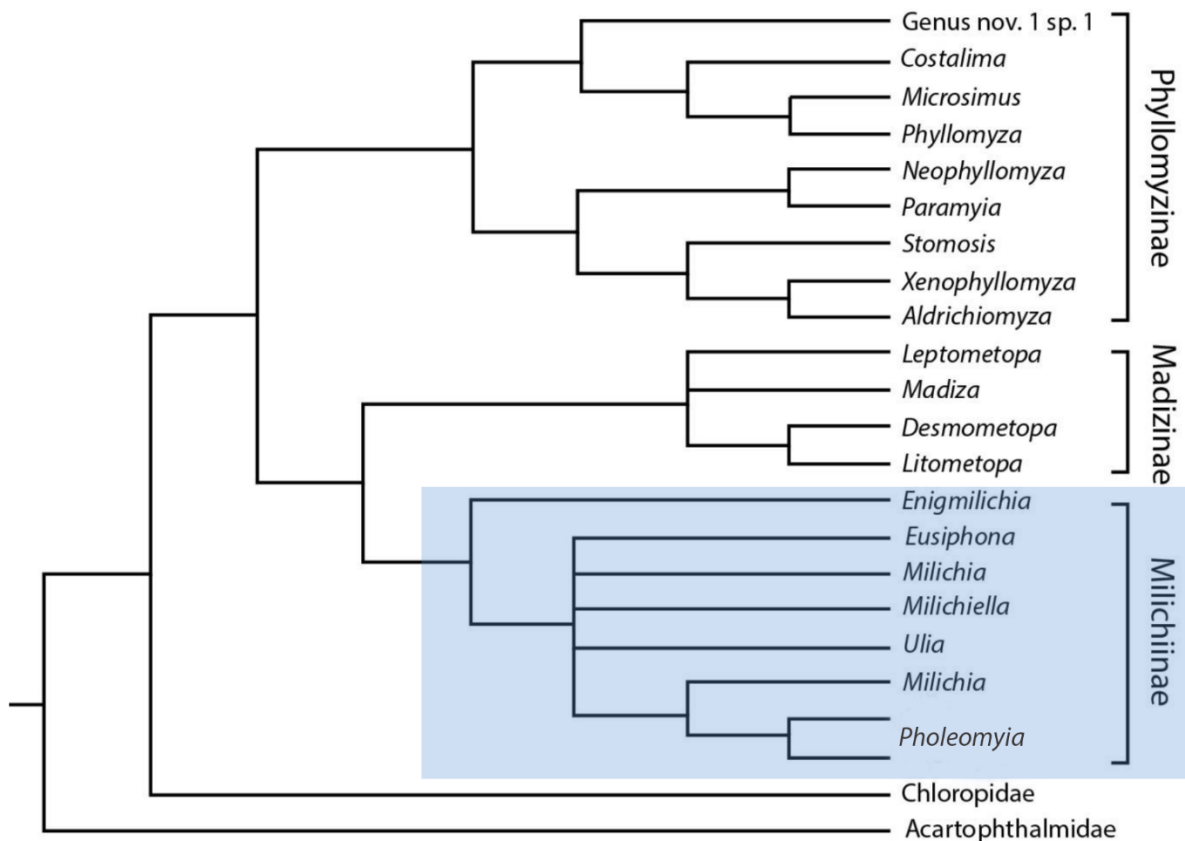
**Figure 1.** Illustrations of some jackal fly biologies. (A) *Paramyia* sp. feeding on the captured stink bug prey of a spider, photo by Steve A. Marshall. (B) *Desmometopa* sp. feeding on the bee prey of a crab spider, photo by Robert Copeland. (C) *Milichia patrizii* Hennig trying to trigger regurgitation of a *Crematogaster* ant, photo by Alex Wild. (D) *Therates labiatus* (Fabricius) with phoretic *Paramyia* sp., photo by Steve A. Marshall. (E) *Milichiella lacteipennis* (Loew) attracted to freshly killed female *Leptoglossus zonatus* (Dallas), photo by Takumasa Kondo. (F) Flowering plant visited by *Paramyia* sp., photo by Steve A. Marshall

It is considerably well established in the literature that Milichiidae is the sister group

of Chloropidae within the Carnoidea. This is supported by some synapomorphies, as the lacinia of maxillae strongly reduced, the proboscis slightly elongated and geniculate, a bare anepisternum, the distiphallus short and glabrous, and the pocket-like ventral receptacle of the female terminalia. There is no dispute that the jackal flies are monophyletic. The most conspicuous synapomorphies of the clade are: upper orbital seta latero-clinate, middle orbital seta latero-clinate, lower orbital seta latero-clinate, two medio-clinate frontal setae, presence of a proclinate setula between supra-antennal seta and eye margin, presence of a pair of setulae on the lunula, presence of four pseudotrachea, and absence of postgonites (Brake, 2000).

In older classifications, the milichiids were divided into the subfamilies Madizinae and Milichiinae. Hennig (1958) pointed out that Madizinae could be paraphyletic. Brake (2000), in fact, recovered in her phylogenetic analysis of the family a paraphyletic Madizinae. She subdivided the Madizinae *sensu lato* into the subfamilies Madizinae and Phyllomyzinae, and recovered Madizinae *sensu stricto* as the sister group of Milichiinae (Figure 2).

Swann (2010) disagreed with the subdivision of Madizinae *s.l.* into two subfamilies. He argued that Brake's (2000) analysis missed information on some characters for several terminals, which ended up as synapomorphies for several Phyllomyzinae taxa. Brake's (2000)



**Figure 2.** Brake's (2000) phylogenetic hypothesis for Milichiidae based on morphological characters.

study is the only formal phylogenetic analysis of the relationships between the milichiid genera. Swann's (2010) comments are helpful as an analysis of the Brake's (2000) study, but are not a formal reanalysis of the group.

Brake (2000) brought a significant advance in the understanding of the evolution of morphological characters in the family, clarifying the phylogenetic relationship between some milichiid genera, but many questions still remain to be solved. Several genera need careful revisions and there are different pending issues on homology in the family, especially of mouthparts and male terminalia sclerites.

## 1.2. The subfamily Milichiinae

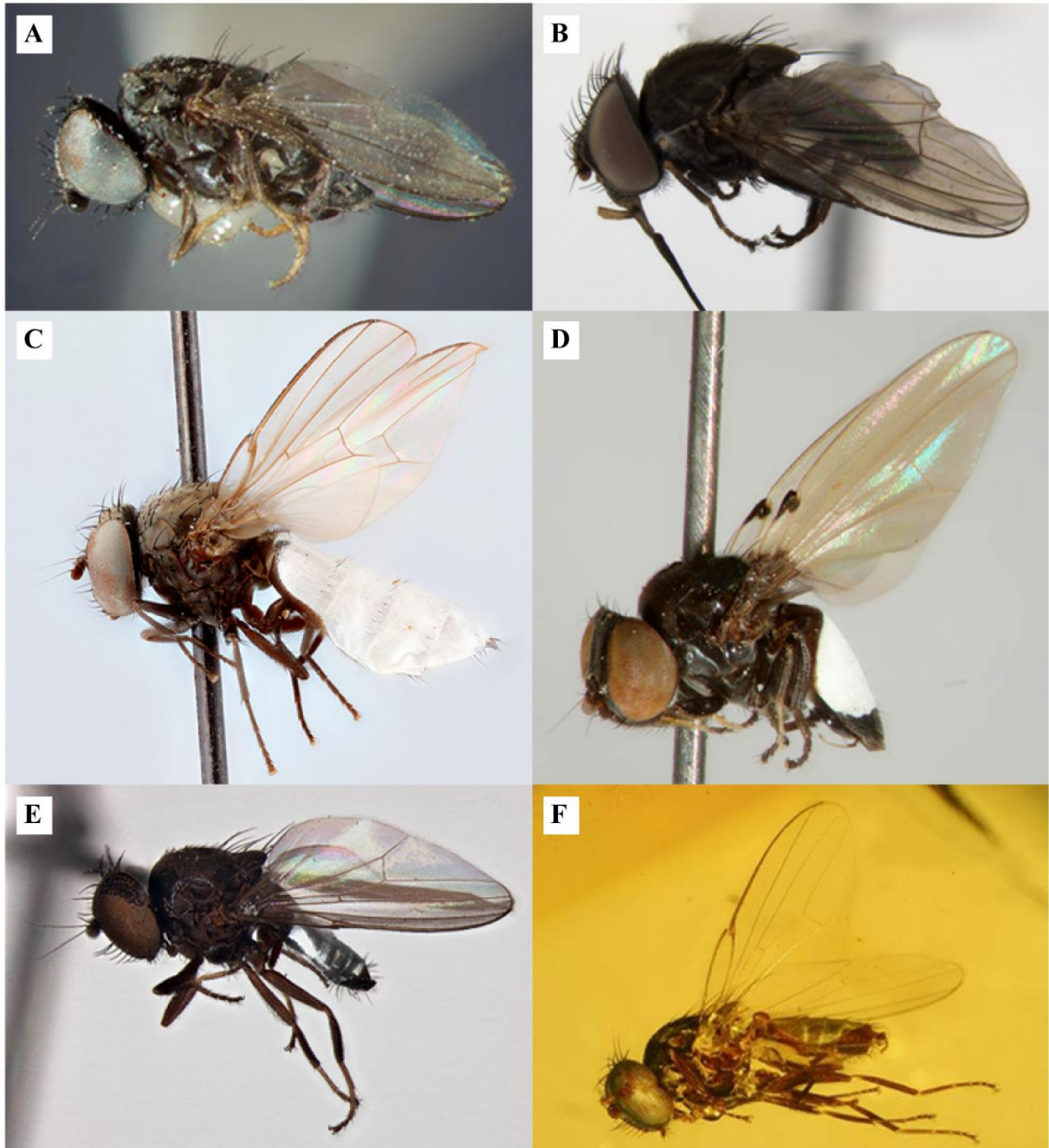
Milichiinae — the milichiid subfamily with largest number of described species — comprises 226 extant species described in five genera in the world (Table 1). It includes the genera *Enigmilichia* Deeming, *Eusiphona* Coquillett, *Milichia* Meigen, *Milichiella* Giglio-Tos and *Pholeomyia* Bilimek (Brake, 2000) (Figure 3). The subfamily also has the largest number of described fossil species: seven *Milichiella* species from Dominican amber (Brake, 2006). In the other subfamilies, only two Phyllomyzinae fossils from Baltic amber and Mexican amber (Hennig, 1967; Sabrosky, 1963) and one Madizinae species from Baltic amber are known (Hennig, 1971).

**Table 1.** Diversity and geographic distribution of extant milichiines. Abbreviations: AF, Afrotropical; AU, Australia; NE, Nearctic; NT, Neotropical; PA, Palearctic; OR, Oriental.

Táxon	Nº of species	AF	AU	NE	NT	PA	OR
<b>Milichiinae</b>	<b>203</b>						
<i>Enigmilichia</i> Deeming, 1981	1	1					
<i>Eusiphona</i> Coquillett, 1897	4			3	1		
<i>Milichia</i> Meigen, 1830	39	21	4			10	6
<i>Milichiella</i> Giglio-Tos, 1895	120	20	14	23	47	6	11
<i>Pholeomyia</i> Bilimek, 1867	39			27	12		

The subfamily is considered monophyletic, a hypothesis supported by the following synapomorphies: enlarged eyes, obsolescent vibrissal angle, vibrissa above lower margin of eye, frons in males narrower than in females, and distal margin of anal cell meeting the anal vein at an acute angle (Brake, 2000). The group has been recovered as monophyletic, but the question of the relationships between the genera and the monophyly of the genera still demand investigation (Figure 2). *Milichia* and *Pholeomyia* are likely to be paraphyletic and the status

of some genera previously described within the subfamily and later synonymized (as *Pseudomilichia* Becker) still remains controversial. It is necessary to increase the taxonomic sampling of all five milichiine genera to properly check their monophyly and address the internal relationships within the subfamily.



**Figure 2.** Representative species of the Milichiinae genera. (A) *Enigmilichia dimorphica* Deeming, Nigeria, photo by Irina Brake. (B) *Eusiphona vittata* Sabrosky, Brazil, photo by Heloísa Flores. (C) *Pholeomyia vockerothi* Sabrosky, United States, photo by Daniel Whitmore. (D) *Milichia formicophila* Deeming, Nigeria, photo by Irina Brake. (E) *Milichiella* sp., Brazil, photo by Heloísa Flores. (F) *Milichiella margaretae* Brake, Dominican Republic amber, photo by Irina Brake.

### 1.3. The genus *Pholeomyia* Bilimek, 1867

According to the world catalog of the family (Brake, 2000), there are 39 species described in *Pholeomyia* (Table 2). The genus is mostly Neotropical, with 11 species known from the Nearctic region. Most *Pholeomyia* species were described by Becker (1907), Hendel (1932), and Sabrosky (1959). The only taxonomic review for the genus in the literature was provided by Sabrosky (1959), which includes all Nearctic and seven of the Neotropical species. Sabrosky (1959) study is the most recent paper to providing descriptions of new species in the genus—no *Pholeomyia* species have been described along the last 60 years.

Adults and larvae of *Pholeomyia* have an interesting biology. Males of most (but not all) of the species of the genus have silvery abdominal tergites, which reflect light while swarming, what allows them to be spotted over long distances (Sabrosky, 1973; Monteith, 1982; Swann, 2010). *Pholeomyia* larvae have been found in nests of *Atta texana*, where they feed on fungal garden debris (Sabrosky, 1959). Larvae of *Pholeomyia* have also been reported in nests of the Megachilidae bee (Sabrosky, 1955).

Brake (2000) recovered *Pholeomyia* as monophyletic in her cladistic analysis of the family, obtaining as synapomorphies the presence of more than one postprotonal setae, the presence of three fronto-orbital setae, and the presence of three or four strong anepisternal setae. Brake (2000) did not include *Pseudomilichia* species as terminal taxa, but synonymized *Pseudomilichia* with *Pholeomyia*. She justified that the three synapomorphies defining *Pholeomyia sensu stricto* would also apply to all *Pseudomilichia* species, and that the emarginate eye of *Pseudomilichia* would be an insufficient character to define the genus.

*Pseudomilichia* was a small genus with only two described species (*Pseudomilichia implicata* Becker, 1907 and *Pseudomilichia schnusei* Becker, 1907), defined mainly by the combination of the presence of an emarginate eye and the presence of three or four strong anepisternal setae. The genus had already been synonymized with *Pholeomyia* by Hendel (1932), but his nomenclatural change was ignored by further authors—e.g., Hennig (1939). Instead, Hennig (1939) synonymized *Macromilichia* Hendel with *Pseudomilichia*. He also suggested that *Macromilichia nigricosta* Hendel, 1932 might be synonymous with *Pseudomilichia schnusei*, a pair of species that Brake (2000) accepted as valid. In the same study, Hennig synonymized *Rhynchomilichia* Hendel with *Pholeomyia*. After the synonymy proposed by Brake (2000), all *Macromilichia*, *Rhynchomilichia* and *Pseudomilichia* species have been included in *Pholeomyia*. It seems well established that *Macromilichia* and



*Rhynchomilichia* would be synonymous of *Pholeomyia*, but questions about *Pseudomilichia* still remain.

Swann (2010) disagreed from Brake's (2000) synonymy and suggested that her synapomorphies for *Pholeomyia* would be homoplastic features evolving inside and outside Milichiinae. He mentioned, for example, that the presence of three or four strong anepisternal setae is also known to occur in some *Eusiphona* species. Following Swann (2000) and Sabrosky (1955), the presence of anepisternal setae may indicate a close relationship between *Eusiphona* and *Pholeomyia*, and even unite *Eusiphona*, *Pholeomyia*, and *Pseudomilichia*. Swann (2010) also points out that the presence of three frontal setae, considered by Brake (2000) as a synapomorphy of the genus, actually corresponds to a range of three to six frontal setae in *Pholeomyia sensu stricto*. Although it is not a unique condition within the subfamily, Swann (2010) considered *Pseudomilichia* a valid genus and the emarginated eye would be one of its defining features.

Brake's (2000, 2009) studies were a major step forward towards resolving the phylogenetic relationships within Milichiinae. Brake (2000) recovered *Pholeomyia* as sister of one of the branches of a paraphyletic *Milichia*. In Brake's (2009) taxonomic review and cladistics analysis of *Milichiella*, the *Pholeomyia* species included was recovered as sister group of *Milichiella*. As both analyses had limited number of species of *Pholeomyia*, a new cladistic reanalysis of the group with a wider taxonomic sampling may bring some shift to the conclusions on the monophyly of the genus and to its position in the system of the Milichiinae.

To properly address Brake's (2000) inferences for *Pholeomyia*, hence, it is necessary to broaden the sampling of species of the genus and properly sampling other Milichiinae genera to root the analysis and test the monophyly of *Pholeomyia*. Also, a detailed study of the male terminalia sclerites would highlight several unsolved questions in the evolution of the terminalia in milichiids. Finally, advances in the understanding of the phylogenetic relationships of the jackal flies shall provide an important background to future studies of evolutionary biology investigating the fascinating life histories of the family.

**Table 2.** Described species of *Pholeomyia* Bilimek, with the corresponding depository institutions of their types. Acronyms for institutions are explained in Material and Methods. Abbreviations: HT, holotype; LT, lectotype, NE, Nearctic; NT, Neotropical; ST, syntype; ♀, female; ♂, male.

Species	Previous generic assignment	HT	LT	ST	♂	♀	Institution	Geographical Distribution
<i>P. aequatorialis</i> Seguy, 1934	-			X	X		MNHNP	NT: Ecuador
<i>P. anomala</i> Hendel, 1933	-	X			X		NMW	NT: Brazil
<i>P. anthracina</i> (Becker, 1907)	<i>Rhynchoimitichia</i> Becker, 1907	X				X	HNHM	NT: Paraguay
<i>P. argyrata</i> Hendel, 1932	-	X			X		NMW	NT: Argentina
<i>P. argyrophenga</i> (Schiner, 1868)	<i>Lobioptera</i> Wahlberg, 1847 <i>Rhynchoimitichia</i> Becker, 1907	X			X		HNHM	NT: Bolivia, Peru
<i>P. comans</i> Sabrosky, 1959	-	X			X		USNM	NE: United States
<i>P. dampff</i> Sabrosky, 1959	-	X			X		USNM	NT: Guatemala, Mexico
<i>P. decorior</i> Steyskal, 1943	-	X			X		USNM	NE: United States
<i>P. dispar</i> (Becker, 1907)	<i>Rhynchoimitichia</i> Becker, 1907			X	X		Coll. Bezzi	NE: United States
<i>P. excelstor</i> (Becker, 1907)	<i>Rhynchoimitichia</i> Becker, 1907			X	X		USNM	NT: Bolivia
<i>P. expansa</i> Aldrich, 1925	-	X			X		USNM	NE: United States
<i>P. fasciventris</i> (Becker, 1907)	<i>Rhynchoimitichia</i> Becker, 1907			X	X		USNM	NT: Bolivia
<i>P. hurdi</i> Sabrosky, 1959	-	X			X		CAS	NT: Mexico
<i>P. implicata</i> (Becker, 1907)	<i>Pseudomitichia</i> Becker, 1907	X			X		USNM	NT: Peru
<i>P. indecora</i> (Loew, 1869)	<i>Lobioptera</i> Wahlberg, 1847 <i>Mitichia</i> Meigen, 1830 <i>Rhynchoimitichia</i> Becker, 1907	X				X	MCZ	NE: United States
<i>P. insecta</i> (Becker, 1907)	<i>Rhynchoimitichia</i> Becker, 1907			X	X		USNM	NT: Bolivia
<i>P. latifrons</i> Sabrosky, 1959	-	X			X		AMNH	NT: Bahamas
<i>P. leucogastra</i> (Loew, 1861)	<i>Lobioptera</i> Wahlberg, 1847 <i>Mitichia</i> Meigen, 1830 <i>Rhynchoimitichia</i> Becker, 1907	X			X		MCZ	NT: Cuba, Mexico

Ativa

Table 2. Continued.

<i>P. leucozoma</i> Blimek, 1867	-	X	X	NMW	NT: Mexico
<i>P. longifacies</i> Hendel, 1933	-	X	X	NMW	NT: Brazil, Paraguay
<i>P. longiseta</i> (Becker, 1907)	<i>Mitichia</i> Meigen, 1830	X	X	HNHM	NT: Argentina, Paraguay, Nicaragua
<i>P. myopa</i> Melander, 1913	-	X	X	USNM	NT: Bolivia, Brazil, Costa Rica, Mexico, Panama, Venezuela
<i>P. nigricosta</i> (Hendel, 1932)	<i>Macromitichia</i> Hendel, 1932	X	X	SMNS	NT: Bolivia
<i>P. nitidula</i> Sabrosky, 1959	<i>Pseudomitichia</i> Becker, 1907	X	X	USNM	NE: United States
<i>P. obscura</i> Sabrosky, 1959	-	X	X	USNM	NE: United States
<i>P. palparis</i> (Becker, 1907)	<i>Rhynchomitichia</i> Becker, 1907	X	X	HNHM	NT: Paraguay
<i>P. pectoralis</i> Hendel, 1932	-	X	X	SMNS	NT: Bolivia
<i>P. politifacies</i> Sabrosky, 1959	-	X	X	USNM	NT: Brazil, Colombia, Costa Rica, El Salvador, Mexico, Panama
<i>P. praeocellaris</i> Hendel, 1932	-	X	X	SMNS	NT: Bolivia
<i>P. praesepta</i> (Becker, 1907)	<i>Rhynchomitichia</i> Becker, 1907	X	X	USNM	NT: Peru
<i>P. prominens</i> (Becker, 1907)	<i>Rhynchomitichia</i> Becker, 1907	X	X	USNM	NT: Bolivia, Peru
<i>P. pseudodecora</i> (Becker, 1907)	<i>Rhynchomitichia</i> Becker, 1907	X	X	Coll. Bezzi	NE: United States
<i>P. quadrifasciata</i> Hendel, 1932	-	X	X	SMNS	NT: Bolivia
<i>P. robertsoni</i> (Coquillett, 1902)	<i>Mitichia</i> Meigen, 1830	X	X	USNM	NE: United States
<i>P. schineri</i> (Hendel, 1932)	<i>Rhynchomitichia</i> Becker, 1907	X	X	NMW	NT: Argentina, Brazil, Paraguay
<i>P. schunzei</i> (Becker, 1907)	<i>Pseudomitichia</i> Becker, 1907	X	X	USNM	NT: Bolivia, Peru
<i>P. sororcula</i> (Becker, 1907)	<i>Rhynchomitichia</i> Becker, 1907	X	X	HNHM	NT: Bolivia, Peru
<i>P. texensis</i> Sabrosky, 1959	-	X	X	USNM	NE: United States
<i>P. vockerothi</i> Sabrosky, 1961	-	X	X	CNC	NE: United States

## 6. CONCLUSIONS

This study addressed the systematics of *Pholeomyia*, one of the most distinctive jackal fly genera, with a phylogenetic hypothesis for the genus. The taxonomic sampling of analysis was considerably extensive, allowing to further discuss the monophyly of *Pholeomyia*, the relationships within the genus, and the relationships between milichiine species.

Our results corroborate the hypothesis of a monophyletic *Pholeomyia* including the species assigned to *Pseudomilichia*. The sampling of the genus *Milichia* also reinforces the idea that this genus is paraphyletic in relation to *Milichiella* and to *Pholeomyia*. Our analysis increased the resolution of the relationships within Milichiinae. This study also highlights the gap in the knowledge of *Pholeomyia* diversity, with up to 30 potentially undescribed species that would add to the 28 described Neotropical species. The phylogeny obtained with 57 species of the genus show four main clades, which position and species composition may find some level of adjustment with the addition of male abdomen information in the data matrix for 12 of the species included as terminals in the analysis.

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