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Phylogeny and biogeography of the enigmatic ghost lineage *Cylindrotomidae* (Diptera, Nematocera)

Iwona Kania-Kłosok¹✉, André Nel², Jacek Szwedo³, Wiktoria Jordan-Stasiło¹ & Wiesław Krzemiński⁴

Ghost lineages have always challenged the understanding of organism evolution. They participate in misinterpretations in phylogenetic, clade dating, biogeographic, and paleoecologic studies. They directly result from fossilization biases and organism biology. The *Cylindrotomidae* are a perfect example of an unexplained ghost lineage during the Mesozoic, as its sister family Tipulidae is already well diversified during the Cretaceous, while the oldest *Cylindrotomidae* are Paleogene representatives of the extant genus *Cylindrotoma* and of the enigmatic fossil genus *Cyttaromyia*. Here we clarify the phylogenetic position of *Cyttaromyia* in the stem group of the whole family, suggesting that the crown group of the *Cylindrotomidae* began to diversify during the Cenozoic, unlike their sister group Tipulidae. We make a comparative analysis of all species in *Cyttaromyia*, together with the descriptions of the two new species, *C. gelhausi* sp. nov. and *C. freiwaldi* sp. nov., and the revision of *C. obdurescens*. The cylindrotomid biogeography seems to be incongruent with the phylogenetic analysis, the apparently most derived subfamily Stibadocerinae having apparently a 'Gondwanan' distribution, with some genera only known from Australia or Chile, while the most inclusive *Cylindrotominae* are Holarctic.

Cylindrotomidae Schinner, 1863¹ together with Limoniidae Speiser, 1909², Pediciidae Osten-Sacken, 1860³, and Tipulidae Latreille, 1802⁴ sensu stricto are classified within Tipuloidea Latreille, 1802⁴, group of insects present in the fossil record since at least 220 Ma (Triassic)⁵. With only 71 extant⁶ and 16 extinct species⁷, this smallest family within Tipuloidea is divided into two subfamilies *Cylindrotominae*, represented mainly in Holarctic Region, and *Stibadocerinae* with an example of vicariant distribution with a sister-group relationship between South American and East Asian taxa, supporting hypothesis an 'ancestral' trans-Pacific biota⁸.

According to phylogenetic synthesis based on combined morphological characters of adult, larvae and pupae, together with nuclear gene sequence data as 28S rDNA or CAD, the *Cylindrotomidae* are found as a sister group of Tipulidae (both being treated as subfamilies in Tipulidae in Ref.⁹. This group of insects is generally indicated as a sister group or being closely related to the Tipulidae by other authors^{10–20}. But, within *Cylindrotomidae*²¹, only the representatives of subfamily *Cylindrotominae* are known from fossil record. The oldest described representatives of *Cylindrotominae* are only known from the Paleogene (56.0–47.8 Ma) by the extant genera *Cylindrotoma* Macquart, 1834²² and *Diogma* Edwards, 1938^{23–33}, plus the extinct genus *Cyttaromyia* Scudder, 1877^{7,34}, while the oldest stem Tipulidae are Jurassic and the oldest crown Tipulidae are Cretaceous^{5,35–37}. Thus, the *Cylindrotomidae* can be considered as a typical ghost lineage during the Mesozoic.

This phytophagous group of craneflies, which immatures lives among mosses and herbaceous plants.

The *Cylindrotomidae* (Supplementary Data S1) probably knew a period of diversification during the Eocene, sufficient to become frequent enough to be found as fossils. Most of the fossil *Cylindrotomidae* are

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know from the Middle Eocene Baltic amber^{31,32,38,39} and the Late Eocene Florissant Formation in USA^{24,39}. Three species of *Cylindrotoma* are known from impressions of the Ypresian Fur Formation; two species from the Ølst Formation of Denmark were described within *Cyttaromyia*²⁹ (Supplementary Table S1). Four species of *Cyttaromyia* were described from the Eocene Green River Formation USA, the other were described from Florissant Formation and Kishenehn Formation in USA, Middle Salt Formation in Alsace (France), Biamo Formation in Russia, and from Baltic amber^{24,27,28,30,31,40}. Here we propose a morphological phylogenetic analysis to define the relationships between extinct genus *Cyttaromyia* and the other taxa in the family. We also describe two new species of *Cyttaromyia* on the basis of new fossils from the same Formation, and new technics of research give us possibility to redescribe *Cyttaromyia obdurescens* Cockerell, 1924²⁷.

Results

Systematic paleontology.

Order Diptera Linnaeus, 1758⁴²

Infraorder Tipulomorpha Latreille, 1802⁴

Family Cylindrotomidae Schinner, 1863¹

Subfamily Cylindrotominae Schinner, 1863¹

Genus *Cyttaromyia*

Type species: *C. yttaromyia fenestrata* Scudder, 1877³⁴, by monotypy.

Key to species of the genus *Cyttaromyia* Scudder, 1877³⁴

1. Wings without distinct patterning.....3.
 - Distinct patterns of coloration on wings.....2.
2. Rs longer than R_{2+3+4} and R_{3+4} combined.....
 -*Cyttaromyia vahldieki* Freiwald, 1991²⁹
Denmark/Ølst Formation
 - Rs shorter than R_{2+3+4} and R_{3+4} combined.....
 -*Cyttaromyia rayona* Freiwald & Krzemiński, 1991³⁰
Russia/Biamo Formation
3. Wings hyaline.....4.
 - Wings pale brownish without conspicuous markings with end of marginal cell apically somewhat clouded (Cockerell, 1924).....
 -*Cyttaromyia reclusa* Cockerell, 1924²⁶
USA/Green River Formation/Roan Moutains/Colorado
4. Vein R_1 well-developed.....5.
 - Vein R_1 reduced.....7.
5. Crossvein m-cu situated beyond bifurcation of Mb on M_{1+2} and M_{3+4} ; Sc terminating in C far beyond fork of Rs.....6.
 - Crossvein m-cu situated at bifurcation of Mb on M_{1+2} and M_{3+4} ; Sc terminating in C just beyond fork of Rs.....
 -*Cyttaromyia gelhausi* sp. nov.
USA/Green River Formation
6. Crossvein sc-r one of its length before tip of Sc; A_1 tip before tip of Sc level.....
 -*Cyttaromyia obdurescens* Cockerell, 1925²⁷
USA/Green River Formation/Roan Moutains/Colorado
 - Crossvein sc-r at least two of its length before tip of Sc; A_1 tip beyond e tip of Sc level.....
 -*Cyttaromyia princetoniana* Scudder, 1894²⁴
USA/Green River Formation
7. Sc elongate, terminating in C well beyond level of fork of Rs.....8.
 - Sc short, terminating in C before level of fork of Rs.....
 -*Cyttaromyia freiwaldi* sp. nov.
USA/Green River Formation

8. Crossvein m-cu situated at most or before fork of Mb on M_{1+2} and M_{3+4}9.
 – Crossvein m-cu situated beyond fork of Mb on M_{1+2} and M_{3+4}
*Cyttaromyia frelloi* Krzemiński, 1998³¹
 Baltic amber
9. Vein r-r (R_2) terminating before level of r'-m', before level of m-m.....10.
 – Vein r-r (R_2) terminating at level of r'-m', behind level of m-m.....
*Cyttaromyia fenestrata* Scudder, 1877³⁴
 USA/Green River Formation
10. d'-cell as long as d-cell or shorter.....11.
 – d'-cell longer than d-cell.....12.
11. M_{3+4} bifurcation on M_3 and M_4 approximately at level of m-m; d-cell 2× as long as M_4 ; m-m beyond level of tip of r-r (R_2).....
*Cyttaromyia lynnae* De Jong, 2019⁴⁰
 USA/Kishenehn Formation.
- M_{3+4} bifurcation on M_3 and M_4 before level of m-m; d-cell 1.5× as long as M_4 ; m-m approximately at level of tip of r-r (R_2).....
*Cyttaromyia fuscula* Cockerell, 1921²⁵
 USA/Green River Formation
12. Fork of Mb at level of fork of Rs; d'-cell narrowed at base.....
*Cyttaromyia quievreuxi* Séguy, 1934²⁸
 Alsace, France/Middle Salt Formation.
- Fork of Mb before level of fork of Rs; d'-cell narrow, but not narrowed at base.....13.
13. Tip of r-r (R_2) beyond level of tip of fork of M_{3+4} on M_3 and M_4 ; Rs at 2× as long as R_{2+3+4}
*Cyttaromyia scudderi* Freiwald, 1991²⁹
 Denmark/Ølst Formation
- Tip of r-r (R_2) before level of tip of fork of M_{3+4} on M_3 and M_4 ; Rs at approximately as long as R_{2+3+4}
*Cyttaromyia rossi* Krzemiński, 2019⁴¹
 UK/Isle of Wight/Bembridge Marls

Cyttaromyia obdurescens Cockerell, 1925²⁷
 (Fig. 1).

Material examined. Holotype No. 26284 (AMNH) (female); American Museum National History; Green River Formation USA, Eocene.

Emended diagnosis. Wing without color spots; Sc elongate, terminating in C well beyond level of fork of Rs, beyond r-m level but far before m-m and r'-m' level; opposite approximately half the length of R_{2+3+4} ; vein r-r (R_2) terminating far before r'-m' and m-m level, at level of basal part of M_3 ; R_1 well-developed; R_{2+3+4} longer than half length of Rs; d'-cell longer than d-cell, narrowed at its base; crossvein m-cu positioned beyond fork of Mb on M_{1+2} and M_{3+4} ; apical section of M_3 almost as long as d-cell; A_1 tip positioned near apex of wing, far behind level of Mb bifurcation on M_{1+2} and M_{3+4} , before r-m level.

Comparison. *Cyttaromyia obdurescens* differs from *C. fenestrata*, *C. freiwaldi* sp. nov., *C. frelloi*, *C. fuscula*, *C. lynnae*, *C. quievreuxi*, *C. scudderi*, and *C. vahldieki* by a well-developed vein r-r (R_2). Wing of *C. obdurescens* is without spot. *C. rayona* and *C. vahldieki* have different patterning of wings. In contrast to *C. gelhausi* sp. nov., crossvein m-cu is situated beyond Mb, while in *C. gelhausi* sp. nov., it is located at Mb bifurcation, in *C. frelloi* just before Mb bifurcation. In *C. obdurescens*, vein sc-r is located one of its length from tip of Sc, tip of A_1 is located before tip of Sc, while in *C. princetoniana* vein sc-r is located at least two of its length before the tip of Sc, A_1 tip is located beyond level of tip of Sc. In *C. obdurescens* crossvein m-cu is positioned beyond fork of Mb measured from base of wing, while in *C. rossi*, m-cu is distinctly before fork of Mb.

***Cyttaromyia gelhausi* sp. nov.** <http://zoobank.org/urn:lsid:zoobank.org:act:9165E3D2-514B-4524-9F85-A9895CBF2A31>.

(Figs. 2, 3).

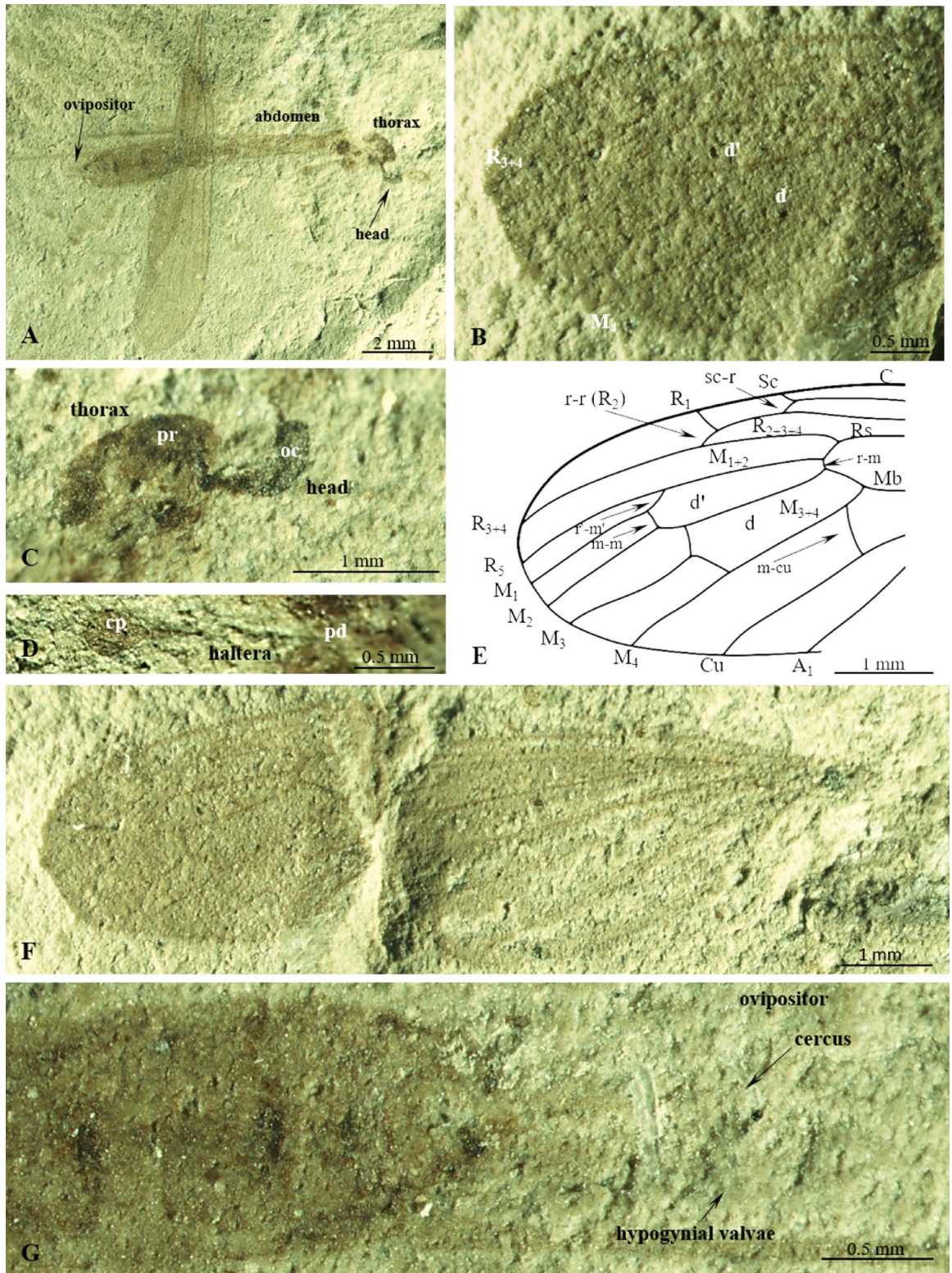


Figure 1. *Cyttaromyia obdurescens* Cockerell, 1925²⁷, holotype No. 26284 (AMNH) (female): (A) habitus, lateral view; (B) apex of wing; (C) head and thorax, enlarged lateral view; (D) haltera; (E) apex of wing, drawing; (F) wing; C. terminal part of abdomen with *ovipositor* visible. cp, *capitellum*, pd, *pedicellus* of haltera, oc, *ocellus*, ped, *pedicellus*, pr, *pronotum*.

Material examined. Holotype MNHN.FA71341 (18a, male); additional material 18b, female; 18c, female (on the same slab with holotype); Muséum national d’Histoire naturelle (MNHN), Paris; Green River Formation USA, Eocene.

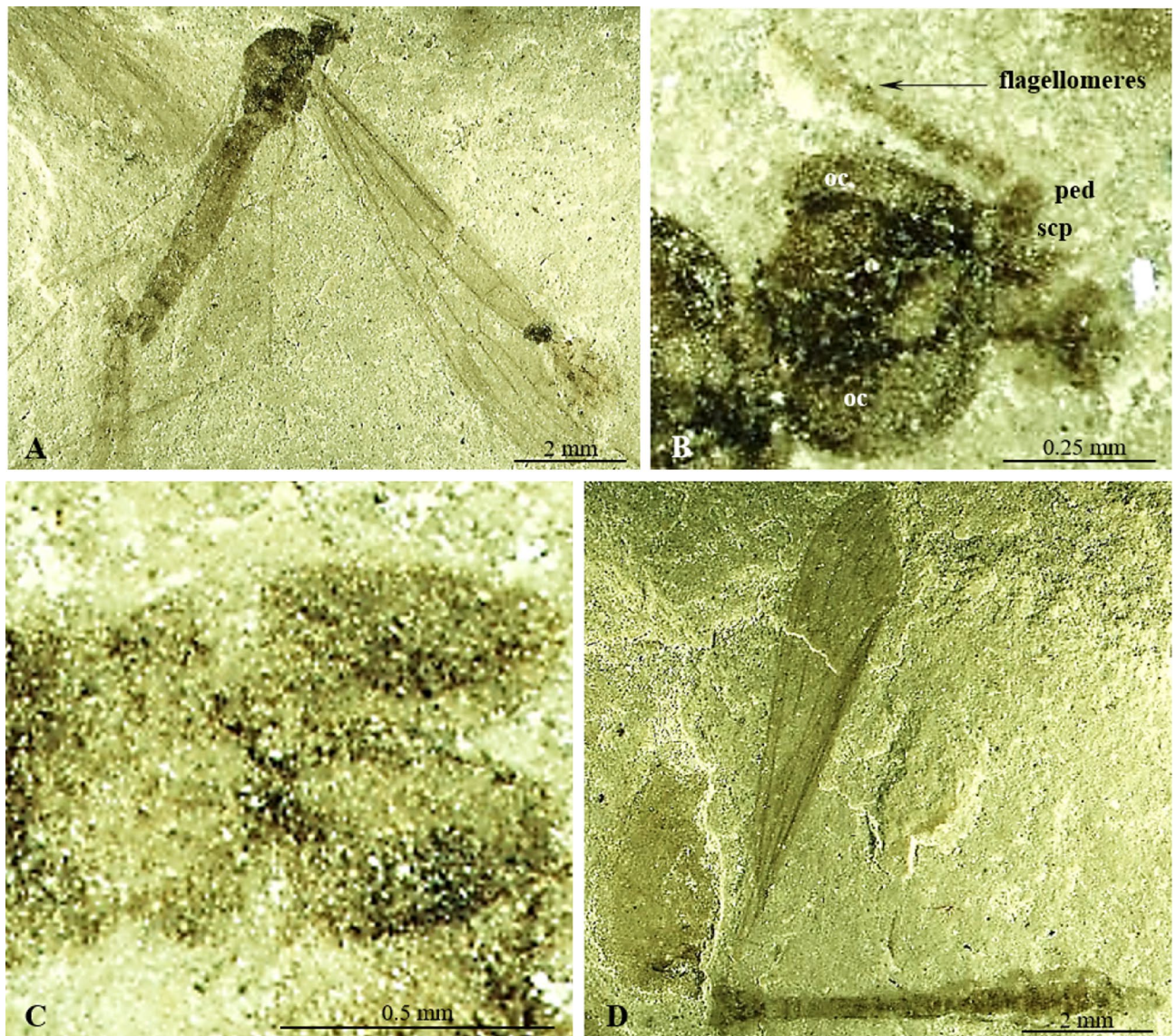


Figure 2. *Cyttaromyia gelhausi* sp. nov.: A.–C. No. MNHN.FA71341 (18a, male) (holotype): (A) habitus, latero-ventral view; (B) head, ventral view; (C) hypopygium, ventral view; D. No. MNHN.FA71341 (18c, female) (additional material), habitus, lateral view. oc, ocellus; ped, pedicellus; scp, scapus.

Etymology. The specific name is given to honor Doctor John Gelhaus (Academy of Natural Sciences of Drexel University), the eminent specialist on extinct and extant insects.

Diagnosis. Flagellomeres short and relatively wide; wing without color spots; Sc not very elongate, terminating in C just beyond level of fork of Rs, opposite level of crossvein r–m; opposite approximately 1/10 length of R_{2+3+4} ; vein r–r (R_2) terminating far before r'–m' and just before m–m level, at level of basal part of M_3 ; R_1 well-developed; R_{2+3+4} longer than half length of Rs; d'-cell longer than d-cell, narrowed at its base; crossvein m-cu positioned at fork of Mb on M_{1+2} and M_{3+4} ; apical section of M_3 almost as long as d-cell.

Comparison. *Cyttaromyia gelhausi* sp. nov. has no distinct color patterning of the wings in contrast to *C. vahldieki* and *C. rayona*. The wing of *C. reclusa* is pale brownish without conspicuous markings²⁶, but the end of the marginal cell and the veins bounding the discal cell apically are somewhat clouded. The body length of *C. gelhausi* sp. nov. is at most 6.83 mm, wing length 7 mm, while the body length of *C. reclusa* is 13.5 mm and wing length 12 mm. In *C. gelhausi* sp. nov. tibial spurs are absent while pattern of tibial spurs of *C. frelloii* is 1:1:2³¹. Vein R_1 is well-developed in *C. gelhausi* sp. nov., while in *C. freiwaldi* sp. nov., *C. frelloii*, *C. fuscula*, *C. lynnae*, *C. quievreuxi*, *C. rossi*, *C. scudderi*, *C. vahldieki* R_1 is reduced. In *C. gelhausi* sp. nov. Sc is not very elongate, terminating in C just beyond level of fork of Rs, opposite the level of crossvein r–m, while in *C. fenestrata*, *C. frelloii*, *C. fuscula*, *C. lynnae*, *C. obdurescens*, *C. princetoniana*, *C. quievreuxi*, *C. rayona*, *C. scudderi*, and *C. vahldieki*, it is terminating far beyond level of Rs. In *C. freiwaldi* sp. nov. Sc terminating in C before Rs level. In *C. fenestrata* and *C. vahldieki*, Sc is very elongated, terminating in C opposite basal part of M_3 . Crossvein m-cu is positioned

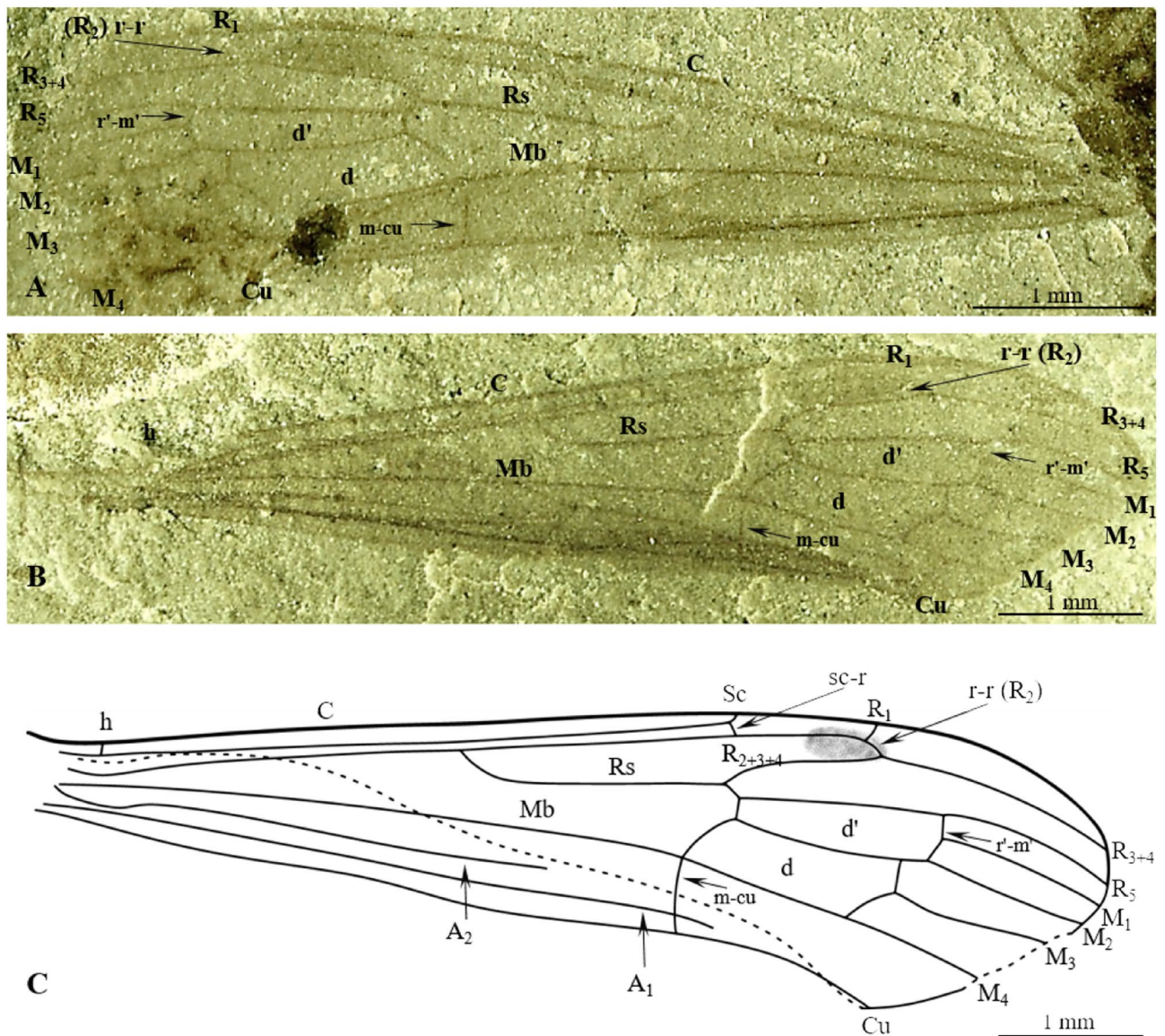


Figure 3. *Cyttaromyia gelhausi* sp. nov.: (A) No. MNHN.FA71341 (18a, male) (holotype), wing; (B), C. No. MNHN.FA71341 (18c, female) (additional material): (B) wing; (C) wing, drawing.

at fork of Mb on M₁₊₂ and M₃₊₄ while in other fossil species of the genus *Cyttaromyia* this vein is situated beyond fork of Mb; in *C. frelloi* this vein is located before fork of Mb. Moreover, in *C. vahldeki* and *C. fenestrata* vein r-r (R₂) terminating in R₃₊₄ beyond d' level, far beyond the level of basal part of M₃, in *C. gelhausi* sp. nov. this vein is terminating at the level of d', at the level of basal part of M₃.

***Cyttaromyia freiwaldi* sp. nov.** (Figs. 4, 5).

<http://zoobank.org/urn:lsid:zoobank.org:act:DB604450-AD47-4644-82E0-A33FEAEB7157>.

Material examined. Holotype MNHN.FA71342 (70A, part/68, counterpart, female), additional material 70B, female, on the same slab as holotype), Muséum national d'Histoire naturelle (MNHN), Paris; Green River Formation USA, Eocene.

Etymology. The new species is dedicated to the German eminent researcher Doctor Andre Freiwald (Institut für Paläontologie, Universität Erlangen).

Diagnosis. Wing without color spots; Sc short, terminating in C before level of fork of Rs, far before level of crossvein r-m; vein r-r (R₂) terminating far before r'-m' level and at m-m level, just beyond level of basal part of M₃; R₁ atrophied; R₂₊₃₊₄ longer than half length of Rs; d'-cell shorter than d-cell, narrowed at its base; crossvein m-cu positioned beyond fork of Mb on M₁₊₂ and M₃₊₄; M₃ shorter than d-cell.

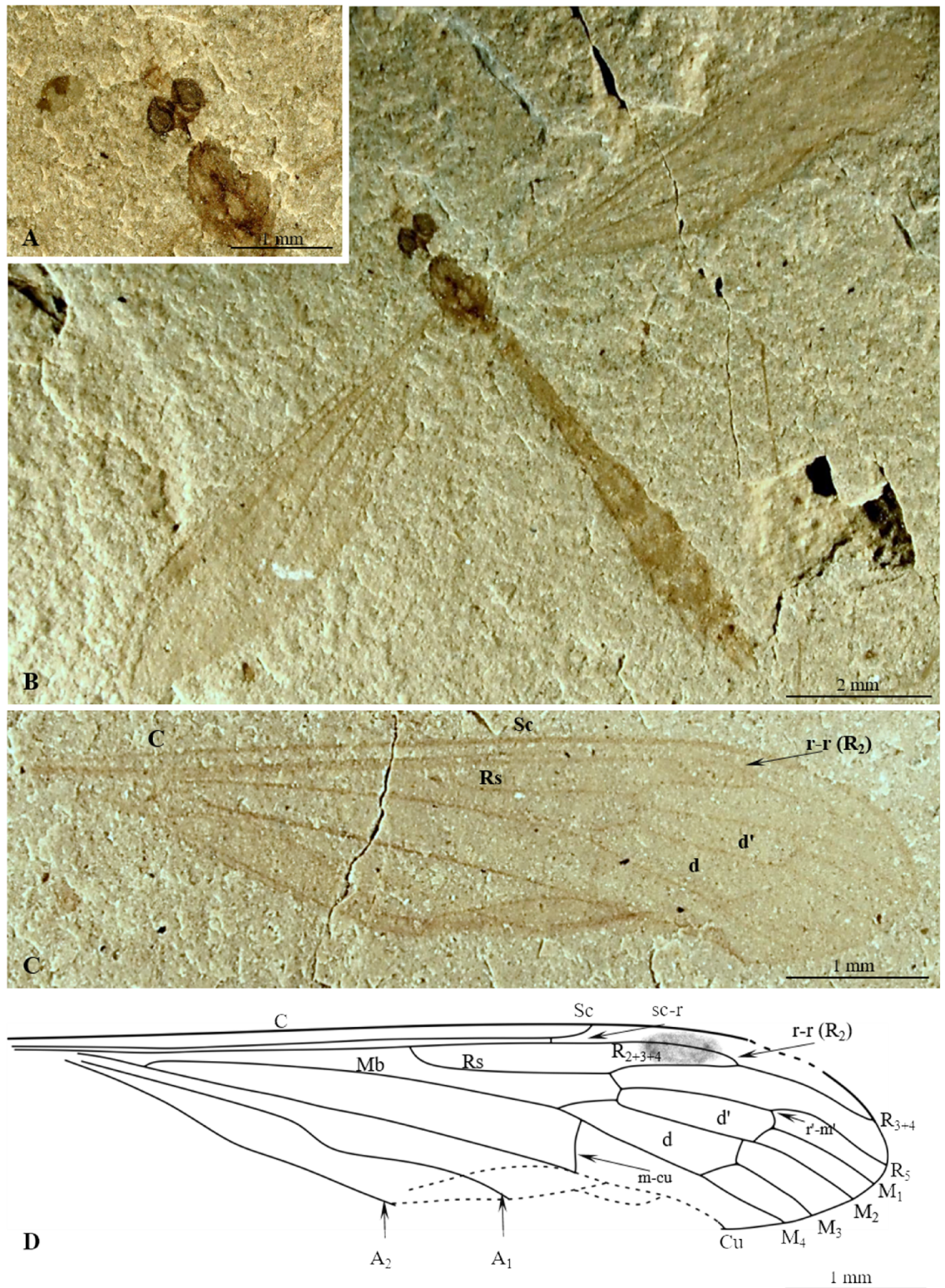


Figure 4. *Cyttaromyia freiwaldi* sp. nov. No. MNHN.FA71342 (70A, part, female), (holotype): (A) habitus, dorsal view; (B) head, dorsal view; (C) ovipositor, dorsal view, (D) wing venation, (E) wing venation, drawing.

Comparison. *Cyttaromyia freiwaldi* sp. nov. has no distinct patterning of the wings in contrast to *C. vahldieki* and *C. rayona*. In contrast to *C. reclusa*, the body length of *C. freiwaldi* sp. nov. is at most 4.8 mm, wing length 6.22 mm, while the body length of *C. reclusa* is 13.5 mm and wing is 12 mm long with the end of the marginal cell and the veins bounding the discal cell apically are somewhat clouded²⁶. *C. freiwaldi* sp. nov. differs from other

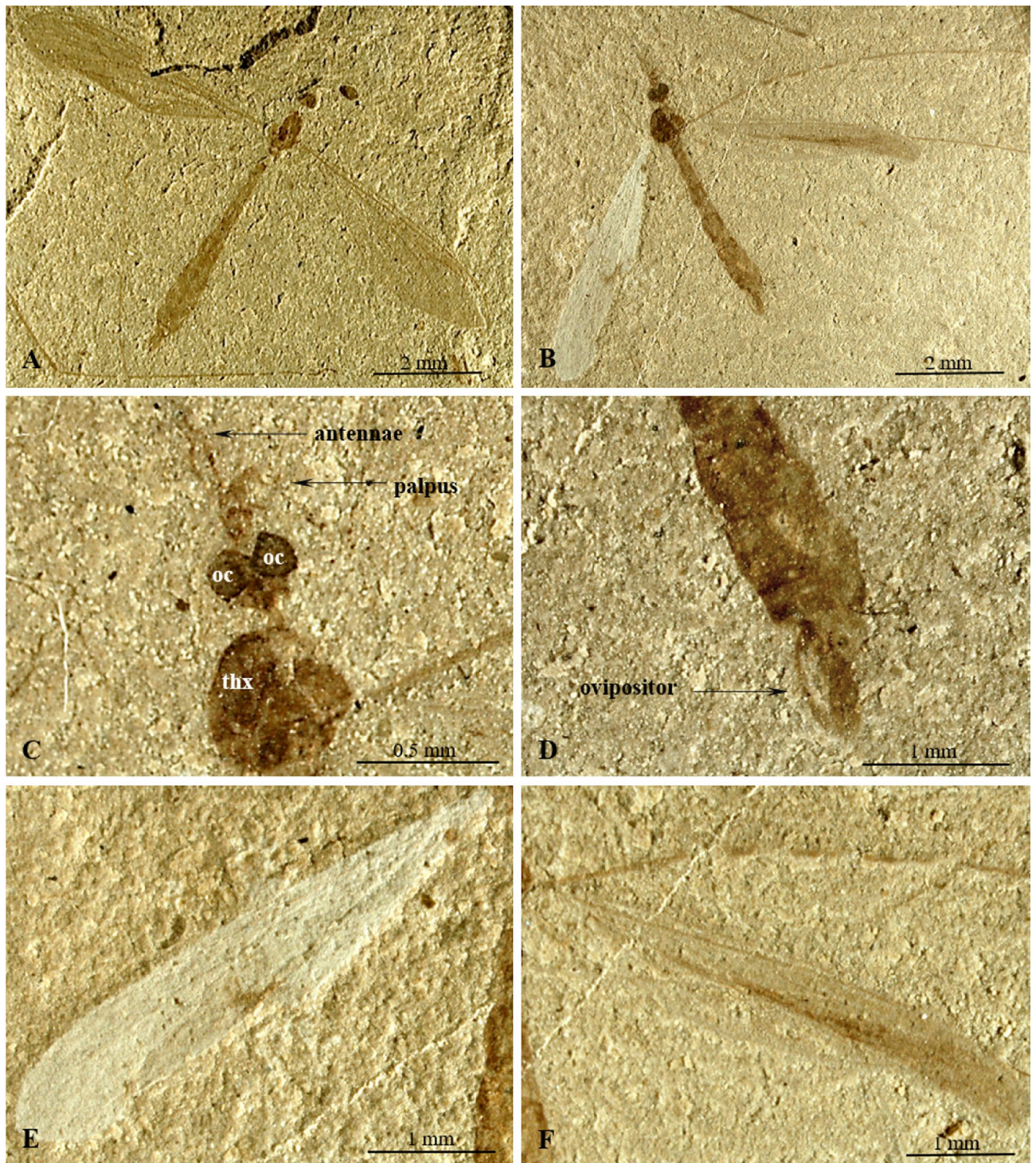


Figure 5. *Cyttaromyia freiwaldi* sp. nov.: (A) No. MNHN.F.A71342 (68, counterpart, female), (holotype), habitus, dorsal view; B–F No. MNHN.F.A71342 (70AB, female), (additional material): (B) habitus, dorsal view; (C) head, dorsal view; (D) terminal part of abdomen with ovipositor visible, dorsal view; (E) left wing; (F) right wing. oc, ocelli; thx, thorax.

fossil species especially by point of termination of Sc. In *C. freiwaldi* sp. nov. vein Sc terminating in C before fork of Rs while in other fossil species, excluding *C. reclusa*²⁷, Sc terminating in C just beyond or far beyond bifurcation of Rs. Moreover, in contrast to *C. gelhausi* sp. nov., *C. obdurescens*, *C. princetoniana*, and *C. rayona*, the vein R_1 in *C. freiwaldi* sp. nov. is reduced. Vein $r-r$ (R_2) of *C. freiwaldi* is terminating in R_{3+4} at d-cell level while in *C. fenestrata* and *C. rayona* beyond this level, in *C. vahldieki* even beyond d'-cell level (Supplementary Data S2).

Phylogenetic position of *Cyttaromyia* within *Cylindrotomidae*. The parsimony analysis yielded three equally most parsimonious cladograms, 53 steps long, with consistency index CI=62, RI=66. Their consensus majority rule cladogram is shown in Fig. 6A. The *Cylindrotomidae* clade is supported by six synapomorphies: presence of petiole (character 15, state 0), relationship of R_3 and R_4 (character 17, state 1), position of crossvein m-cu relative to the bifurcation of M_{3+4} (character 18, state 1), shape of d-cell (character 20, state 1), position of tip of A_2 (character 25, state 0), morphology of aedeagus (character 28, state 2). *Cyttaromyia* is supported on consensus tree by the 'presence of supernumerary crossvein connecting vein R_{4+5} with M_1 near its origin, to produce two discal cells' (character 16, state 1). The clade (*Cylindrotominae* + *Stibadocerinae* (Hennig, 1973)⁴³) is supported by two synapomorphies, relationship of M_1 and M_2 (character 14, state 1), position of tip of A_2 (character 26, state 0). The clade [*Phalacrocerca replicata* Linnaeus, 1758⁴¹ + (*Liogma nodicornis* Osten Sacken, 1865⁴⁴ + *Triogma trisculata* Shummel, 1829⁴⁵)] is supported by one synapomorphy, viz. position of crossvein m-cu relative to the bifurcation of Mb (character 19, state 1). *Triogma trisculata* appears as the sister-group to *Liogma nodicornis*, the clade (*Liogma nodicornis* + *Triogma trisculata*) being supported by three synapomorphies: shape of flagellomeres (character 2, state 1), position of R_5 (character 13, state 1), degree of reduction of crossvein r-m (character 24, state 1). The clade *Stibadocerinae* [= (*Stibadocerodes australiensis* Alexander, 1922⁴⁶ + (*Stibadocera bullans* Enderlein, 1912⁴⁷) + (*Stibadocerella pristina* Brunetti, 1918⁴⁸ + *Stibadocerina chilensis* Alexander, 1928⁴⁹))] is supported by two synapomorphies: the number of branches of R_s reaching wing margin (character 9, state 1), relationship of R_3 and R_4 (character 17, state 2) (Supplementary Data S3).

Discussion

The subfamily *Cylindrotominae*²¹ currently contains more species and genera, compared to the *Stibadocerinae*. The oldest record of the extinct genus *Cyttaromyia* is Paleogene, as for the genus *Cylindrotoma*. The fossil record of the *Cylindrotominae* dated back to at least 56.0 Ma²⁹, with no evidences on older occurrences. Other representatives of *Cylindrotominae* are known in the Eocene^{24,25,27,28,31,32,38}, but most of them are strictly modern⁶ (Supplementary Figs. S1, S2).

According to our parsimony analysis (consensus tree), *Cyttaromyia* falls as sister group of all the extant genera of *Cylindrotomidae*, and thus belongs to the stem group of the family, and could correspond to a different subfamily. Also, the extant *Cylindrotominae* appear paraphyletic in respect to the *Stibadocerinae* because the two genera *Cylindrotoma* and *Diogma* fall in an unresolved polytomy with this subfamily plus a clade that contains the other *cylindrotomine* genera. Nevertheless, these results are preliminary and would need to be completed by the addition of characters, in particular molecular.

The *Cylindrotominae* (and also *Cyttaromyia*) have a Holarctic distribution, while the *Stibadocerinae* have a more disjunctive distribution in Indo-Malaysia, Australo-Papua and Southern Neotropics (Taiwan, China, Indonesia, Malaysia, India, Papua New Guinea, Philippines, Australia, and Chile). Such distribution resembles that of an ancient Gondwana group, with 'relic' taxa in Australia and Chile; but the present phylogenetic analysis would contradict this hypothesis, as the only known stem representative of the family is also Holarctic. Further analyses together with discoveries of fossil *Cylindrotomidae* in the Southern Hemisphere shall be necessary to clarify this complex, strange situation.

From a taxonomic point of view, *Architipula* is characterized by the occurrence of vein Sc tip beyond fork of R_s level, subequal to or a little shorter than veins R_{2+3} and R_3 combined, distinctly inclined crossvein m-m between M_{1+2} and M_3 , usually short and straight vein A_2 ⁵¹. *Cyttaromyia* is characterized by the occurrence of two discal cells (d-cell and d'-cell), supernumerary crossvein r'-m' connecting vein R_5 with M_1 near its origin, to produce two discal cells. Some similarities are present in the wing venations of *Cyttaromyia* and *Cylindrotoma*: separate M_1 and M_2 and relatively long vein M_1 . In *Cylindrotoma* the crossvein r'-m' is atrophied, but the base of vein M_1 is strongly arched and only one discal cell (d-cell) is present. In all other *Cylindrotominae*, like *Diogma* (recorded from the Middle Eocene)³², or other genera with a younger fossil record, the crossvein r'-m' is reduced and only one discal cell (d-cell) is present (Fig. 6B,C; Supplementary Figs. S3, S4).

Conclusion

The revision of *Cyttaromyia obdurescens* and the description of two new species *Cyttaromyia gelhausi* sp. nov. and *Cyttaromyia freiwaldi* sp. nov., allowed us to propose a key to the species of this genus. We have also made the first morphological phylogenetic analysis of the *Cylindrotomidae*, with in the rather surprising result of the putative paraphyly of the *Cylindrotominae* and a position of *Cyttaromyia* in the stem group of this family.

Material and methods

The study was based on material from the collection of the Muséum national d'Histoire naturelle (MNHN), Paris (five specimens) and American Museum National History (AMNH) (one specimen). The imprints from sediments of Green River Formation USA (age 50.3–46.2 Ma⁴²) were studied using a Nikon SMZ 1500 stereomicroscope equipped with a Nikon DS-Fi1 camera in University of Rzeszów. The microphotographs and measurements were taken with NIS-Elements D 3.0 software. Drawings were completed by tracing the photographs, nomenclature of wing venation was used³¹.

The Eocene Green River Formation USA (50.3–46.2 Ma) (Rocky Mountains, Colorado, Wyoming, Utah) is one of the most famous Eocene palaeontological sites of the World. The sediments include mainly calcium carbonate, calcite and aragonite. The occurrence of different types of sludge varies with the geological levels. Tipton Shale Member in Greater Green River Basin is the oldest rock formation of the Green River⁵².

Placement of the genus *Cyttaromyia* within *Cylindrotomidae* was tested with the use of Maximum Parsimony (MP) criterion, implemented in TNT 1.5 software package, with the 'Traditional Search' options^{53,54}, with memory to store 99,999 trees, 10,000 replications, with 100 trees to save per replication; utilizing

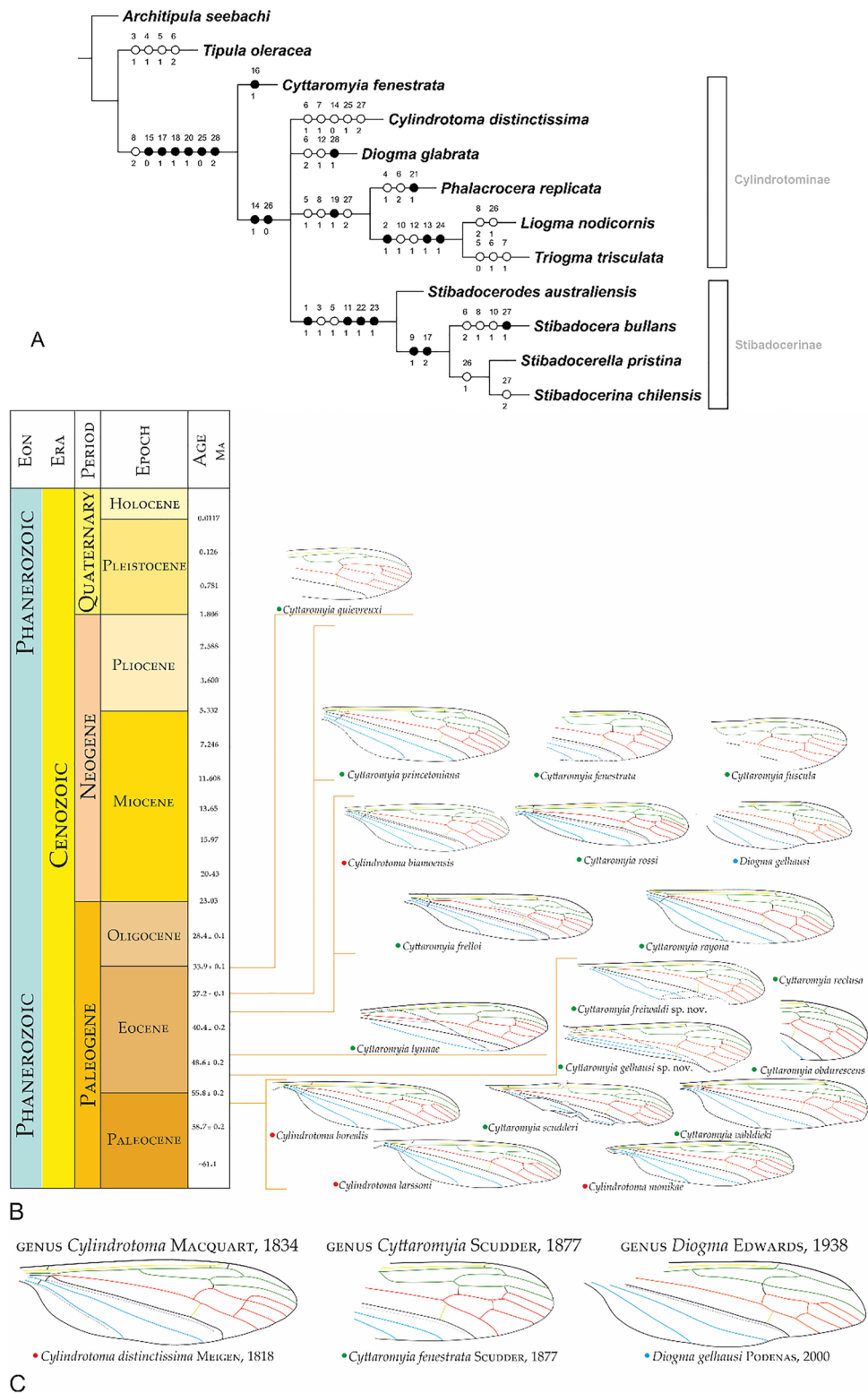


Figure 6. (A) Consensus relationships tree of genera of subfamily Cylindrotominae. Filled circles indicate synapomorphies or autapomorphies; open circles indicate plesiomorphies. Number of character given above circles, states of characters below circles; (B) Wing venation of fossil Cylindrotominae with chronostratigraphic distribution view; (C) Wing venations of representatives of genera: *Cylindrotoma*, *Cyttaromyia*, *Diogma*, represented in fossil record. Wing venation redrawing^{24,25,27–29,31,34,42,50}. Stratigraphic chart according to International Stratigraphic Chart, International Commission of Stratigraphy (v. 2021/05) <https://stratigraphy.org/chart>.

tree-bisection-reconnection (TBR) algorithm and collapsing zero length branches. The type species of extinct and extant genera of family Cyndrotomidae were included in the analysis. *Architipula seebachi* (Geinitz, 1884)⁵⁵—type species of the genus *Architipula* Handlirsch, 1906⁵⁶, was selected as outgroup because the Architipulinae are closely related to Cyndrotominae. *Tipula oleracea* Linnaeus, 1758⁴¹ was used as a type species of the genus *Tipula*, in the family Tipulidae, currently considered as the sister family of the Cyndrotomidae. The morphological data to the matrix were compiled in the Nexus file using Mesquite v. 3.61 build 927⁵⁷. All 28 characters of the imagines used in the analysis were treated as unordered and unweighted (Supplementary Table S2). Equal weighting analysis (EW) was performed⁵³; the trees received were viewed and their features studied using WinClada 1.00.08 and ASADO 1.61, with Unambiguous Changes Only, Fast Optimization (ACCTRAN) and Slow Optimization (DELTRAN) options^{57–59}. Tree files received were adjusted using Corel Draw X3 and Photo-Paint Software. The 28 morphological characters of the imago observed in the fossil and recent material and used for analysis are listed below. The data matrix given is partly based on used morphological features^{8,21,34,41,43–45,50,56}.

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I.K.-K. conceived and designed the study, lead and performed the data analysis, interpretations and writing, making photographs, graphical figures, analysis; correspondence. A.N. writing and corrections of the manuscript. J.S. corrections of phylogenetic analysis methods. W.J.-S. making photographs and drawing, graphical figures, analysis. W.K. corrections of the manuscript.

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