



A second genus of Progonocimicidae (Hemiptera: Coleorrhyncha) from the Early Toarcian of Luxembourg

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Abstract

Liassoprogonocimex bascharagensis gen. et sp. nov. of Progonocimicidae (Coleorrhyncha: Progonocimicomorpha) are described. It is one of two genera from the Early Jurassic of Luxembourg, the other being *Indutionomarus* Szwedo, 2011. This new occurrence increases the morphological diversity of Coleorrhyncha during the Mesozoic and helps us to discuss the systematics of extinct Coleorrhyncha as well as their evolutionary history.

Keywords: Insecta, Jurassic, moss bugs, fossil record, Progonocimicomorpha

Introduction

Coleorrhyncha Myers & China, 1929 or moss bugs is a hemipteran lineage considered as a relict group, represented nowadays by only one family (i.e., Peloridiidae Breddin, 1897) with only 37 species in 17 genera, distributed only in the sub-Antarctic rainforests of the southern Hemisphere (Burckhardt, 2009). Ye *et al.* (2019) has linked the biogeographical pattern of these moss bugs to the successive break ups of Gondwana throughout the Late Cretaceous.

Although extant Coleorrhyncha are not especially diverse, their fossil record indicates that their past biodiversity was much greater, with numerous occurrences during the Mesozoic (Wang *et al.*, 2009). The current classification of Coleorrhyncha comprised two infra-orders. Firstly, the extinct Progonocimicomorpha Popov, 1980 and its unique family the Progonocimicidae Handlirsch, 1906 ranging from Permian to Late Cretaceous

with two recent occurrences from the mid-Cretaceous Kachin amber (Jiang *et al.*, 2019; You *et al.*, 2024). Secondly, the extant Peloridiomorpha Popov, 1980 that encompassed two extinct families: Hoploridiidae Popov & Shcherbakov, 1991 and Karabasiidae Popov, 1985; and the extant family Peloridiidae. Recently, Burckhardt *et al.* (2023) described the earliest family of Coleorrhyncha with Permoridiidae Burckhardt *et al.* 2022 from Early Permian, pushing back the origin of moss bugs and trying to define putative synapomorphies for Coleorrhyncha, leaving the new family unranked within Coleorrhyncha systematics. According to recent phylogenomic analyses, moss bugs may well predate the Permian; they may have appeared during the Early Carboniferous [as is the case with all major hemipteran lineages, approx. 350 million years ago (Johnson *et al.*, 2018)].

The Progonocimicidae are known from the Permian to mid-Cretaceous (Popov & Shcherbakov, 1991; Dong *et al.*, 2014; Jiang *et al.*, 2019; You *et al.*, 2024) with more than 20 genera distributed within two subfamilies: Progonocimicinae Handlirsch, 1906 and Cicadocorinae Becker-Migdisova, 1958. The first subfamily is known from the Permian to the Jurassic, whereas Cicadocorinae are Jurassic to Cretaceous. In particular, eight Cretaceous genera and 11 species are described (see Jiang *et al.*, 2019: fig. 7). Several specimens of Progonocimicidae have been reported among the rich Crato Formation but have never been formally described (Bechly & Szwedo, 2007). They represent the first records of Coleorrhyncha in the New World. The presence of this lineage in the New World is crucial to undercover the biogeographical history of moss bugs. However, the Jurassic seems to be a linchpin point in the evolutionary history of the Coleorrhyncha with the

decline of their representatives in Europe, probably linked to climate changes during this period. Subsequently, depicting the diversity of Coleorrhyncha in the Jurassic is important to better understand their decline. To this end, we describe the progonocimicid new genus and species *Liassoprogonocimex bascharagensis* gen. and sp. nov. from the Toarcian of Luxembourg, the second genus and species in this area after *Indutionomarus treveriorum* Szwedo, 2011. This new occurrence helps to discuss the general taxonomy and evolutionary history of Coleorrhyncha.

Material and methods

The new fossil was collected by one of us (MH) in the carbonate nodules in Bascharage, Grand-Duché-du-Luxembourg (see map in Ansorge, 2003: Fig. 1). The carbonate nodules are intercalated in the bituminous shales of the Elegantulum subzone, Lower Serpentinum Zone (Thuy *et al.*, 2025). This outcrop has already provided many insects (Nel, 1989; Nel & Henrotay, 1992, 1994; Nel *et al.*, 1993; Henrotay *et al.*, 1997; 1998; Nel *et al.*, 2003, Fleck *et al.*, 2003; Szwedo, 2011).

The specimen was photographed with a Canon EOS 70D camera equipped with a Novoflex Micro-Tube M26 achromatic lens and a Mitutoyo M Plan Apo 5×/0.14 microscope lens. The camera was mounted on a semi-automatic Cognis Rail macro Stack Shot controlled by Helicon Remote software. We also conducted UV-a light imaging of the specimen following the setup and protocol described in Boderau *et al.* (2024).

Photographs were digitally stacked using Helicon Focus 6.7 software and were exported in TIF format. We used a camera lucida mounted on an Olympus SZX9 stereomicroscope to perform the drawing lines. Final images and original drawing lines were enhanced using Pixelmator Pro 3.6.12 Archipelago software.

We follow the systematic framework of Szwedo (2018) for the Coleorrhyncha higher ranks classification, the taxonomic attribution within Progonocimidae follows Popov & Shcherbakov (1991), Dong *et al.* (2014), and the Palaeobiology Database (<https://paleobiodb.org/#/>; last access on January, 7th 2025). The forewing venation nomenclature follows the general pattern for Acercaria proposed by Nel *et al.* (2012) modifying the interpretations of Popov & Shcherbakov (1991). We adopted the cell nomenclature of Bourgoin *et al.* (2015).

Forewing venation abbreviations are as follow: (CA) costa anterior; (CP) costa posterior; (Pc) praecosta; (bScP) basal portion of the subcosta posterior; (ScP) subcosta posterior; (RA) radius anterior; (RP) radius posterior; (M) media; (CuA) cubitus anterior; (CuP) cubitus posterior;

(PCu) postcubitus vein; (A_1) first anal vein. Each crossvein between two main veins is written in lower case.

Systematic palaeontology

Order Hemiptera Linnaeus, 1758

Suborder Coleorrhyncha Myers & China, 1929

Infraorder Progonocimicomorpha Popov, 1980

Superfamily Progonocimicoidea Handlirsch, 1906

Family Progonocimicidae Handlirsch, 1906

Subfamily uncertain

Genus *Liassoprogonocimex* gen. nov.

Type species. *Liassoprogonocimex bascharagensis* sp. nov.; by monotypy.

Etymology. The generic name is a combination of ‘Liassic’, other name for Lower Jurassic and ‘Progonocimex’, the type genus of the family. Gender: masculine.

Diagnosis. Tegmen with ScP + RA₁ diverging before end of clavus; distance between ScP+RA₁ and RA₂ nearly the same as distance between RA₂ and RP at their bases; RA₂ not continuous with R-stem; *rp-m* crossvein very short; alignment of basal part of CuA₂, *m-cua*, and basal part of M₁₊₂; PCu strongly diverging from CuP before its fusion with A₁; PCu₂+A₁ very short, ending into posterior margin of wing well basal to apex of clavus; presence of a *m-cua* crossvein between M₃₊₄ and CuA₁.

Liassoprogonocimex bascharagensis sp. nov.

(Fig. 1)

Material. Holotype MNHN.F.C02460 housed in the collection of Palaeontology Department in the Muséum national d’Histoire naturelle (Paris, France).

Etymology. The specific epithet is referring to the type locality from where the specimen originated.

Diagnosis. As for the genus (*vide supra*).

Locality and horizon. Bascharage; Grand-Duché-du-Luxembourg (Henrotay *et al.*, 1998: Fig. 1); lower Toarcian; Lower Jurassic.

Description. Tegmen *ca.* 2.04 mm long, rounded apically; apex of clavus reaching roughly half of total tegmen length; postclaval area slightly wider than preclaval area; CP+PC long reaching end of clavus; basal cell closed apically by a long transverse *cua-cup*; basal portion of ScP (bScP) very weak, reaching R+M+CuA before basal half of clavus; ScP+R diverging from M+CuA and M diverging from CuA at level of basal half of clavus; ScP+R stem twice shorter than M-stem and CuA-stem; ScP+RA₁ nearly transverse and diverging from RA₂+RP slightly before level of end of clavus; RA₂ subparallel to

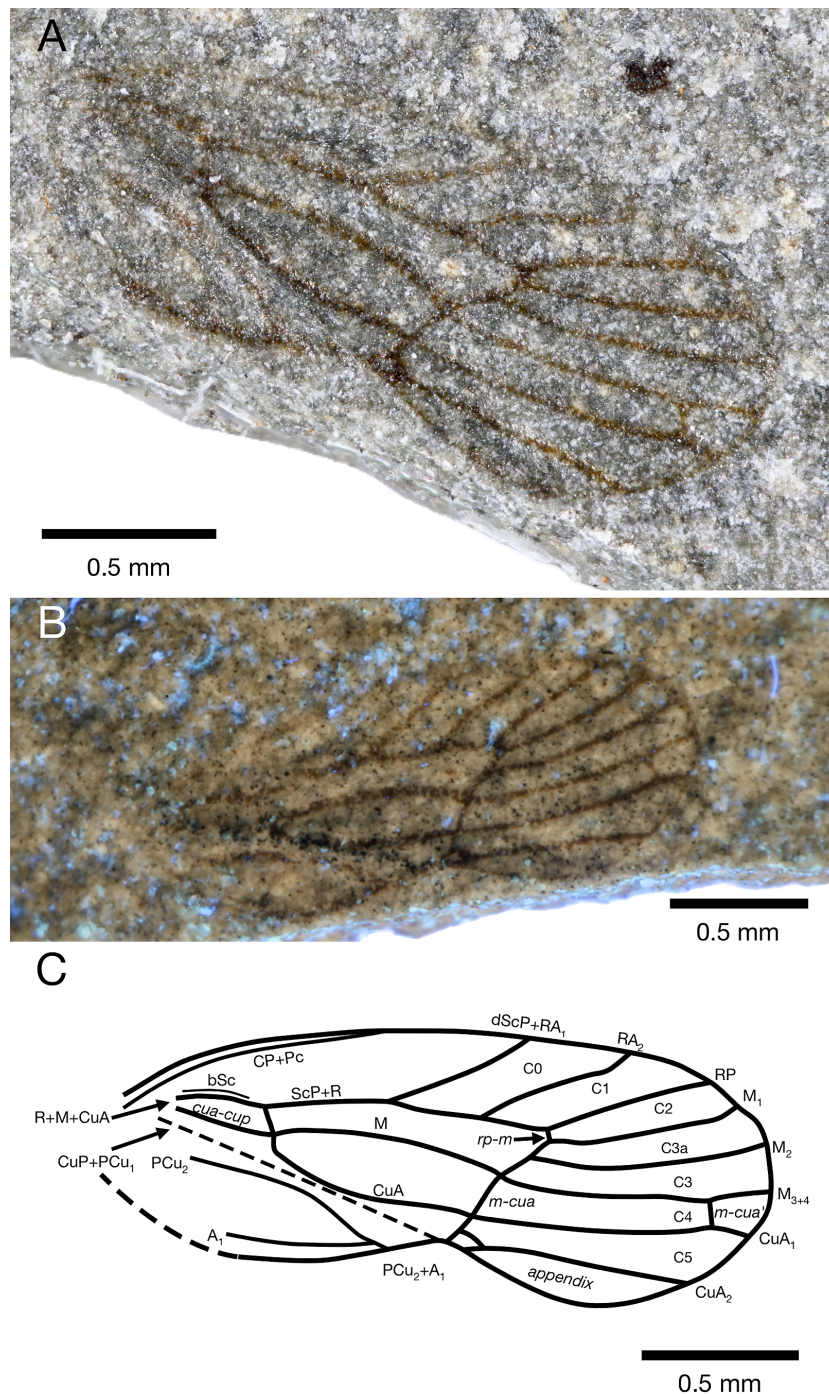


FIGURE 1. *Liassoprogonocimex bascharagensis* gen. et sp. nov. holotype MNHN.F.C02460. **A**, Photograph of forewing. **B**, Photograph of forewing under UV-light. **C**, Interpretive drawing line.

RP, diverging closer to ScP+RA₁ than to RP; M trifurcated, each fork parallel to each other, without a common stalk between M₁₊₂ and M₃₊₄ but with a common stalk of M₁ and M₂, *r-m* between RP and M₁ very short, *ca.* 0.005 mm; terminal M₃₊₄ distinctly longer than terminals RP, M₁, and M₂, subequal in length to terminal CuA₁; terminals of M anteriorly curved apically; terminals of CuA slightly curved apically towards posterior margin of wing; M₃₊₄ and CuA₁ separated by a short and thick crossvein; CuA₂ not parallel to appendix margin, this margin being

medially curved; CuP+PCu₁ straight forming distinct claval suture, adpressed to CuA, distinctly thinner than other main veins; PCu₂ subparallel to CuP+PCu₁, strongly curved distally and fused with A₁ *ca.* 0.5 mm from wing base, distal vein PCu₂+A₁ extremely short, about ten times shorter than free portion of A₁, and ending well before apex of clavus; cell C0 elongate, subequal to cell C1; cells C3a to C5 subrectangular, with C5 the longest and the widest cell.

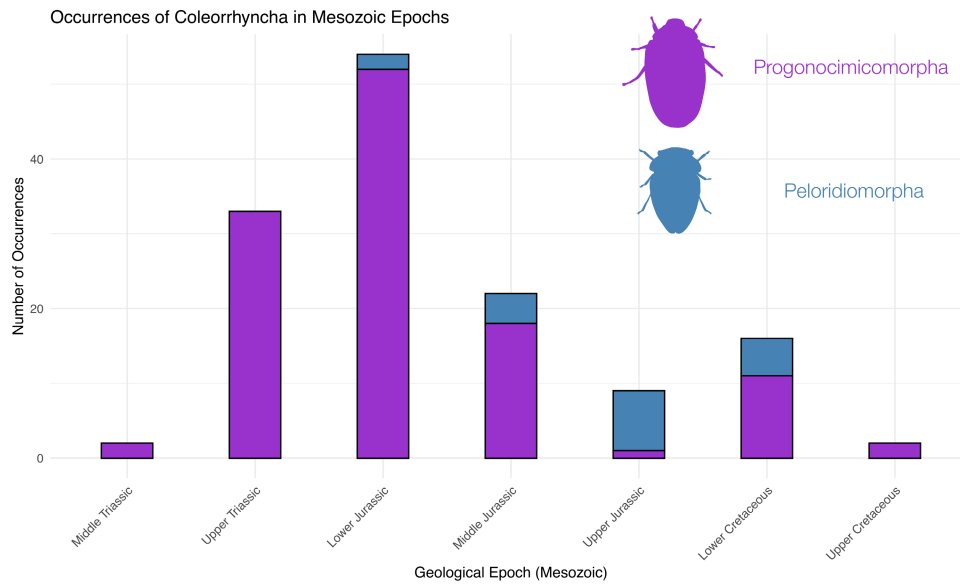


FIGURE 2. Diversity of Coleorrhyncha throughout Mesozoic. Data compiled from ‘The Paleobiology Database’ (<https://paleobiodb.org/#/>). Insect silhouettes drawn by M.B.

Discussion

The new fossil can be attributed within Progonocimicidae based on the tegmen with eight apical cells, without areolae and long ScP incorporated into R-stem (Popov & Shcherbakov, 1991; Wang *et al.*, 2009). Progonocimicidae are divided into two subfamilies: Cicadocorinae Becker-Migdisova, 1958 and Progonocimicinae Handlirsch, 1906.

Popov & Shcherbakov (1991: 217) separated these two groups as follows: (1) A_1 not shorter than half of claval suture in Progonocimicinae *vs.* shorter in Cicadocorinae; (2) PCu_2 nearest to the suture before joining A_1 in Progonocimicinae *vs.* diverging from suture before joining A_1 in Cicadocorinae; (3) precostal carina [CP+Pc] deflected dorsad or obsolete in Progonocimicinae *vs.* horizontal or deflected ventrad (if dorsad, base of costal area narrow) in Cicadocorinae; (4) base of costal area not narrow in Progonocimicinae; (5) R-stem not continued by RP in Progonocimicinae *vs.* continued in Cicadocorinae; (6) and/or $dScP+RA_1$ at least partly curved anteriorly in Progonocimicinae *vs.* straight or curved posteriorly in Cicadocorinae.

Character (1) is not preserved in some taxa (*e.g.*, *Pelorisca* Popov & Shcherbakov, 1991 and *Woottonia* Popov & Shcherbakov, 1991); furthermore, its definition is ambiguous as A_1 is fused for a long distance with posterior margin of wing in its basal part in these wings. In the new fossil, its distal free part is half of claval suture, which is in-between the two states.

Character (2) is also not preserved in some taxa (*e.g.*, *Pelorisca* and *Woottonia*). In the genus

Cicadocoris Becker-Migdisova, 1958 (type genus of the Cicadocorinae), PCu_2 is straight and, at most, very slightly diverging from the suture in its distal part (Becker-Migdisova, 1958; Dong *et al.*, 2014; Jiang *et al.*, 2016; Jiang & Huang, 2017; Xu & Huang, 2023). In the new fossil, it is strongly diverging from the suture.

Character (3) is quite difficult to determine whether it is deflected dorsal *vs.* ventral in compression fossils. Its state is unknown in the new fossil.

Character (4) can be present in both groups after Popov & Shcherbakov (1991).

Character (5) is also difficult to estimate because the angle between the radial stem and RP can be pronounced. In the new fossil, it is very weak, weaker than in the progonocimicine *Pelorisca*, suggesting an attribution to the Cicadocorinae.

Character (6): in the new fossil, $dScP + RA_1$ is distinctly curved anteriorly. This character would support an attribution to the Progonocimicinae. But in *Progonocimex* Handlirsch, 1906, $dScP+RA_1$ is as straight as in some species attributed to *Cicadocoris*, *e.g.*, *C. henannensis* Xu & Huang, 2023 (Popov & Wootton, 1977: fig. 1, 2; Xu & Huang, 2023).

As a result, these characters have three defaults: one is not present in the type genus of the Cicadocorinae, some are ambiguous, and they are not polarized. Thus, the monophyly *vs.* paraphyly of these two groups of taxa are not demonstrated. The new fossil would be ‘intermediate’ between them. Therefore, we avoid putting it into a precise subfamily. These would need to be redefined after a phylogenetic analysis. This is confirmed by Dong *et al.*

(2014) who did not recover the monophyly of the two subfamilies in a partial phylogeny of the Coleorrhyncha.

Therefore, we need to compare it with all the genera of Progonocimicidae. The new fossil differs from *Progonocimex* in the distinctly curved PCu₂ vs. straight (Popov & Wootton, 1977). It differs from *Woottonia* and *Heterojassus* Evans, 1961 by the appendix separated by straight vein subparallel to CuA₁ (Popov & Shcherbakov, 1991) Affinities with *Pelorisca*, *Actinoscytina* Tillyard, 1926, *Triassodoecus* Evans, 1963, and *Heteroscytina* Evans, 1963 are excluded based on the imaginary continuation of R-stem closer to RP than RA₂. Unlike *Indutionomarus* Szwedo, 2011, the new specimen has the crossvein rp-m between RP and M₁ vs. between RP and M₁₊₂ (Szwedo, 2011). The new fossil differs from *Platyscytinella* Evans, 1956 by lacking a common stalk formed by M₂ and M₃₊₄ (Evans, 1956).

Mesoscytina Tillyard, 1919 and *Eocercopis* Handlirsch, 1939 differ from the new fossil in the M-stem and R-stem forming curved lines vs. straight (Lambkin, 2016; Zhang *et al.*, 2024). In *Mesocimex* Hong, 1983 and *Eocercopis*, PCu₂ is parallel to CuP+PCu₁ up to its apex (Wang *et al.*, 2009), vs. strongly diverging apically in the new fossil. Also *Mesocimex* has a longer PCu₂+A₁ than in the new fossil and ending at apex of the clavus (Szwedo, 2011). In *Onokhoia* Popov, 1988 and *Popovus* Özdikmen & Demir, 2007 (substitute name for *Progonus* Popov, 1986), PCu is less diverging from CuP than in the new fossil, and PCu₂+A₁ ends at apex of clavus vs. reaches posterior margin of wing basal of apex of clavus in the new fossil (Popov, 1986, 1988). *Archicercopis* Handlirsch, 1939 has PCu close to claval fracture, and M₁₊₂-forks beyond r-m. *Cicadocoris* also has PCu₂+A₁ ending at apex of clavus and PCu₂ straight. Its type species also strongly differs from the new fossil in the shape of the distal part of stem of M distal of m-cua and position of M₃₊₄ (Becker-Migdisova, 1958: fig. 2), even if Dong *et al.* (2014) described another species with a pattern of M and CuA similar to that of the new fossil.

Hexascytina Wootton, 1963 is based on a small fragment of a forewing. It is supposed to have a RP making a very strong zigzag, and the crossvein rp-m very long (Wootton, 1963). *Microscytinella* Wootton, 1963 and *Platyscytinella* Wootton, 1963 also have a very long rp-m. *Popovigocimex* Martins-Neto & Gallego, 2003 strongly differs from the new fossil in the presence of five parallel anterior branches of ScP+R (Martins-Neto *et al.*, 2003). *Yurigocimex* Martins-Neto & Gallego, 2003 differs from the new fossil in the base of M + CuA opposite the base of ScP+RA₁, and very short r and m cells. *Ovicimex* Hong & Wang, 1990 has a forewing venation strongly different from those of the other Progonocimicidae and probably does not belong to this family (Hong & Wang, 1990: figs

66, 67). *Triscytina* Evans, 1956 strongly differs from the new fossil in the absence of alignment of the basal part of CuA₂, m-cua, and basal part of M₁₊₂, plus PCu closely parallel to CuP (Evans, 1956). *Triassodoecus* Evans, 1963 is based on an incomplete forewing. It differs from the new fossil in PCu parallel to CuP and PCu₂+A₁ ending at distal angle of clavus (Evans, 1963). *Heteronella* Evans, 1961 is based on an incomplete forewing, and its affinities are uncertain. It differs from the new fossil in the vein ScP+RA emerging at level of emergence of M + CuA (Evans, 1961).

Gakasha Jiang, Wang & Szwedo, 2019 strongly resembles the new fossil in the shape of PCu strongly diverging from CuP in its distal part, PCu₂+A₁ ending in posterior wing margin well basal to apex of clavus, pattern of the distal parts of veins RP, M, and CuA. It differs from the latter in distance between ScP+RA₁ and RA₂ much longer than distance between RA₂ and RP at their bases vs. of nearly the same, and PCu₂+A₁ much longer than in the latter (Jiang *et al.*, 2019).

Awanis You *et al.*, 2024 shares with our new fossil the short rp-m but differs from it based on R-stem not continued by RP; very short cua-cup. Additionally, the forking of M is strikingly different, in the new fossil M-stem is continuous with M₃₊₄ branch whereas M-stem is between M₂ and M₃₊₄ branches in *Awanis*. Furthermore, C5 is almost rectangular in *Awanis*, while C5 is enlarged distally in *Liassoprogonocimex* **gen. nov.**

In addition to the aforementioned differences, this new fossil exhibits unique features such as: a C0 cell more anterior than in any other Progonocimicidae genera with the division of ScP+RA₁ occurring before the end of the clavus; the rp-m crossvein is very short with the base of RP and M₁ very close to each other, and the presence of a distal crossvein between M₃₊₄ and CuA₁, a feature never recorded in any Progonocimicidae. We considered that all these differences are sufficient to erect a new genus among Progonocimicidae with *Liassoprogonocimex* **gen. nov.**

Shcherbakov (1996) proposed that 'Progonocimicidae derive from the 'Ingruidae' (in the paraphyletic unit Ingruomorpha Szwedo, 2018). The Ingruidae putatively constitutes the 'ancestors' of Coleorrhyncha lineages (Popov & Shcherbakov, 1996). Such hypotheses of 'descendant' taxa in Coleorrhyncha are often cited but should be taken very carefully as a lineage cannot be derived from another, otherwise they are all paraphyletic. "Ingruomorpha" representatives are probably nested within Euhemiptera, with which they share the PCu fused with A₁, and the typical Y-shaped vein in the claval area. However, they strongly differ from the Progonocimicidae in the forking of the CuA vein, with two terminals very close to each other, whereas in Progonocimicidae the two

branches of the CuA vein are very long and, in most cases, subparallel to each other.

Progonocimicidae are currently grouped with the Coleorrhyncha. However, as discussed by Burckhardt *et al.* (2023), there is no clear synapomorphy for moss bugs in wing venation. The Progonocimicidae do not share with Permoridiidae and Peloridiidae the ScA well separated from C with branches and an elongate and well-defined ScP. They also differ from Karabasiidae and Hoploridiidae by the ScP incorporated into R. Additionally, Popov & Shcherbakov (1996), and Shcherbakov & Popov (2002: 147, 148) proposed a list of ‘putative synapomorphies’ for Coleorrhyncha with: ‘suprantennal ledge continuous below median ocellus; forewings overlapping in repose (left one uppermost); forewing with enlarged posterior apical cells, short transverse CuA₂ and broad appendix; coupling fold very short in forewing; fore and middle tarsi two-segmented; pygophore barrel-shaped, parameres protruding and elbowed’. These characters are very difficult to observe in most Progonocimicidae fossils or not preserved. However, the unique specimen described from mid-Cretaceous Kachin amber, *Gakasha calcaridentata* Jiang, Wang & Szwed, 2019, definitely shows a suprantennal ledge continuous below median ocellus, and two-segmented fore and middle tarsi (but a feature present in numerous different acercarian lineages). The wing venation features proposed by Popov & Shcherbakov (1996) are not sufficient as they are present in other clades of Hemiptera. This highlights that a significant effort of systematics within Coleorrhyncha is required to delimit phylogenetically robust units.

The evolutionary history of Coleorrhyncha is complex. They originated more than 295 Myr ago (Burckhardt *et al.*, 2023) but they are quite uncommon in Late Paleozoic to Early Mesozoic outcrops, most notably because these periods are less documented (Schachat & Labandeira, 2021). The Coleorrhyncha are diverse in rocks of Upper Triassic to Middle Jurassic age (Fig. 2), corresponding to a high diversity of Progonocimicidae. The Progonocimicidae are widely spread between Triassic to Early Cretaceous, with occurrences in Australia, Eurasia, and South America (Martins-Neto *et al.*, 2003). They strongly declined during the late Jurassic, in terms of occurrences. The Jurassic ones are currently restrained to Eurasia, but in the Cretaceous they are recorded in South America, Burmese paleo-island in the middle of the Tethys Ocean, and Eurasia. The last occurrence of Progonocimicidae is from the mid-Cretaceous Kachin amber, suggesting an extinction of the family related to the global floristic changes occurring during the late Cretaceous with the rise of the angiosperms and the subsequent decline of gymnosperms, their putative host-plants (Popov & Shcherbakov, 1996). But future

discoveries in the latest Cretaceous could change this scenario.

The other main coleorrhynchan lineage is the Peloridiomorpha, known from the early Jurassic to the early Cretaceous (Fig. 2), their geographic distribution is limited to Asia, and they are morphologically closer to the only extant family of Coleorrhyncha (Peloridiidae). Recently, Ye *et al.* (2019) investigated the timing of diversification in Coleorrhyncha and the biogeographical history of Peloridiidae. However, their ingroup consists of the extant Peloridiidae, supposedly diverging from the outgroups (Heteroptera, Fulgoromorpha, Cicadomorpha, Sternorrhyncha) during the Carboniferous. This early divergence could strongly impact on the phylogenetic reconstruction and hamper molecular time divergence analyses. Additionally, the biogeographical analysis was only based on the Gondwanan distribution of Peloridiidae, but we illustrated that the Coleorrhyncha were widely spread throughout the Mesozoic. It would be interesting to have a total-evidence approach to study the diversification and biographical history of the Coleorrhyncha clade, by including the Progonocimicomorpha and the extinct Peloridiomorpha, to overcome the limitations due to their extant relict diversity. However, most of the extinct Coleorrhyncha are known by forewings, or poorly preserved bodies, which could lead to an issue of morphological data incompleteness between extant and extinct Coleorrhyncha.

Conclusion

We describe the second genus and species of Progonocimicidae of the Early Toarcian of Luxembourg. Intensive fieldwork in this locality has delivered interesting insect fossils, but this outcrop will need to be investigated, namely because a putative and particular entomofauna, linked to a short and fast period of global warming, occurring synchronously with the Toarcian-Oceanic Anoxic Event. Depicting the biogeographical history of Coleorrhyncha is challenging because of their relict diversity, hampered by their lacunar fossil record, which needs additional discoveries to propose enhanced hypotheses.

Acknowledgments

We thank two anonymous referees for their comments on the first version of the paper. We are also grateful towards Antoine Mantilleri and Christophe Rivier (MNHN) who gave access to the Entomology laboratory’s photographic equipment, and for their kind advice on photography

acquisition and to Valérie Ngô-Muller (UPC) for early discussions on the specimen. This work is a contribution of the 2D/3D photonics lab of ISYEB lead by RG, with MNHN and ISYEB funding.

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