A Triassic-Jurassic window into the evolution of Lepidoptera

Timo J. B. van Eldijk,¹ Torsten Wappler,² Paul K. Strother,³ Carolien M. H. van der Weijst,¹ Hossein Rajaei,⁴ Henk Visscher,¹ Bas van de Schootbrugge¹*

On the basis of an assemblage of fossilized wing scales recovered from latest Triassic and earliest Jurassic sediments from northern Germany, we provide the earliest evidence for Lepidoptera (moths and butterflies). The diverse scales confirm a (Late) Triassic radiation of lepidopteran lineages, including the divergence of the Glossata, the clade that comprises the vast multitude of extant moths and butterflies that have a sucking proboscis. The microfossils extend the minimum calibrated age of glossatan moths by ca. 70 million years, refuting ancestral association of the group with flowering plants. Development of the proboscis may be regarded as an adaptive innovation to sucking free liquids for maintaining the insect’s water balance under arid conditions. Pollination drops secreted by a variety of Mesozoic gymnosperms may have been non-mutualistically exploited as a high-energy liquid source. The early evolution of the Lepidoptera was probably not severely interrupted by the end-Triassic biotic crisis.

INTRODUCTION

Lepidoptera (moths and butterflies) represent one of the most admired and studied insect groups, not least for their remarkable associations with flowering plants. However, despite their important role in terrestrial ecosystems, the early evolutionary history of these insects remains murky and mired in an exceedingly poor fossil record (1). Current evolutionary concepts are largely based on molecular phylogenetic analyses, suggesting that Lepidoptera diverged from their sister group Trichoptera (caddisflies) during Permian (2, 3) or (Late) Triassic (4–6) times. The large discrepancies in divergence time are mainly due to competing molecular dating methods and the choice of calibration fossils for providing age constraints. However, in any case, age estimates are substantially older than the oldest known stem-group lepidopteran fossil Archaeolepis mame [Early Jurassic; Sinemurian; ca. 195 million years ago (Ma); Dorset, UK] (7) and the oldest known crown-group representative Parasabatinca atrimacrai (Early Cretaceous; Barremian; ca. 129 Ma; Lebanon) (7).

To contribute to a possible reduction of the gap between molecular and fossil dates, we explore for the first time the phylogenetic potential of dispersed lepidopteran wing scales encountered in sedimentary organic matter. Lepidoptera are characterized by, and named after, their dense covering of chitinous scales on bodies, legs, and wings. Detached scales can be transferred by wind and water action to depositional areas for burial in terrestrial or even marine sediments, from which they may be recovered by palynological methods (8, 9). Because the structure of the scales, particularly the wing scales, is taxonomically informative (10), well-preserved fossil specimens could have clade-level morphological characteristics relevant to more accurate calibration of divergence-time estimates in molecular lepidopteran phylogenies. We studied fossilized scales encountered as rare palynological elements (Fig. 1) in Triassic-Jurassic boundary sediments from the cored Schandelah-1 well, drilled in northern Germany near Braunschweig.

RESULTS

The scales were found discontinuously within a 26-m stratigraphic interval embracing the Triassic-Jurassic (Rhaetian-Hettangian) transition (Fig. 2). About 70 scales and scale fragments, in various states of degradation, could be analyzed. Exceptionally well-preserved specimens were recovered from just above the palynologically defined Triassic-Jurassic boundary. Taxonomic identification of the fossil scales has been based on relevant literature data on scale morphology and structure of extant Lepidoptera and other scale-bearing hexapods, supplemented by the analysis of additional scanning electron microscopy (SEM) images (see the Supplementary Materials). Our survey of extant scale types and a compilation of the principal morphological characteristics (Table 1A) suggest that most hexapods, other than Lepidoptera, may be excluded as a source for the fossil scales (Table 1B). There is also little affinity with the scale types of the extinct neuropteran family Kalligrammatidae (11) and Tarachoptera, a recently proposed extinct order of the Amphiesmenoptera (12).

Fig. 1. Lepidopteran scales in palynological preparations, as seen in transmitted light. (A) Serrated scale from the Hettangian [316.70 m below surface (mbs)]. (B) Scale with a rounded apical margin from the Rhaetian (337.50 mbs). Scale bars, 20 μm.
Highlighting their diversity, the encountered scales could be categorized into four broad morphological groups with different overall shapes and scale margins, including triangular to rounded margins, serrated margins, elongated shapes, and fringed margins (Fig. 3, A to F). On the basis of SEM analyses and histological sections of well-preserved specimens, we identified two distinctive scale types showing decisive evidence of lepidopteran affinity. Type I scales (Fig. 4, A and B) are "solid" (fused upper and lower laminae) and have an unserrated, rounded, or slightly triangular apical margin. Areas between longitudinal ridges on the upper surface are sculptured with a faint herringbone pattern. In marked contrast, type II scales (Fig. 4, C to E) are "hollow" (upper and lower laminae connected by columnar trabeculae) and usually have serrated apical margins. The inter-ridge areas of the upper lamina contain a relatively dense covering of cross ridges, interspersed with small circular perforations. The top of the longitudinal ridges is adorned with microribs. A single type II scale also bears oblique apical crests (Fig. 4B). In addition to these lepidopteran scales, several scale types, which we have been unable to attribute to any of the modern scale-bearing hexapod groups, were encountered. The most characteristic of these unidentified types is a fringed, probably solid, scale without herringbone patterns (Fig. 3F and 4F).

The general morphology and internal structure of both type I and type II scales is consistent with scale morphologies found in the so-called non-ditrysian Lepidoptera, a paraphyletic group of extant families reflecting early lepidopteran phylogeny (13) but embracing only 1 to 2% of all lepidopteran species. The characteristic herringbone patterns, as observed in type I scales, are only present today in those lineages that diverge at the three earliest nodes in this phylogeny: Micropterigidae, Agathiphagidae, and Heterobathmiidae (10). The fossil scales share their solid structure with both Micropterigidae and Heterobathmiidae but differ from the hollow wing scales of the Agathiphagidae (10). These three primitive families represent relict lineages of small moths with mandibulate, chewing mouthparts (14). Morphologically related Jurassic fossils have been included in four extinct mandibulate families: Archaeolepidae, Eolepidopterygidae, Mesokristenseniidae, and Ascolelopodopterygidae (15). Only the Archaeolepidae, with the single-wing species A. mane, are characterized by the presence of relatively well-preserved, probably solid, wing scales, but no herringbone pattern has been observed (16). The Rhaetian-Hettangian type I scales would corroborate recognition of a successful and probably diversified clade of mandibulate Lepidoptera during the Early Mesozoic.

The affinity of type II scales is clearly associated with the morphological clade Coelolepida, defined principally by hollow wing scales with perforations in the inter-ridge areas of the upper lamina, characteristic of the vast majority of extant Lepidoptera (10). Serrated apical margins are additional evidence for a coelolepidan relationship. On the basis of the number and size of the perforations and the density of cross ridges, numerous type II scales show a resemblance to scales of non-dytisian Lepidoptera present.
families of the Coelolepida, such as the Acanthopteroctetoidae, Adelidae, Incurvariidae, and Cecidosidae (10). Unfortunately, morphological information on modern wing scales is still too restricted to judge whether the fossil material originates from crown- or stem-group members of the Coelolepida. Modern occurrences of oblique apical crests, noted in a single type II specimen, are limited to the non-coelolepidan Eriocraniidae (10). Nonetheless, the fossil scale is clearly hollow and has the hallmark perforations of the Coelolepida. Because the family Eriocraniidae is

<table>
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<th>Group</th>
<th>Apical margin</th>
<th>Cross ridges</th>
<th>Microribs</th>
<th>Structure</th>
<th>Perforations</th>
<th>Herringbone pattern</th>
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<tr>
<td>A</td>
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</tr>
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<td>Dense</td>
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usually considered to be the sister group to the Coelolepida (13), a tantalizing possibility is that this scale represents a stem-group coelolepidan, still retaining a character from its non-coelolepidan ancestors.

DISCUSSION

Although their monophyly is questioned (17), coelolepidan lineages are nested in the Glossata (13), the huge clade that includes all moths and butterflies having a sucking proboscis, a sophisticated feeding device fundamentally adapted to fluid uptake from droplets and surface films (14). On the basis of an undescribed larva from Lebanese amber characterized by a spinneret, a silk-spinning organ apomorphic of Glossata (18), the minimum age of fossil representatives of this clade could thus far be constrained with any certainty as mid Early Cretaceous (Barremian; ca. 129 Ma) (1). The Rhaetian-Hettangian scale record pushes back the calibrated age by ca. 70 million years, supporting molecular estimates of a mid Late Triassic (Norian; 212 Ma) divergence of the Glossata from non-glossatan, mandibulate ancestors (4). The record falsifies an Early Cretaceous (Berriasian; ca. 141 Ma) origination of the Glossata, which followed from the comprehensive phylogenomic treatment of insects by Misof et al. (6).

The transition to exclusively feeding on liquids was most likely an evolutionary response to widespread heat and aridity during the Norian (19). When flying in dry air, the high ratio of surface area to volume inherent in the small body size of basal moths would intensify evaporative losses of body moisture (20). Because free liquid drinking is an efficient technique to replenish lost moisture and survive desiccation stress, substitution of mandibulate mouthparts by a sucking proboscis could be seen as an adaptation to adequate maintenance of body water balance of small, short-lived moths. Like in the most basal extant lineages, such as the Eriocraniidae and the Mnesarchaeidae (14, 21), short and simply composed proboscides of ancestral Glossata must have been used initially to drink from water droplets or sap from injured leaves.

In association with the rise and diversification of crown-group angiosperms during the Cretaceous (22), possession of the proboscis facilitated feeding from concealed floral nectaries, thus paving the way
for the evolution of the multitude of pollination mutualisms between Lepidoptera and flowering plants. In modern non-ditrysian Glossata, long proboscides adapted to nectar feeding occur in members of the more advanced clade Eulepidoptera (14). However, preceding an angiosperm connection, early radiation of non-ditrysians evidently occurred parallel to the increasing diversity of gymnosperms during the Triassic and Jurassic. As in the majority of modern representatives (23), ovules of many Triassic-Jurassic gymnosperm species secreted pollination drops to capture airborne pollen grains and trigger their germination (24). Similar to angiosperm nectar, the sugary droplets offered a high-energy nutritional source (25), which could attract adult glossatan moths and other Mesozoic proboscate flying insects (26). To a large extent, early pollination drop exploitation must have been unidirectional, benefiting only moths over their plant hosts. Only with the evolution of bisexual reproductive structures, a prerequisite for effective cross-pollination by flying insects, gymnosperm taxa would open the prospect of reciprocal plant-pollinator benefits. Among Mesozoic gymnosperms, bisexuality has only been confidently identified in permineralized cones of the Bennettitales. The overall architecture of these structures provides strong evidence of self-pollination, whereas regular traces of beetle predation on pollen suggest fortuitous, incidental entomophily (27). Any mutualistic association between Triassic-Jurassic glossatan moths and Bennettitales remains unlikely.

Pollinator interaction with Gnetales seems more plausible option. Many extant members of this order produce female (ovuliferous) and bisexual (ovuliferous and polliniferous) cones on separate plants (28). The ovules on bisexual cones are usually sterile but can still secrete sugary pollination drops. Pollination is accomplished either by wind, by insects, or by a mixed wind/insect pollination mode (amphiboly). A variety of flying proboscate insects has been observed feeding on the pollination drops of both fertile and sterile ovules of Gnetum (29), Ephedra (30), and Welwitschia (31). Although their organization is still imperfectly known, (probably Middle) Permian bisexual cones described as Palaeognetaleana auspicia could well be regarded as possible gnetalean stem relatives (32). In situ pollen grains correspond to the long-ranging, essentially gnetalean form genus Ephedripites. Similar ephedroid pollen is regularly detected in Triassic and Jurassic palynological records, mostly as a subordinate element. Among insects visiting modern Gnetales, the Lepidoptera are represented by glossatan moths belonging to the large ditrysian families Pyralidae and Geometridae (33). Similar to other insect groups (46), early Lepidoptera may have been essentially immune to the end-Triassic crisis, probably because both larval and adult moths were thriving on long-ranging plant species. Potential host-plant availability can be inferred from those gymnosperm pollen types that cross the Triassic-Jurassic boundary in the Schandelah-1 core (Fig. 2). Most of this pollen corresponds to conifers (Perinopollenites, Ovalipollis, and Ricciisporites), which may represent a regional vegetation predominantly composed of conifers, peltasperms, and cycads. Concomitant proliferation of fern spores assignable to the Matoniaceae and Schizaeaceae is likely to be indicative of stressed habitats (43, 44).

Subsequently, a gradual return to gymnosperm woodland in the earliest Hettangian is evidenced by increasing amounts of conifer pollen, particularly Araucariacites, Perinopollenites, and Cerebropollenites (42, 45). Similar to other insect groups (46), early Lepidoptera may have been essentially immune to the end-Triassic crisis, probably because both larval and adult moths were thriving on long-ranging plant species. Potential host-plant availability can be inferred from those gymnosperm pollen types that cross the Triassic-Jurassic boundary in the Schandelah-1 core (Fig. 2). Most of this pollen corresponds to conifers (Araucariacites, Araucariacites; Cheirolepidiaceae, Classopollis; Cupressaceae, Perinopollenites), cycads (Cycadales, Chasmatosporites), and pteridosperms (Caytoniales, Vitrisporites). However, among these groups, the Araucariaceae and Cheirolepidiaceae were unlikely hosts for proboscate moths. Modern Araucariaceae (47) and, most probably, the extinct Cheirolepidiaceae (48) are characterized by extra-ovular pollen germination, not requiring secretion of a pollination drop. Rare occurrences of Ephedripites may indicate the persistent presence of entomophilous) Gnetales in the regional vegetation.

By analogy with quantitative records of dispersed wing scales from young lake sediments (9), it may perhaps be hypothesized that the scale-bearing intervals of the Schandelah-1 core reflect severe outbreak of defoliating lepidopteran larvae at the time of the end-Triassic ecological crisis. It is also conceivable that the crisis presented taphonomic constraints for the preservation of the scales. Dieback of woody vegetation could have promoted increased runoff and rapid burial of organic materials under low-oxygen or even euxinic conditions, ensuring long-term preservation of delicate chitinous structures. A search for further Mesozoic wing-scale records is needed to fully assess the potential of the palynological microfossils as a useful source of evolutionary and ecological information, but it seems safe to conclude that our Rhaetian-Hettangian data already offer a new window on the timing of basal lepidopteran divergences.
MATERIALS AND METHODS
The Schandelah-1 well, covering a Late Triassic (Rhaetian) to late Early Jurassic (Toarcian) shallow-marine succession, was drilled close to the village of Schandelah in the vicinity of Braunschweig, Lower Saxony (site coordinates, 52°18′23″N/10°42′66″W; elevation, 84 m; well depth, 338 mbs). The recovered sediment core is stored at the German core repository for scientific drilling in Berlin-Spandau, managed by the German Research Centre for Geosciences (GFZ), Potsdam. The Rhaetian-Hettangian interval (Fig. 2) is composed mainly of organic-rich shales and claystones, interspersed with several massive sandstone beds. Chronostratigraphic control is provided by palynology and ammonite biostratigraphy. At 335 mbs, the presence of the dinoflagellate cyst Lunnomadium scanisense confirms a Rhaetian age for the basal part of the studied section. Together with a marked quantitative palynofloral turnover, last occurrences of the typically Triassic pollen types Lunatisporites rhaeticus and Ovalipollis ovalis at 319.50 mbs approximate the Rhaetian-Hettangian boundary. This boundary is tentatively placed at 318.60 mbs, at the base of gray-brown sandstone beds corresponding to the Psilonoten Sandstone, a regional marker horizon. Within the Hettangian, the Planorbis, Liassicus, and Angulata chronozones of standard NW European ammonite zonation can be recognized.

Initial palynological investigation of the core revealed the presence of scales in two separate intervals: in the Rhaetian section between 337.50 and 336.80 mbs, and in the Hettangian section between 317.90 and 316.40 mbs. A single scale was noted at 312.00 mbs. For the purpose of the present study, five fractions, prepared at the palynological laboratory of the Goethe University, Frankfurt, were still available for detailed analysis; these correspond to depths of 337.50, 336.80, 317.90, 317.10, and 316.40 mbs. In addition, eight new fractions from samples of the Schandelah-1 core were prepared at the Laboratory of Palaeobotany and Palynology of Utrecht University; these preparations correspond to depths of 337.40, 336.70, 336.20, 317.70, 317.30, 316.80, 316.60, and 316.20 mbs. All fractions were chemically prepared following standard palynological protocols, including successive HCl and HF treatment and storage in a glycerol-water solution. Subsequently, individual scales were microscopically traced in uncovered preparations and transferred to separate slides using a nose hair–tipped needle. All isolated scales were photographed using light microscopy. Apart from some specimens selected for sectioning, the scales were then mounted on SEM stubs for imaging with a VEGA TESCAN TS130LM scanning electron microscope. Both scale isolation and imaging were performed at Bonn University.

As an aid to taxonomic identification of the fossil scales, we studied SEM images showing scale morphology and structure of representatives of recent Lepidoptera and other scale-bearing hexapod groups (see the Supplementary Materials). Scales for SEM analysis were removed from specimens representing the Colembola, Archaeognatha, Zygentoma, Thysanoptera, Psocoptera, Coleoptera, Lepidoptera, and Culicidae, made available by Museum Koenig, Bonn. Imaging was performed at Bonn University. Additional images representing lepidopteran families were provided by the State Museum of Natural History, Stuttgart.

SUPPLEMENTARY MATERIALS
Supplementary material for this article is available at http://advances.sciencemag.org/cgi/content/full/4/1/e1701568/DC1

Supplementary Text: Comparative scale morphology
fig. S1. SEM images of scales of Colembola and Archaeognatha.
fig. S2. SEM images of scales of Zygentoma and Psocoptera.

REFERENCES AND NOTES
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fig. S4. SEM images of scales of coelolepidan Lepidoptera.
fig. S5. SEM images of scales of Culicidae.
fig. S6. SEM images of scales of Zygentoma and Psocoptera.
fig. S7. SEM images of scales of Collembola and Archaeognatha.
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fig. S9. SEM images of scales of coelolepidan Lepidoptera.
fig. S10. SEM images of scales of Culicidae.
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fig. S17. SEM images of scales of coelolepidan Lepidoptera.
fig. S18. SEM images of scales of Culicidae.


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