

Article

Updated Taxonomic Key of European Nycteribiidae (Diptera), with a Host-Parasite Network

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Abstract: Current taxonomical keys are essential for better biodiversity knowledge. Moreover, accurate identification of groups of organisms known to act as disease vectors is vital in today's world of change and the spread of zoonotic disease threats. This is especially relevant to bats and their parasites, given the events of recent years. The available keys of European Nycteribiidae (Diptera) are outdated and do not cover all the species currently known from Europe. Therefore, a summary key of 16 European species is provided in this paper. Based on published data from Europe, a total of 173 host-parasite interactions were observed between 16 European bat fly species and 31 host species (bats). The highest number of associations with different host species and the lowest specialization was observed in *Penicillidia (P.) dufourii dufourii*, followed by *Nycteribia (N.) kolenatii*, *N. (N.) schmidlii* and *Phthiridium biarticulatum*; most of the other species also had low specialization. Most species are oligoxenous, parasitizing on several families, and three species are stenoxenosus. Only one nycteribiid species, *Basilia daganaiae*, was linked with only one host (*Pipistrellus kuhlii*). *Myotis myotis* has 12 associations with different bat fly species, followed by *Plecotus auritus* and *Myotis daubentonii* with 12 associations. A relatively high number of bat species were associated with only one bat fly species. In addition, we performed a bipartite analysis complemented by network indices and host specificity at the species and whole network levels, yielding new information in terms of ecology and the host-parasite relationships related to these groups of potential vectors (Nycteribiidae) and sources of zoonoses (bats). The parasite-host network composed of 16 nycteribiid species and 31 bat host species was characterized by a low network level specialization, low modularity, and low nestedness. Our findings also suggest a high probability of associated Nycteribiidae in bat taxa with a similar roosting ecology and phylogenetic relationship.



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1. Introduction

Bats (Mammalia: Chiroptera) are the second most numerous mammalian group after rodents, with more than 1430 species (<https://batnames.org/>; accessed 19 March 2023) [1], and they provide crucial ecosystem services as seed dispersers, pollinators, controllers of insects, and nutrient recyclers [2]. On the other hand, bats are also considered to be natural reservoirs of a wide range of microorganisms, especially viruses [3]. The fact that they are reservoirs for a staggering number of viruses and other potential pathogens is due to their biology, ecology, and behavior (e.g., diversity of food, colonial sociality, mobility, and migration, etc.) [4]. Although bats on average harbor a higher diversity of parasites and pathogens than any other mammalian group, the likelihood of bat flies vectoring disease agents across host species of bats may be relatively small [5].

Many ectoparasites, such as acarines (ticks, chiggers, and mites), bat flies, and fleas, specialize in bats as hosts. Bat flies (Diptera: Nycteribiidae) form a fascinating and largely

disregarded group of obligate, blood-feeding, highly specialized ectoparasites of bats [6,7]. This family is included in the superfamily Hippoboscoidea, along with the families Hippoboscidae (keds or louse flies), Glossinidae (tse-tse flies), and Streblidae [8]. Phylogenetic studies suggest monophyly among Hippoboscoidea members and that the ancestor of this superfamily was a free-living insect feeding on mammals' blood [9,10]. One of the most interesting characteristics of bat flies is that they are obligate parasites of bats [11]. Their ecological origins are obscure, but Jobling [12] envisioned an ancestral association with caves and guano accumulations before their switch to an ectoparasitic lifestyle. Bat fly morphology reflects their role as ectoparasites. The wings of many species are strongly reduced or absent, and long limbs facilitate locomotion in the host body. Nycteribiids possess several ctenidia or combs. Various theories have been published on the function of ctenidia. One theory claims that the ctenidia help them to hold onto the fur and wing membrane [13,14]. Observations by other authors suggest that ctenidia function not as organs of attachment, but rather to protect highly mobile joints and their associated membranes, which are not otherwise protected [15]. One of the most consistent morphological features of bat flies is the rudimentary manifestation of their visual system [16]. Their eyes are completely missing or reduced to relatively few facets and ocelli are absent [17]. Worldwide, three subfamilies (Archinycteribiinae Maa, 1975, Cyclopodiinae Maa, 1965, Nycteribiinae Westwood, 1835), 11 genera, and more than 276 nycteribiid species are known [18]. In Europe, only 16 species from the subfamily Nycteribiinae have been recorded [19–21].

Adult bat flies of both sexes spend their lives on bats where they can be observed in the fur or on the wing membranes. Bat flies reproduce via viviparous puparity [16]. Eggs are internally fertilized and all larval stages develop within the females [22]. Larvae molt twice inside the female, and gravid females deposit a single terminal (3rd-instar) larva on the roosting substrate. Once deposited, the larva (referred to as a prepupa) immediately forms a puparium. The pupal stage lasts 3–4 weeks [23]. Bat flies deposit their larva on substrates, such as the host roost wall. After larviposition, females return to their host. When the offspring emerge, they actively search for bat hosts [24].

On the behavioral side, bat flies have been observed to react to sudden strong light flashes. Upon exposure, they consistently respond with a startle reflex resulting in flight. Bat flies are covered in setae, many of which seem to function as mechanoreceptors, aiding in the sensation of air currents or vibrations. Little is known about the olfactory faculties and preferences of bat flies, but presumably, olfactory cues are important in host finding. Some bat flies exhibit strong plumose modification of the arista, which in other Diptera is known to contain thermo- and hygroreceptors [25].

As obligate blood-feeding parasites, bat flies seem to be excellent candidates for vectors of zoonotic agents, and recent studies have shown that bat flies could carry pathogens, such as bacteria and viruses, and transmit them among bat individuals in a colony [1,2,26–33]. The likelihood of disease transmission by bat flies to distantly related mammals, such as humans, is considered to be relatively small [5]. Further, interspecies transmission via bat flies is unlikely under normal conditions, because Nycteribiids are usually highly specific parasites, with each species of bat fly parasitizing a single bat species or genus [15] or rarely, families, as in the case of *Penicillidia fulvida* [34]. Bats occasionally form species-specific groups in colonies [35], which may lead to low ectoparasite overlap between bat species. However, they very often form mixed species colonies [36].

Although identification and knowledge of the ecology and biology of bats are at a high level, this is not the case for some of their parasites (such as Nycteribiidae) [37].

Moreover, apart from the epidemiological perspective, the study of host-parasite interactions is an opportunity to understand the ecological relationships between organisms. Parasites, together with their hosts, constitute an ideal system for investigating the patterns and mechanisms of community structure and dynamics [38]. Analyses of the relations between parasites and hosts are possible using so-called ecological network analysis, which is a representation of interactions within the studied ecosystems [39–41]. Ecological networks can inform us about the evolutionary and ecological processes that generate and

shape biodiversity. The structure of ecological networks can also provide information on the vulnerability of ecological communities to various kinds of disturbances: from climate change to the illegal hunting of keystone species and extinctions of species-to-species invasions [42], or they can help to better understand the phoretic relationships between arthropods [43]. Ecological networks and host–parasite associations remain understudied in bats and their parasites, including nycteribiids. Despite growing knowledge on diversity and bat-Nycteribiidae associations [35,38,44–50] and the existence of literary records on the host species of bat flies from Europe [7,51,52], many gaps need to be filled.

2. Materials and Methods

2.1. Updating of Identification Key

Using the records of Szentivanyi et al. [21] as a starting point, we checked the literature on the host associations of European bat louse for new, overlooked, or misattributed records [53–142]. In addition, we conducted a literature search using the keywords ‘Nycteribiidae*’, ‘European bat flies*’, ‘host associations *’, ‘bats *’, and ‘Europe*’ in Scopus and Google Scholar [7,51,52,143–154]. Figures are redrawn and modified based on the original illustration by Theodor (1954) [155] (pp. 66 a. Nycteribiidae. Taf. I–IV, VI–VII, IX, XIII–XVI); Theodor (1967) [57] (pp. 1–506, figures compared with [50]); Hůrka (1970) [53]. (p. 243: Figure 1); Hůrka (1972) [54] (p. 711: Figure 1); Beaucournu and Noblet (1685) [68] (p. 637: Figure 1).

2.2. Host Specificity, Bipartite Networks, and Statistics

For the host associations of European bat flies, we used information published by Szentivanyi et al. [21] and the recent literature [7,51,52,143–154]. The terminology for the host specificity of particular bat fly species follows [156,157].

To illustrate the parasite–host relationship between bat flies and bats, we used a bipartite network. The boxes of the bipartite graph represent species, and their width represents the summary of interactions among these species. The number of interactions is shown by lines linking the species, and the number of lines means the number of species interactions. To do this, we used a function visualizing a bipartite interaction named matrixplotweb, through which a two-dimensional matrix is plotted as a bipartite graph [158]. Another function of the bipartite package for R—plotModuleWeb takes an object of class in moduleWeb and plots the modules found by function computeModules(...) function into the graph [159].

To carry out the analyses of patterns in the host–parasite network, we used the bipartite package (version 2.18) for R software [160]. For quantitative indices, the number of bat fly species infesting each bat species was used. We prepared matrices with bat species (host) in the columns and bat flies in the rows. In the next step, we calculated these bipartite indexes: connectance (C), network specialization (H_2'), nestedness (N), modularity (Q), and species' specialization metrics (d').

Connectance (C) is one of the most popular and earliest metrics proposed to characterize species interaction networks: it is defined as the proportion of realized interactions from the pool of all possible interactions between the species of a network [161]. The Networklevel function of the bipartite package analyzes bipartite webs at the level of the entire network and calculates a variety of indices and values for a bipartite network, including connectance [162].

The index of network specialization (H_2') reflects the deviation of a species' realized number of interactions and the expected number from each species' total quantity of interactions [163]. The value of this index varies from 0 to 1, where 0 suggests high specialization and 1 implies low specialization. To calculate this index, we used the H2fun function of the bipartite package with two arguments: web, which is a matrix representing the interactions observed between higher trophic level species in columns and lower trophic level species in rows, and H2_intriger, which is a logical argument indicating whether web entries are integer numbers [164].

Nestedness (N) is a measure of the structure of an ecological system and expresses how many interactions realized by specialists are a subset of those performed by generalists. The unit of nestedness is the nestedness temperature T , ranging from 0 to 100°. However, in this study, we used a binary system, where metrics define $N = (100 - T)/100$. It measures the departure from a perfectly nested interaction matrix [165]. In the range of 0–1, value 1 implies maximum nestedness [166–168]. To compute this index, we used a function of the bipartite package named nestedness, which calculates matrix temperature using the BinMatNest approach of Miguel Rodriguez-Girones [169].

Modules are aggregated sets of species that interact. Their defining feature is that interactions within a module are more common than interactions between modules [170–172]. Modularity (Q) is calculated as likelihood, which is implemented in the bipartite computeModules function. This function takes a bipartite weighted graph and computes modules by applying Newman's modularity measure in a bipartite weighted version to it [173].

The last index we calculated was species' specialization metrics (d'). This index measures interactions at the species level and refers to each bat fly species related to bats in the network [174]. To determine this index, the function dfun was used, which returns the specialization index d' for the lower level, which expresses how specialized the species is in relation to what higher-level partners are on offer. The mentioned function has two arguments: web, which is a matrix representing the interactions observed between higher level species (columns) and lower-level species (rows), and abuns, which is a vector of abundance for the higher level, usually from independent information [175]. For a more detailed description of these methods, see [157].

3. Results

(Figures 1–5)

Identification Key for European genera of Nycteribiidae

1 eyes absent (Figure 1(1,4)) ... 2

– eyes present (Figure 1(2,3)) ... 4

2 tibiae longer and narrower, 4.5 to 5 times longer than wide (Figure 1(6)) ... *Phthiridium biarticulatum* Hermann, 1804

– tibiae short, curved ventrally ... 3

3 tibiae nearly semicircular, 2 to 2.5 times longer than wide (Figure 1(5)) ... *Nycteribia* (*Nycteribia*)

– tibiae more slender, 3.5 times as long as wide ... *Nycteribia* (*Acrocholoidia*) *vexata* Westwood, 1835

4 eyes pigmented, with at least 2 lenses, specimens to 3 mm ... *Basilia*

– unpigmented eyes with one lens (Figure 1(2,3)), specimens from 3.5 to 4 mm ... *Penicillidia*

Key for sex determination

1 ventral abdomen with claspers (Figure 1(8)) ... males

– ventral abdomen without claspers (Figure 2(15–20); Figure 3(27–30)) ... females

Keys to species of European genera of Nycteribiidae

Genus *Basilia* Miranda Ribeiro, 1903

Males

1 sternite 5 with one row of spines ... 2

– sternite 5 with two or more rows of spines ... 4

2 sternite 5 with 6–8 spines with convex posterior margin (Figure 1(9)) ... *Basilia nattereri* Kolenati, 1857

– sternite 5 with more than 8 spines ... 3

3 sternite 5 with 8–10 short spines (Figure 1(10)) ... *Basilia nana* Theodor & Moscona, 1954

– sternite 5 with 11–13 spines with convex posterior margin (Figure 1(11)) ... *Basilia mediterranea* Hurka, 1970

4 sternite 5 convex, 4 times wider than long, with a group of about 20–25 soft spines in 2 rows in the middle of posterior margin ... 5

– sternite 5 rectangular, 3 times wider than long, with a group of about 15 spines in a double row (Figure 1(12)) ... *Basilia daganaiae* Theodor & Moscona, 1954

5 sternite 5 significantly convex, with a group of about 20 soft spines in 2 rows in the middle of the posterior margin (Figure 1(13)) ... *Basilia italicica* Theodor, 1954

– sternite 5 slightly convex, with a group of about 25 soft spines in 2 rows in the middle of the posterior margin (Figure 1(14)) ... *Basilia mongolensis nudior* Hurka, 1972

Females

1 tergal plate 2 trapezoidal ... 3

– tergal plate 2 parallel ... 2

2 tergal plate 2 transversely rectangular, lower edge 2 times narrower than upper (Figure 2(15))
... *Basilia nattereri* Kolenati, 1857

– tergal plate 2 square (Figure 2(16)) ... *Basilia nana* Theodor & Moscona, 1954

3 posterior margin 2 times narrower than upper, with a row of 18–22 setae at the posterior margin which has a small indentation in the middle (Figure 2(17)) ... *Basilia italicica* Theodor, 1954

– the posterior margin 3 times narrower than upper ... 4

4 tergal plate 2 with 6 long setae and a second row of small setae at the posterior margin which has a big indentation in the middle (Figure 2(18)) ... *Basilia daganiae* Theodor & Moscona, 1954

– tergal plate 2 with one row of 6 or fewer long setae in the middle of the posterior margin, which has an indentation in the middle ...

5 tergal plate 2 with 6 long setae in the middle of the posterior margin, which has a strong indentation in the middle (Figure 2(19)) ... *Basilia mediterranea* Hurka, 1970

– tergal plate 2 with 4 long setae in the middle of the posterior margin, which has a strong indentation in the middle (Figure 2(20)) ... *Basilia mongolensis nudior* Hurka, 1972

Genus *Nycteribia* Latreille, 1796

Subgenus *Acrocholidia* Kolenati, 1857—one species only

Nycteribia (Acrocholidia) vexata Westwood, 1835

Male—sternite 5 with a row of 8–10 strong spines of concave posterior margin (Figure 3(21)). Aedeagus with a rounded end, which is as wide as the base, narrower in the middle. Parameres slender, triangular, three-quarters the length of the aedeagus.

Female—genital plate consisting of a strongly sclerotised arc with about 4 (to 7) short spines. Tergite 1 with a row of long setae posteriorly (Figure 3(22)).

Subgenus *Nycteribia* Latreille, 1796

Males

1 sternite 5 three to 4 times wider than long ... 2

– sternite 5 two times wider than long, strongly convex posteriorly, with a double row of 14–16 short, thick spines (Figure 3(23)). Anal segment very long, nearly parallel-sided. Aedeagus with a long bifid tip which is curved backwards and has scales at the base of the dorsal membrane ... *Nycteribia (Nycteribia) schmidlii* Schiner, 1853

2 sternite 5 with a row of 14–18 spines in the middle of the concave posterior margin (Figure 3(24)). Phallobase conical, aedeagus conical with a ventral tooth near the tip ... *Nycteribia (Nycteribia) latreillii* (Leach, 1817)

– sternite 5 with a row of 9–12 short spines in the middle of the convex posterior margin (Figure 3(25)). Phallobase with a marked dorsal bulge. Aedeagus wide and short, with a rounded end and a ventral tooth at about the distal third ... *Nycteribia (Nycteribia) pedicularia* Latreille, 1805

3 sternite 5 with a row of 7–8 spines in the middle of the row, posterior margin (Figure 3(26)). Phallobase conical, without the dorsal bulge. Aedeagus slender, with a ventral tooth at the distal fifth ... *Nycteribia (Nycteribia) kolenatii* Theodor & Moscona, 1954

Females

1 dorsal sternite 5 with 2 small lateral plates, with 2 long setae posteriorly. Sternite 6 with 2 lateral plates narrowed toward the centre. Sternite 7 with plate as long as the width of the plates on sternite 6. Ventral genital plate wide. Tergite 2 of abdomen about as long or longer than the width of tergite 1, strongly convex posteriorly. Dorsal genital plate triangular with

truncated sides and 11–15 strong spines (Figure 4(27)) ... *Nycteribia (Nycteribia) latreillii (Leach, 1817)*

– dorsal sternite 5 with 2 lateral plates, with 5 and more long setae ... 2

2 dorsal sternite 5 with 2 lateral plates with 7 to 9 long setae. Sternite 6 with 2 lateral plates touching each other. Sternite 7 with plate as long as two-thirds of the plates on sternite 6. Ventral genital plate narrow. Tergite 2 similar to that of *N. latreillii* or pointed posteriorly. Dorsal genital plate triangular, with 8–10 long setae (Figure 4(28)) ... *Nycteribia (Nycteribia) kolenatii Theodor & Moscona, 1954*

– dorsal sternite 5 with 2 lateral plates, with 5–7 long setae centrally. Sternite 6 with 2 lateral plates not touching each other. Sternite 7 with plate longer than two-thirds of the plates on sternite 6, concave at front. Ventral genital plate narrow, length 2.5 mm. Tergite 2 shorter in the middle than the width of tergite 1. Dorsal genital plate triangular with a row of about 12 short spines (Figure 4(29)) ... *Nycteribia (Nycteribia) pedicularia Latreille, 1805*

3 dorsal sternite 5 with 2 lateral plates, with 15–17 long setae in two lines. Sternite 6 with 2 lateral plates not touching each other. Sternite 7 with plate longer than two-thirds of the plates on sternite 6, concave at front. Genital plate elliptical with a row of very short spines posteriorly. Tergite 3 about three-quarters as wide as tergite 2, surface covered with short spines (Figure 4(30)) ... *Nycteribia (Nycteribia) schmidlii Schiner, 1853*

Genus Penicillidia Kolenati, 1863

Subgenus Penicillidia Kolenati, 1863

Males

1 posterior angle of sternite 5 pronounced. About 15 spines in 1–2 rows at each side of the posterior margin (Figure 4(31)). Apex of the triangle bare. Surface of the sternite bare except for groups of setae at the lateral corners. Aedeagus with an apical dorsal tooth ...

Penicillidia (Penicillidia) conspicua Speiser, 1901

– posterior angle of sternite 5 not pronounced ... 2

2 sternite 5 with two lateral broad, rounded processes covered with numerous short, thick spines and with only a median group of spines between the lateral processes (Figure 4(32)). Frons not prolonged (Figure 1(2)) ... *Penicillidia (Penicillidia) dufourii dufourii (Westwood, 1835)*

– sternite 5 with two shorter lateral broad, rounded processes covered with numerous short, thick spines and with a group of spines between the lateral processes (Figure 4(33)). Frons prolonged into a long horn (Figure 1(3)) ... *Penicillidia (Penicillidia) monoceros Speiser, 1900*

Females

1 tergite 1 triangular, with a few short spines at the apex. Tergal plate 2 large, rounded, with a marginal row of about 20 long setae and covered with short hairs on the surface (Figure 5(34)). Sclerotised processes lacking. Genital plate with an angular row of 10–14 setae ... *Penicillidia (Penicillidia) conspicua Speiser, 1901*

– tergite 1 rounded posteriorly ... 2

2 tergite 1 with a row of short, thick setae with a gap in the middle. Tergal plate 2 rectangular, bare on the surface, with 2 groups of long spines at the posterior margin. Two sclerotised processes with a sharp posterior edge on the connexivum behind tergal plate 2 (Figure 5(35)). Genital plate small, rounded, with 6–8 setae. Frons not prolonged (Figure 1(2)) ... *Penicillidia (Penicillidia) dufourii dufourii (Westwood, 1835)*

– tergite 1 with a row of short, thick setae with a gap in the middle. Tergal plate 2 rectangular, bare on the surface, with 2 groups of long spines at the posterior margin. Two sclerotised processes with a sharp posterior edge on the connexivum behind tergal plate 2. Genital plate small, rounded, with 6–8 setae. Frons prolonged into a long horn (Figure 1(3)) ... *Penicillidia (Penicillidia) monoceros Speiser, 1900*

Genus Phthiridium Hermann, 1804

Males

1 sternite 5 four times wider than long, with a row of 15–18 strong spines in the middle and an upper row with soft spines of strong convex posterior margin (Figure 5(36)) ...

Phthiridium biarticulatum Hermann, 1804

– sternite 5 three times wider than long, with a row of about 25 small spines in 1–2 rows of slightly convex posterior margin (Figure 5(37))... *Phthiridium integrum* (Theodor and Moscona, 1954)

Females

1 tergite 2 very long, covering about half the abdomen. Tergite 6 very large, triangular, with some short setae on the surface (Figure 5(38)) ... *Phthiridium biarticulatum* Hermann, 1804

– tergite 2 shorter, covering about one-third of the abdomen, broadly rounded. Tergite 6 smaller, rectangular, with a bare surface, with 6 long and 4 short setae posterior (Figure 5(39))... *Phthiridium integrum* (Theodor and Moscona, 1954)

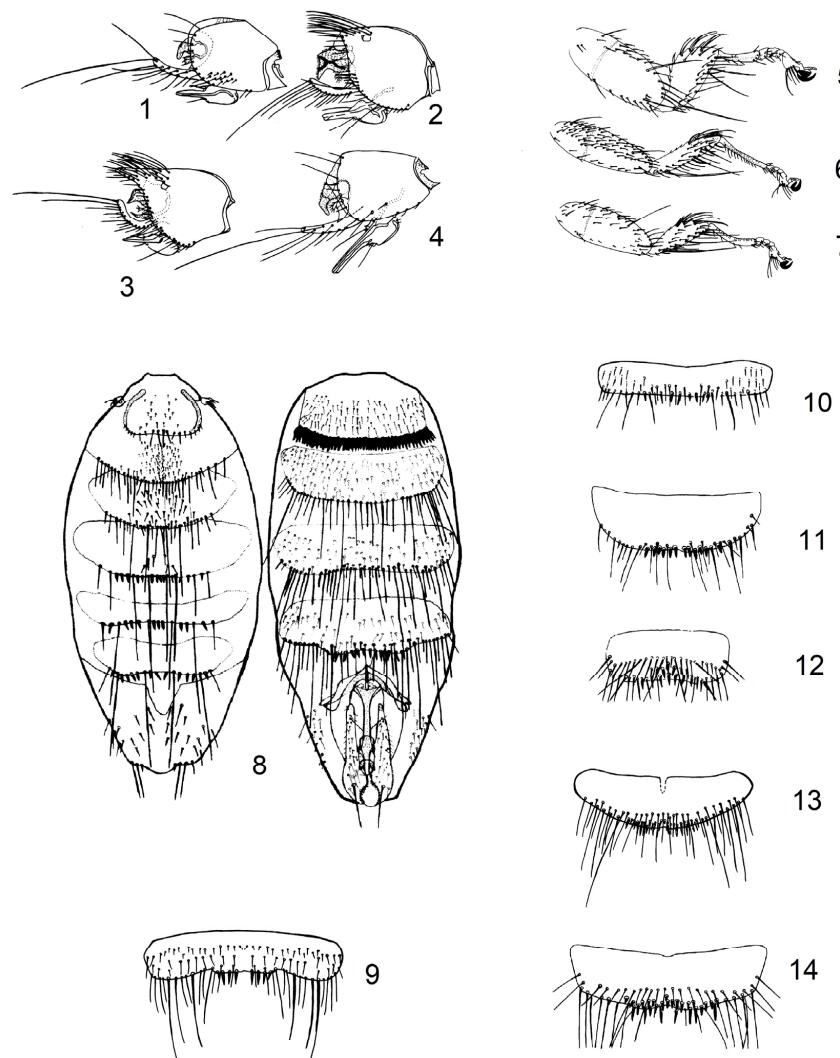


Figure 1. (1) *Nycteribia (Nycteribia) latreillii* (Leach, 1817), head; (2) *Penicillidia (Penicillidia) dufourii* (Westwood, 1835), head; (3) *Penicillidia (Penicillidia) monoceros* Speiser, 1900, head; (4) *Nycteribia (Acrocholidia) vexata* Westwood, 1835, head; (5) *Nycteribia (Nycteribia) kolenatii* Theodor and Moscona, 1954, leg; (6) *Phthiridium biarticulatum* Hermann, 1804, leg; (7) *Basilia nana* Theodor and Moscona, 1954, leg; (8) *Nycteribia (Nycteribia) pedicularia* Latreille, 180, male, ventral abdomen with claspers; (9) *Basilia nattereri* Kolenati, 1857, male, sternite 5; (10) *Basilia nana* Theodor and Moscona, 1954, male, sternite 5; (11) *Basilia mediterranea* Hurka, 1970, male, sternite 5; (12) *Basilia daganiae* Theodor and Moscona, 1954, male, sternite 5; (13) *Basilia italicica* Theodor, 1954, male, sternite 5; (14) *Basilia mongolensis nudior* Hurka, 1972, male, sternite 5.

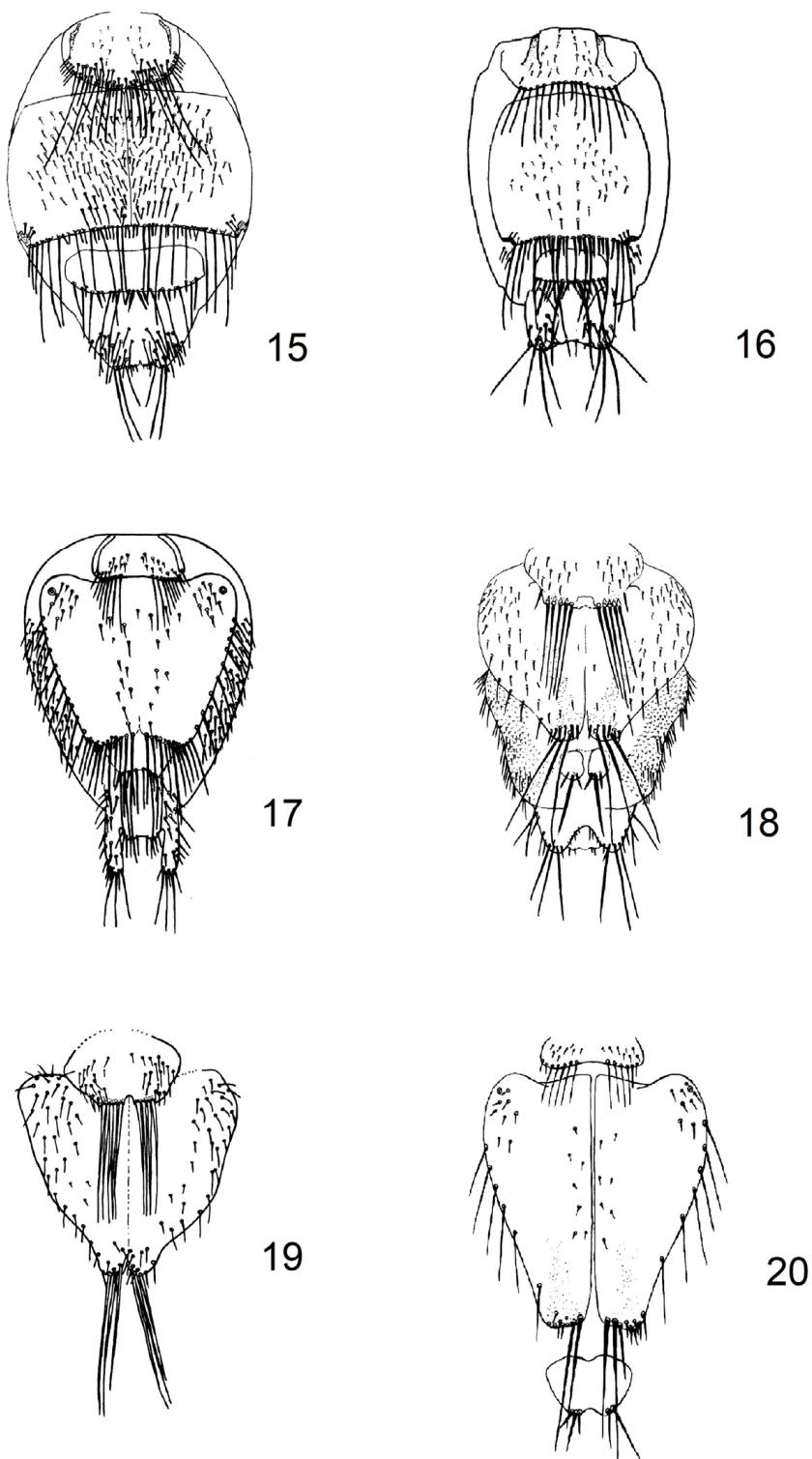


Figure 2. (15) *Basilia nattereri* Kolenati, 1857, female, abdomen; (16) *Basilia nana* Theodor and Moscona, 1954, female, abdomen; (17) *Basilia italicica* Theodor, 1954, female, abdomen; (18) *Basilia daganiae* Theodor and Moscona, 1954, female, abdomen; (19) *Basilia mediterranea* Hurka, 1970, female, abdomen; (20) *Basilia mongolensis nudior* Hurka, 1972, female, abdomen.

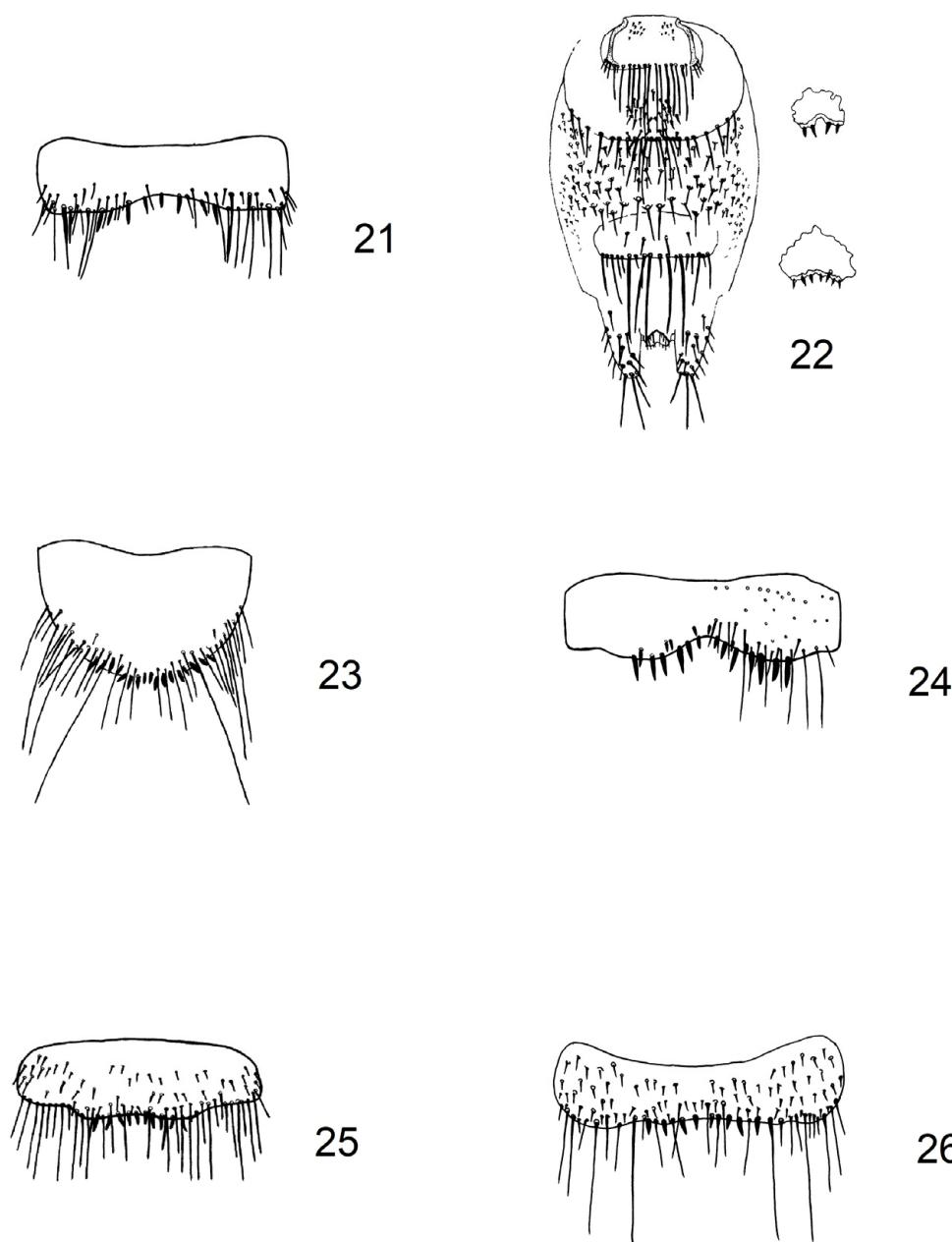


Figure 3. (21) *Nycteribia (Acrocholidia) vexata* Westwood, 1835, male, sternite 5; (22) *Nycteribia (Acrocholidia) vexata* Westwood, 1835, female, abdomen, genital plate; (23) *Nycteribia (Nycteribia) schmidlii* Schiner, 1853, male, sternite 5; (24) *Nycteribia (Nycteribia) latreillii* (Leach, 1817), male, sternite 5; (25) *Nycteribia (Nycteribia) pedicularia* Latreille, 1805, male, sternite 5; (26) *Nycteribia (Nycteribia) kolenatii* Theodor and Moscona, 1954, male, sternite 5.

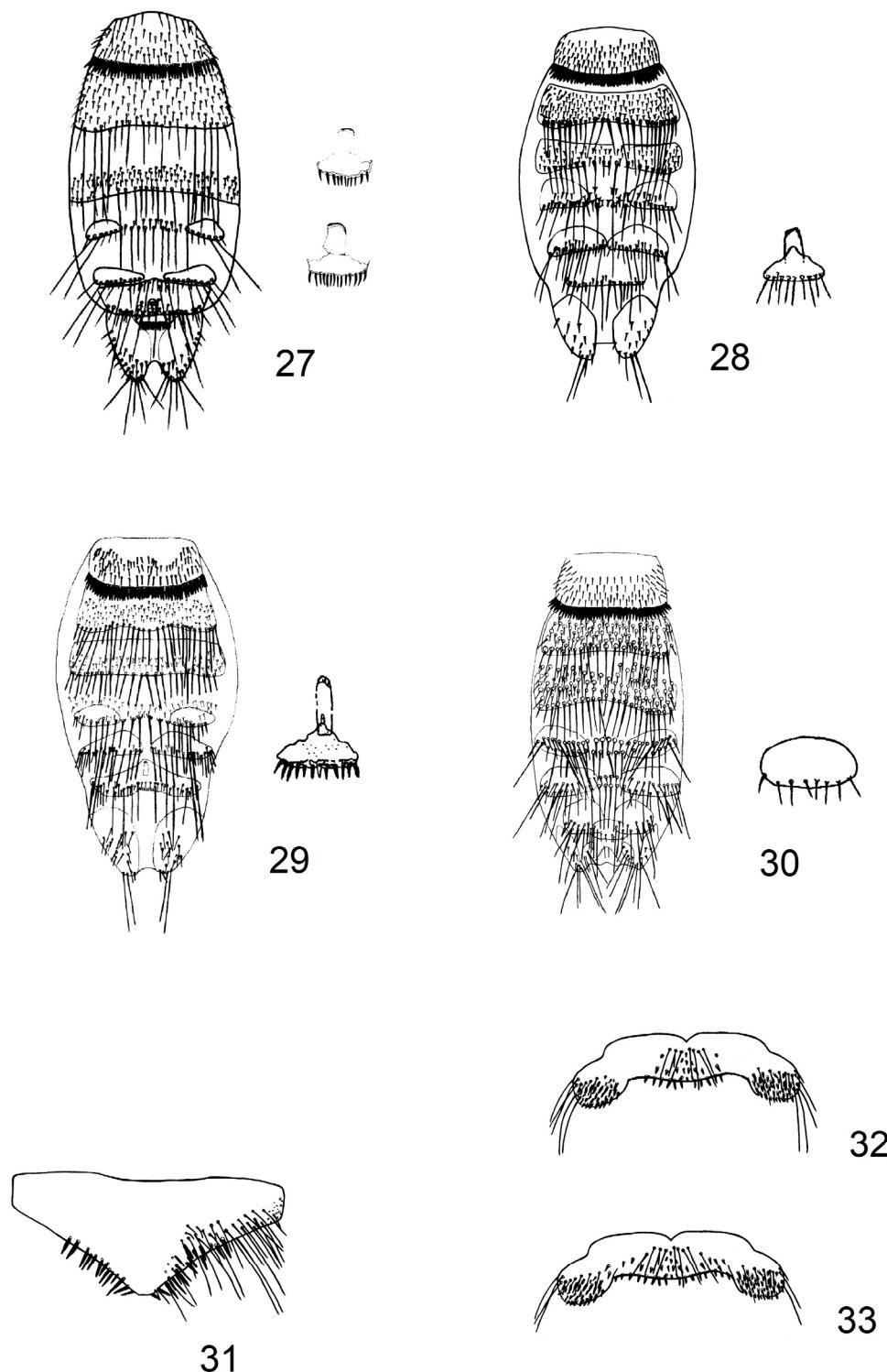


Figure 4. (27) *Nycteribia (Nycteribia) latreillii* (Leach, 1817), female, abdomen, genital plate; (28) *Nycteribia (Nycteribia) kolenatii* Theodor and Moscona, 1954, female, abdomen, genital plate; (29) *Nycteribia (Nycteribia) pedicularia* Latreille, 1805, female, abdomen, genital plate; (30) *Nycteribia (Nycteribia) schmidlii* Schiner, 1853, female, abdomen, genital plate; (31) *Penicillidia (Penicillidia) conspicua* Speiser, 1901, male, sternite 5; (32) *Penicillidia (Penicillidia) dufourii dufourii* (Westwood, 1835), male, sternite 5; (33) *Penicillidia (Penicillidia) monoceros* Speiser, 1900, male, sternite 5.

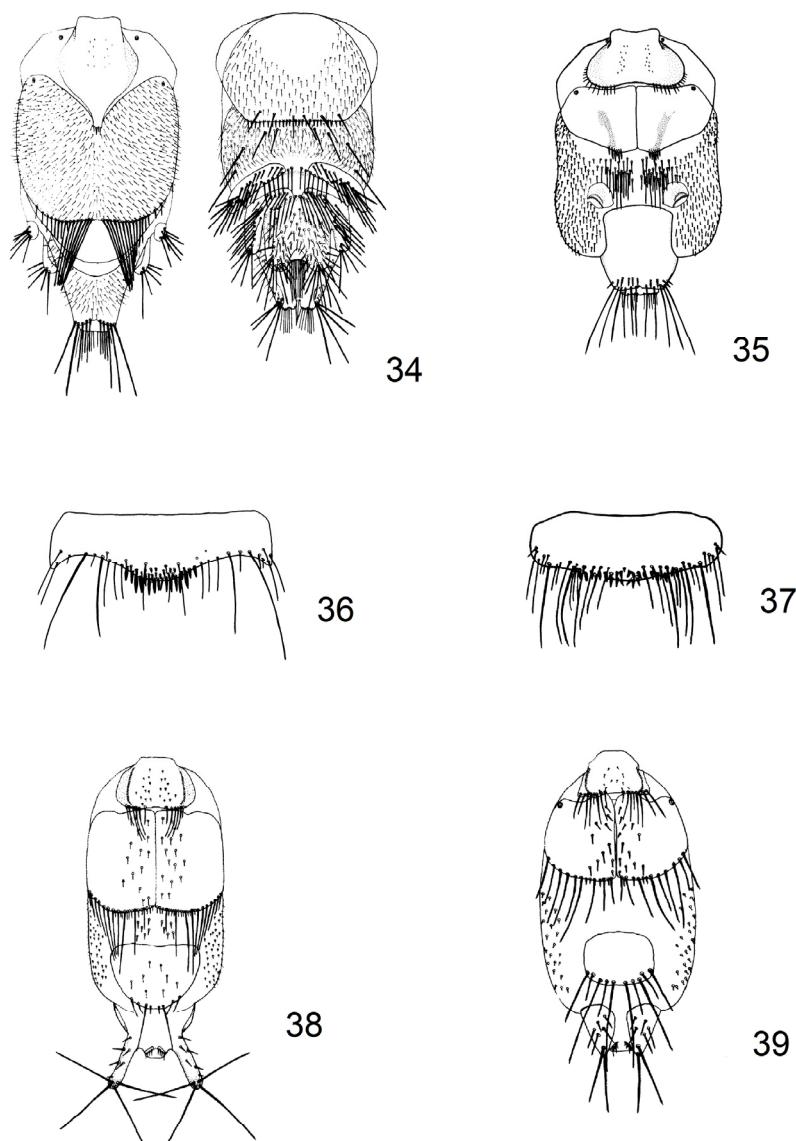


Figure 5. (34) *Penicillidia (Penicillidia) conspicua* Speiser, 1901, female, abdomen; (35) *Penicillidia (Penicillidia) dufourii dufourii* (Westwood, 1835), female, abdomen; (36) *Phthiridium biarticulatum* Hermann, 1804, male, sternite 5; (37) *Phthiridium integrum* (Theodor and Moscona, 1954), male, sternite 5; (38) *Phthiridium biarticulatum* Hermann, 1804, female, abdomen; (39) *Phthiridium integrum* (Theodor and Moscona, 1954), female, abdomen.

Bat Fly-Bat Species Associations and the Bipartite Network

Altogether, 173 host-parasite interactions were observed between 16 European bat fly species and 31 host species (bats) (See also Supplementary). The highest number of associations (17) with different host species and the lowest value of the index of specialization (d') was observed in *Penicillidia (Penicillidia) dufourii dufourii*, followed by *Nycteribia (Nycteribia) kolenatii* (17 host spp.), *N. (N.) schmidlii* (16; host spp.), and *Phthiridium biarticulatum* (16; host spp.). Most of the other species have low or relatively low specialization (< 0.5), and most species are oligoxenous, parasitizing on several families of one order [156]. Three species (*B. maditerranea*, *Basilia mongolensis nudior*, *Phthiridium integrum*) are stenoxenous, parasitizing more than one genus of hosts, but are restricted to one family [5,156], and four species (*B. italica*, *B. mediterranea*, *B. mongolensis nudior*, *Phthiridium integrum*) have high specialization. Only one nycteribiid species (*Basilia daganiae*) was monoxenous with only one host (*Pipistrellus kuhlii*) and $d' = 1$ (Figure 6 and Table 1).

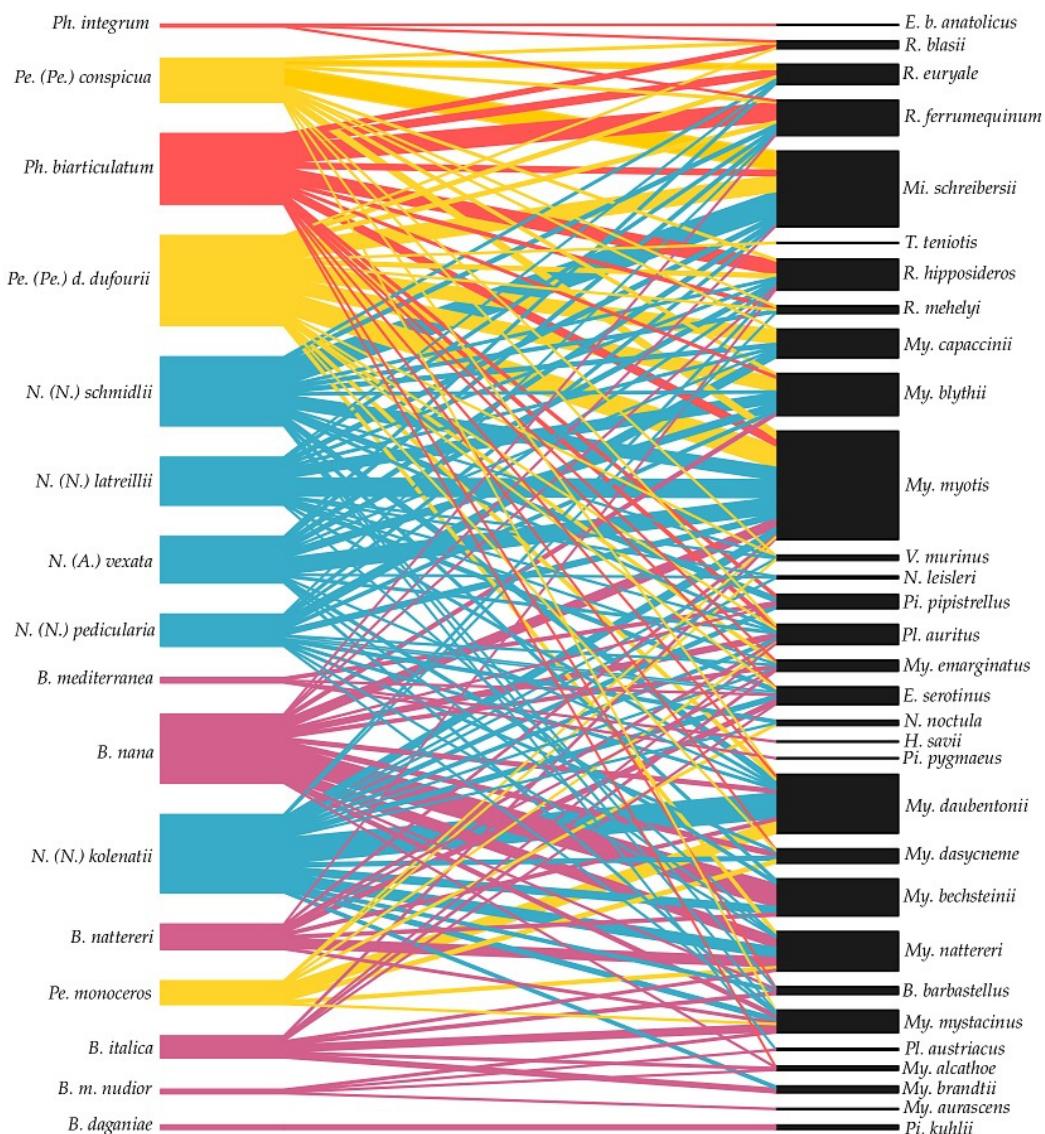


Figure 6. Bipartite network graph of interactions between bat flies (left) and bats (right). Each color of bat fly species means one genus (*Basilia*—purple, *Nycteribia*—blue, *Penicillidia*—yellow, and *Phthiridium*—red). The overlap of lines is minimized and parasite species are ordered so that the closest are those that have the most common hosts, and on the other hand, the parasites are ordered by similarity in terms of common hosts.

Three bat species, namely *Myotis myotis* (12 associations with different bat fly species) and *Plecotus auritus* with *Myotis daubentonii* with 12 associated species, were found. On the other hand, a relatively high number of bat species were associated with only one bat fly species (*Eptesicus bottae anatolicus*, *Hypsugo savii*, *Myotis davidi*, *P. kuhlii*, *P. pygmaeus*, and *Tadarida teniotis*) from Europe, according to the publications (Figure 6, Table 1, and Supplementary).

According to the results of bipartite analysis, the network of bats and bat flies has low connectance ($C = 0.31$), low specialization ($H2' = 0.30$), and a low degree of nestedness ($N = 0.31$ with nestedness temperature = 29.19). The specialization on the species level (d') ranged between 0.14 and 1.00 (Table 1). The strength of the interactions between the parasites (bat fly) and hosts (bats) is illustrated in the bipartite graph (Figure 6), as the thickness of the line connecting parasites with host species reflects the number of records published. The thickness of the box of individual bat and bat fly species represents the

relative number of records of particular parasites and host species. As the network is very complex, different colors represent different bat fly genera.

Table 1. The number of hosts published and the index of bat flies' specialization at the species level (d'), and the specificity of European bat fly species. Grey rows mean high specialization and dark grey rows mean the highly specialized monoxenous species.

Bat Fly Species	No. of Hosts spp. Published	d'	Specificity
<i>Basilia daganaiae</i>	1	1.00	monoxenous
<i>Basilia italica</i>	9	0.50	oligoxenous
<i>Basilia mediterranea</i>	4	0.67	stenoxyenous
<i>Basilia mongolensis nudior</i>	4	0.65	stenoxyenous
<i>Basilia nana</i>	15	0.32	oligoxenous
<i>Basilia nattereri</i>	9	0.30	oligoxenous
<i>Nycteribia (Acrocholida) vexata</i>	13	0.14	oligoxenous
<i>Nycteribia (Nycteribia) kolenati</i>	17	0.30	oligoxenous
<i>Nycteribia (Nycteribia) latreillii</i>	14	0.18	oligoxenous
<i>Nycteribia (Nycteribia) pedicularia</i>	14	0.20	oligoxenous
<i>Nycteribia (Nycteribia) schmidlii</i>	16	0.18	oligoxenous
<i>Penicillidia (Penicillidia) conspicua</i>	13	0.24	oligoxenous
<i>Penicillidia (Penicillidia) dufourii dufourii</i>	17	0.19	oligoxenous
<i>Penicillidia (Penicillidia) monoceros</i>	8	0.41	oligoxenous
<i>Phthiridium biarticulatum</i>	16	0.35	oligoxenous
<i>Phthiridium integrum</i>	3	0.56	stenoxyenous

4. Discussion

There is a growing interest in bat flies due to their importance in the study of parasitic and hyperparasitic relations [38,176,177], as well as their vectorial role in pathogen transfer [32,33,178,179]. Identification and knowledge of the ecology and host–parasite network of Nycteribiidae are critical. In order to contribute to the knowledge of this interesting and important group of parasites and to fill knowledge gaps, we developed a key for the identification of European nycteribiid species. Based on published data, we analyzed their hosts, their degree of specialization, and the communities (modules) that they form within parasite–host relationships.

The outdated keys of European Nycteribiidae (Diptera) were one of the complications making it difficult to study their ecology and possible importance for pathogen transmission [180]. Therefore, we have herein provided a summary key of 16 nycteribiid species currently known from Europe.

With this method, we aimed to find broader contexts, and more generally, valid patterns, and we succeeded in doing so. For example, disturbance may in turn cause an increase in accidental or transient associations of bats and their parasites with non-primary hosts. Such dynamics may increase the likelihood of host switching, and thus the movement of pathogens to new host species and the expansion of their range, increasing the risk of transmission to other mammals, including humans [5]. Moreover, according to published papers, the fact that specialization within parasite–host networks varied over time and space in the different communities analyzed (the influence of season and associated changes on the abundance and species' richness of bats, the influence of latitude, etc. according to [35,47,50], and differences in host–parasite associations between published studies [21] and the recent literature [7] (e.g., [51,52] and many references there)), this indirectly demonstrates the greater potential for bat flies to parasitize more bat species than can be detected in a sub-study devoted to individual communities. This is consistent with the findings of our more comprehensive analysis.

Thus, our study demonstrated that the bat flies included generally showed variable degrees of host specialization, and Nycteribiidae seem to be less specialized than was supposed by Marshall [15] and Verrett et al. [34]. The information published in these two papers, that Nycteribiids are usually highly specific, with each species of bat fly parasitizing

a single bat species or genus or rarely, families, contradicts the other published records from Europe used in this study. Only one bat fly species (*B. daganae*) was recorded from only one host species (*P. kuhlii*), possibly because there are only five reports and research in the area where this species occurs. It could potentially parasitize other host species; however, we do not know as we do not have data in the area. In contrast, most of the European species are oligoxenous (parasitizing more genera from one family) according to the published records [21]. Which intrinsic and extrinsic (biotic and abiotic) factors are crucial in terms of the ecological specialization of nycteribiids to particular bat host species are debatable. Sharing resources, including roosts, by bats can involve costs as well as benefits [181].

Dolabela Falcão et al. [50] claimed that more abundant bat species had a greater richness of associated ectoparasites. On the other hand, in addition to the population density of the bat host species, other factors may influence the diversity of associated ectoparasites. The age structure of the observed bat host population, the fitness of the animals in the subpopulation and population, and the season relative to the phenology of both the host and parasite may also be important [182–184].

The results of the bipartite network analysis shed more light onto the complexity of the relationships between bat flies (parasites) and bats (their hosts). Similarly to other bat fly–bat network analyses [35,47,50], our results of bipartite network analysis, based on the published records from Europe, show low specialization, low modularity, and low nestedness. This seems to be a general pattern in bat fly–bat networks, although the nestedness, connectance, and the number of compartments (modules, communities) are related to bat richness [50]. When comparing our results with Durán et al. [39], the connectance and network specialization are very similar, but the modularity was much lower in our study. This difference may also be caused by different species' richness in these studies. Despite the low specialization (H_2' and d'), according to the results of our study, some of our results indicate the importance of the phylogenetic and ecological relatedness of hosts when phylogenetically related bat species support similar associated ectoparasite communities. Such findings indicating the importance of phylogenetic relationships are consistent not only in bat flies and bats [50], but also among different types of interactions and taxonomic groups [185–188], even though comparing results from different regions and between different taxa is problematic. In our study, Nycteribiidae showed an affinity towards particular bat species that are phylogenetically related, e.g., *B. m. nudior* and *B. italica* towards three cryptic whiskered bats (*Myotis alcathoe*, *M. mystacinus*, *M. brandtii*). It has been observed that bat species that share roosts are prone to becoming infested by ectoparasites of other species [189], and roosting site preference and roosting behaviors of bats may be factors that significantly affect the dynamic of bat–bat flies interactions [190]. This can also be assumed based on the data in the present study. For example, *N. (A.) vexata*, *N. (N.) latreilliid*, and *P. d. dufourii* presented predominant associations with *M. myotis*, *M. blythii*, and *Rhinolophus* species, which prefer underground or attic spaces for roosting and often together exploit the same roosting site [191]. Additionally, taxons *N. (N.) schmidlii*, *P. conspicua*, and *Ph. biarticulatum* are associated with three species of the genus *Rhinolophus* and with *M. schreibersii*, showing a similar shelter preference [192,193].

These findings suggest a high probability of specific Nycteribiidae in bat taxa with similar roosting ecology. As the complexity of the problematics of pathogen transmission by nycteribiids have not been exhaustively elucidated to date, it is extremely important to further investigate the relationships between specific species of nycteribiids and bats [11,194,195].

5. Conclusions

Current knowledge, complemented by our results, points to the importance of further research focusing on ecological networks and parasite–host relationships, particularly, but not exclusively, in groups at high risk of epidemiologically serious diseases transmissible to other mammals, including humans. This may facilitate targeted research on bat fly–bat networks, bat fly specialization, and nycteribiid-transmitted bat pathogens. At the same time, however, the legislative conditions that regulate sampling of their hosts (Chiroptera)

must be respected in order to ensure minimal impact of the research on populations of protected taxa. In addition, collection methods are challenging [180].

Supplementary Materials: The Overview of the Recorded Interactions. The following supporting information can be downloaded at: <https://www.mdpi.com/article/10.3390/d15040573/s1>.

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