Report

Current Biology

Beetle Pollination of Cycads in the Mesozoic

Highlights

- A specialized beetle-mediated pollination mode is reported from Burmese amber
- The mid-Cretaceous boganiid beetle has many pollenfeeding adaptations
- The fossil boganiid was probably a pollinator of Encephalarteae cycads
- This suggests a probable ancient origin of beetle pollination of cycads in the Mesozoic

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In Brief

Cai et al. report a new mid-Cretaceous boganiid beetle with specialized pollenfeeding adaptations. The phylogenetic analyses and associated pollen grains indicate that it was probably a pollinator of early cycads. This suggests an ancient origin of beetle pollination of cycads long before the rise of flowering plants.



Beetle Pollination of Cycads in the Mesozoic

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SUMMARY

Cycads, unlike modern wind-pollinated conifers and Ginkgo, are unusual in that they are an ancient group of gymnosperms pollinated by insects [1-3]. Although it is well documented that cycads were diverse and abundant during the mid-Mesozoic, little is known about their biogeography and pollination before the rise of angiosperms. Direct fossil evidence illuminating the evolutionary history of cycads is extremely rare [4, 5]. Here we report a specialized beetle-mediated pollination mode from the mid-Cretaceous of Myanmar, wherein a new boganiid beetle, Cretoparacucujus cycadophilus, with specialized pollen-feeding adaptations in its mouthparts and legs, was associated with many pollen grains of Cycadopites. Phylogenetic analyses indicate Cretoparacucujus as a sister group to the extant Australian Paracucujus, which pollinate the cycad Macrozamia riedlei. Our discovery, along with the current disjunct distribution of related beetle-herbivore (tribe Paracucujini) and cycad-host (tribe Encephalarteae) pairs in South Africa and Australia, indicate a probable ancient origin of beetle pollination of cycads at least in the Early Jurassic, long before angiosperm dominance and the radiation of flowering-plant pollinators later in the Cretaceous.

RESULTS AND DISCUSSION

Cycads (Cycadales) are a non-speciose group of ancient living seed plants with remote roots in the Permian [1]. They are remarkable gymnosperm remnants, dominating terrestrial ecosystems during the Mesozoic and dwindling to their current diversity of approximately 330 species as angiosperms rose to dominance [2]. Cycads are dioecious gymnosperms, and most, if not all, share obligate mutualisms with specialist insect pollinators such as beetles (Coleoptera) and thrips (Thysanoptera) [3]. Understanding pollination modes of Mesozoic cycads is significant for elucidating the early diversification of cycads and insect-plant associations before angiosperm dominance.

With >380,000 named living species constituting almost 25% of all known lifeforms on our planet, beetles exhibit an astonishing morphological, taxonomic, and ecological diversity [6]. The beetle family Boganiidae is a small, monophyletic, and relict cucujoid group, with six extant genera and 15 species restricted to southern Africa, southwestern and eastern Australia, and New Caledonia [7]. Adult boganiids, characterized by head frons with median endocarina and mandibles with a dorsal setose cavity near the base, are distinctive among Cucujoidea [7]. Boganiidae currently comprise two subfamilies: Boganiinae (two genera) and Paracucujinae (four genera). In Paracucujinae, two closely related but geographically widely separated genera, Metacucujus and Paracucujus (tribe Paracucujini), feed on cones of the cycad tribe Encephalarteae (Zamiaceae) in southern Africa and the southwest of Western Australia [4, 7, 8]. This distribution suggests a Gondwanan vicariance during the Middle Jurassic resulting in the isolated distribution for these poorly dispersing beetles and their cycads [4, 9]. Here, we report on a cucujoid beetle belonging to the Boganiidae preserved within a piece of mid-Cretaceous Burmese amber (approximately 99 Ma [10]), which also harbors many cycad pollen grains alongside the beetle. The phylogenetic placement of the beetle along with the presence of specialized mandibular cavities known in modern cycad-pollinating boganiids for the transport of pollen reveal the fossil to be an early-cycad-visiting species. This specialized beetle-gymnosperm association represents the first probable insect pollination mode for Cycadales during the mid-Mesozoic.

Systematic Paleontology

Order Coleoptera Linnaeus, 1758.

Family Boganiidae Sen Gupta and Crowson, 1966.

Subfamily Paracucujinae Endrödy-Younga and Crowson, 1986.

Cretoparacucujus cycadophilus gen. et sp. nov. Cai and Escalona.



ZooBank LSID (*Cretoparacucujus*): urn:lsid:zoobankorg:act: 578961B8-5BDC-4FFF-92FC-8F0F6FB65B58.

ZooBank LSID (*C. cycadophilus*): urn:lsid:zoobank.org:act: 377DF250-3F15-4A47-AAD4-EEE71FE3909D.

Material

Holotype, NIGP166883, female. Mid-Cretaceous amber (ca. 99 million years ago), Tanai, Kachin State, northern Myanmar.

Etymology

The generic name is a combination of *creto-* and the genus *Paracucujus*. The specific epithet is a combination of Greek *kykas* (meaning, cycad) and *philia* (meaning, friendly love or affection).

Diagnosis

Cretoparacucujus is distinguished from other boganiids by the following combination of characters: upper body surface sub-glabrous; head large, slightly wider than pronotum; antenna fili-form, without antennal club; clypeus sub-triangular, apex widely notched medially; frontal carina meeting frontoclypeal sulcus; mandible long, nearly straight; maxillary palpus elongate, with maxillary palpomere 4 much shorter than palpomere 3; protibial apex not expanded; and elytral punctation seriate.

Description

Refer to Data S1 for a complete description.

Figure 1. Photomicrographs of *Cretoparacucujus cycadophilus* and Mouthparts of Extant *Paracucujus rostratus*

(A) General habitus, dorsal view, under epifluor-escence.

(B) Head of *C. cycadophilus*, dorsal view, under normal reflected light.

(C) Right mandible of *P. rostratus*, showing the characteristic setose mandibular cavity, under scanning electron microscope (SEM).

(D) Mouthparts of *C. cycadophilus*, dorsal view, under normal reflected light.

(E) Mouthparts of *P. rostratus*, dorsal view, under SEM.

(F) Left maxillary palpus of *C. cycadophilus*, under epifluorescence.

(G) Right protarsus of *C. cycadophilus*, lateral view, under reflected light.

Abbreviations: ca, mandibular cavity; cl, clypeus; en, endocarina; ey, compound eye; fs, frontoclypeal sulcus; ma, mandible; mp2–4, maxillary palpomeres 2–4; pt1–5, protarsomeres 1–5; se, adhesive setae. See also Figures S1 and S2.

The fossil species *Cretoparacucujus cycadophilus* (Figures 1A, 2, S1A, and S1B) displays many diagnostic morphological features of Boganiidae, including mandibles with large, setose cavities on the dorsal surface (synapomorphy of the family; Figures 1B and 1D), elongate clypeus (Figure 1D), and tarsi with a greatly reduced tarsomere 4 and ventrally lobed tarsomere 3 (Figures 1G and S2F)

[7]. Cretoparacucujus cycadophilus can be further placed in one of the two extant subfamilies, namely Paracucujinae, by the lateral pronotal carina without an anterior callosity. long and parallel-sided apical maxillary palpomere (Figure 1F), and seriate elytral punctation (Figures 1A and S1A). To investigate the precise phylogenetic position of the new genus, we analyzed an integrated morphological data matrix combing our fossil and representatives of all extant genera using both parsimony and Bayesian approaches. Results under parsimony, using both equal weights and implied weighting, all coincided in the monophyly of Boganiidae. All analyses strongly supported Cretoparacucujus as a sister group to Paracucujus, and they together as sister to Metacucujus (Figure 3). The alternative Bayesian analysis produced a similar tree, supporting Cretoparacucujus as a sister group to Paracucujus (Figure 3). To date, Paracucujinae comprise four genera (Athertonium, Dzumacium, Metacucujus, and Paracucujus) [7]; Cretoparacucujus can be assigned to the derived Metacucujus + Paracucujus clade based on a single synapomorphy, i.e., filiform antennal flagellum without distinct antennal club (Figure S1C) [7]. A close affinity with the Australian Paracucujus is supported by the striate elytra and apex of clypeus with small teeth [7].

The most impressive feature of *Cretoparacucujus* is the large prognathous head with large compound eyes, sharp mandibles, and extremely long maxillary palpi (Figures 1A, 1B, 1F, S1D, S1F, and S2A). These features, close to those of



Paracucujus (Figures 1E, S1E, and S1G) and the Jurassic Parandrexis (Parandrexidae), probably suggest an open habitat, and most likely habitation on the surface of the strobiles of gymnosperms [11]. The mandible of Cretoparacucujus bears a dorsal, round, and setose cavity (Figures 1D and S1D) near its base, and the clypeus is basally constricted (Figure 1D), making the mandibular cavity clearly visible in dorsal view. This specialized modification of the mandible (Figure 1C) has been known to function in containing and most likely transporting pollen grains, as both adults and larvae of extant Boganiidae feed on pollen [7, 12]. In Cucujoidea, similar analogous mandibular cavities are known in the extant pollen beetles (Nitidulidae: Meligethinae), which are pollenophagous and possible pollinators [13]. Observations of the mandibles of some extant boganiids such as Athertonium and Boganium have provided direct evidence that the peculiar mandibular cavities are capable of carrying a few pollen grains [13]. As in most boganiids, the tarsi of Cretoparacucujus have dense normal adhesive setae on the ventral surfaces of the basal three tarsomeres (Figures 1G and S2F), which is a specialization for holding and climbing on the surfaces of plants. In addition, the maxilla of Cretoparacucujus, with an apically expanded and densely setose galea (Figure S2D), is typical of many modern boganiids, and it may be used to collect small particles such as pollen and direct them into the mouth.

In Paracucujinae, the life history of *Dzumacium* (New Caledonia) is unknown, and their feeding habit remains elusive. Adults of *Athertonium* from eastern Australia are pollenophagous, associated with angiosperms such as Lauraceae, Elaeocarpaceae, Cunoniaceae, and Meliaceae [7]. Species of *Metacucujus* from South Africa are dependent on male cones of various cycads in the genus *Encephalartos* [7]. There is evidence indicating that *Metacucujus encephalarti*, as well as an erotylid beetle, are probably the main pollinators of *Encephalartos* [8, 14]. The

Figure 2. Photomicrographs of Cycad Pollen Grains Associated with *C. cycadophilus* (A) General view of *C. cycadophilus* and aggregations of pollen grains by the beetle.

(A') Enlargement of an aggregation of three pollen grains.

(A'') Enlargement of a single grain.

(B) Enlargement of three larger aggregations of pollen grains.

(C) Enlargement of (B), showing 14 aggregated pollen grains.

(D) Enlargement of (B), showing six aggregated pollen grains.

monotypic genus *Paracucujus* from southwestern Australia occurs in large numbers on male cones of *Macrozamia riedlei* and sometimes in sticky traps on female cones [7, 13]. As such, *Paracucujus rostratus* is probably a pollinator of *Macrozamia riedlei* [3, 13]. Recent phylogenetic analyses indicated that the extant *Metacucujus* and *Paracucujus* are close extant sister groups [7], as are the two associated cycad host-plant genera,

which belong to the same tribe, Encephalarteae (Zamiaceae). This remarkable distribution indicates a Gondwanan vicariance that began during the Middle Jurassic for these poorly dispersing beetle and cycad pairs [4, 9]. Apparently, the beetle-cycad interaction was established during the Mesozoic as supported by the disjunct distribution of the beetle and cycad pairs. Our discovery of a *Paracucujus*-related genus from the Cretaceous suggests that *Cretoparacucujus* may have a similar feeding habit to that of *Paracucujus*, feeding on pollen grains of cycads, as this seems to be a biological trait for the clade as a whole.

It is remarkable that several aggregations of exquisitely preserved pollen grains are located along the left side of the fossil beetle and with two pollen grains close to the head (Figure 2). All pollen grains associated with C. cycadophilus are boat shaped, prolate, and monosulcate. They display an elliptical outline and rounded polar margin in polar view and subcircular shape in equatorial view. Average polar axis length and equatorial diameter of the palynospecies (N = 21) are 20.65 μ m (20.07– 21.04 µm) and 14.30 µm (14.20-14.37 µm), respectively. The length-width ratios range from 1.1 to 1.4. The sulcus is elongate, extending almost the entire length of the grain. The sulcus exhibits rounded ends and it is much broader at their ends $(\sim 4.62 \ \mu m)$ than at the mid-point $(\sim 2.12 \ \mu m)$. The ornamentation is psilate, and the pollen exine is approximately 1 μ m thick. Based on a combination of the shape, sulcus structure, and ornamentation, the pollen can be referred to Cycadopites, a form-genus of polyphyletic origin occurring in sediments from the late Palaeozoic to Holocene [1], which can be also produced by modern cycads. It is challenging to affiliate Cycadopites that are taphonomically deformed to a particular group of gymnosperms, as they are comparable to pollen grains of a wide range of plants, including Bennettitales, Cycadales, Czekanowskiales, Ginkgoales, Peltaspermales, Pentoxylales, and a few basal angiosperm lineages. However, these amber-entombed pollen



Figure 3. Phylogenetic Positions of *Cretoparacucujus* from Bayesian Inference and Maximum Parsimony Analyses

Posterior probabilities above 0.5 are shown on branches in the Bayesian inference (BI) tree (left); bootstrap support values over 50% are shown on the maximum parsimony (MP) tree (right) (strict consensus of two trees: length = 121 steps, consistency index [CI] = 0.65, retention index [RI] = 0.67). See also Figure S3, Table S1, and Data S1.

grains exhibit three-dimensional features with high fidelity that allow for a more accurate systematic attribution. These pollen grains differ from those of Ginkgoales by their oval outline, rounded ends, and the colpus form [15]. Some pollen grains of extinct Bennettitales also belong to *Cycadopites*, but they are distinguished by their large size and a more spindle-like outline [15]. The present pollen grains are comparatively small in size (~21 µm long), close to some species of *Cycas, Macrozamia*, and *Zamia*, but considerably smaller than those of *Encephalartos* and *Lepidozamia* [16].

Another fact supporting this beetle-mediated pollination is that these *Cycadopites* pollen grains are in multiple aggregations, which comprise 3–14 grains in the specimen (Figures 2A' and 2B–2D). Many modern entomophilous cycad pollens and pollens of insect-pollinated angiosperms adhere in large aggregations, whereas wind-pollinated, or anemophilous, pollens are dispersed as single grains, or monads [17, 18]. The Burmese amber has yielded a diverse Cretaceous flora including moss, liverworts, ferns, conifers, and angiosperms, but cycads remain unknown. Our discovery of the *Cycadopites* pollen represents the first evidence of cycads from Burmese amber.

Given the feeding functional morphology and phylogenetic placement of Cretoparacucujus, as well as the identification of the associated cycad pollen aggregations, the Cretaceous C. cycadophilus, like the related P. rostratus, was probably a cycad pollen feeder and most likely a pollinator for Macrozamia-related cycads that belong to the tribe Encephalarteae (Figure S3). Moreover, the establishment of this type of boganiid-cycad association may be significantly older, probably extending back prior to the eventual breakup of Gondwana during the Early Jurassic, some 167 million years ago [19]. The hypothesis is also supported by the remarkable vicariance of two poorly dispersing cycad-host and beetle-herbivore pairs from widely separated Gondwanan continents (Figure 4) [9]. The species of Encephalartos from southern Africa and Macrozamia riedlei from Western Australia, belonging to the same tribe Encephalarteae (Zamiaceae), are closely related [20], so are their corresponding pollenophagous beetles: Metacucujus and its modern sister group, Paracucujus [7].

Cycads have a rich fossil record, extending to the earliest Permian [21] and peaking in diversity during the Jurassic and Cretaceous periods [1]. Unfortunately, fossil cycads are usually preserved as fragmentary stems or leaves, and rarely with reproductive organs, which yield non-overlapping character datasets, hindering phylogenetic analyses [22]. The earliest cycad with an affinity to Zamiaceae, Plagiozamites oblongifolius, comes from Late Permian of China [23]. Its stomatal architecture, especially the prominent thickenings of the subsidiary cells, resembles that of some extant Macrozamia and Encephalartos [23], both belonging to the entomophilous Encephalarteae. More convincing fossils placed in Encephalarteae are from the Cretaceous of Antarctica, India, and Patagonia [22]. Cenozoic representatives of Encephalarteae from the early Eocene of Patagonia and the middle Eocene of Australia suggest that the tribe occurred widely across Gondwanan landmasses until the final breakup of the former supercontinent [22].

Recent fossil-calibrated molecular dating estimates produced various ages for the origin of crown cycads, ranging from ca. 200 Ma (Early Jurassic) to 274.5 Ma (Late Permian), but all suggested that the extant diversity of cycads derives from mostly Miocene radiations and postdates the Cretaceous-Paleogene boundary [2, 20, 24]. Encephalarteae, as traditionally defined based on morphology, was recovered as a monophyletic group by almost all recent molecular phylogenies [2, 20, 24], except in a maximum-likelihood analysis based on a single nuclear gene [2]. It is unexpected that these molecular divergence estimates suggested an origin of stem-group Encephalarteae during the Late or Early Cretaceous [2, 20, 24]. This young age for origin appears to be unlikely, because at that time Africa (western Gondwana) was widely separated from eastern Gondwana, which comprises Antarctica, India, Madagascar, Australia, and New Zealand [19]. Considering the low dispersal ability of early cycads and their potential beetle pollinators, Encephalarteae and their cycad-beetle associations were most likely established before the separation of the Gondwanan landmasses, a time during the Early Jurassic or earlier. Additionally, fossil evidence of late Permian cycad cuticles and a late Triassic whole-plant from China [23, 25], both with apomorphies of modern Encephalarteae, suggest a much longer evolutionary history for the tribe than implied by methods relying solely on indirect inferences from molecular clocks.

Modern cycads are principally pollinated by beetles, and rarely by thrips or moths [3]. Their beetle pollinators mainly include cucujoids (Biphyllidae, Boganiidae, Erotylidae, and Nitidulidae), weevils (Anthribidae, Belidae, Brentidae, and Curculionidae), and, unusually, tenebrionids [3]. Cycadothrips, a lineage of basal thrips, participate in the pollination of some species of Macrozamia in Australia. Based on the biogeography of Cycadothrips and the antiquity of the family to which it belongs (Aeolothripidae), thrips were most likely among the earliest pollinators of these plants [4, 26], although fossil evidence is lacking. Weevil pollinators of cycads, as revealed by extensive comparative studies, are probably derived from angiosperm-dwelling ancestors, rather than from the older gymnosperm feeding weevils [27], and are therefore correspondingly and comparatively younger in age. It is noteworthy that the Triassic obrieniids with a weevil-like rostrum (nose) and their frequent co-occurrence with cycad remains have been hypothesized as evidence of a potential Triassic beetle-cycad association [28]. Such early interactions have been



exemplified by a Middle Triassic cycad cone that contains pollenladen coprolites possibly produced by beetles [28]. This find provides the earliest evidence for a cycad-insect interaction, illuminating the early stage in the establishment of complex entomophily in cycads [28]. Among the cucujoid pollinators, Nitidulidae have their oldest known fossil records from the Early Cretaceous, whereas Biphyllidae and Erotylidae date back to the mid-Eocene [29-31]. By contrast, the first boganiid fossil, Palaeoboganium jurassicum, has been recently documented from the Middle Jurassic (approximately 165 Ma) of northeastern China [32]. Morphology-based phylogenetic analyses recovered Palaeoboganium as a sister group to Metacucujus + Paracucujus, which implied the fossil beetle as "a good candidate" for a pollinator of cycads [32]. The feeding habits of P. jurassicum are undoubtedly difficult to determine due to the insufficient preservation of compression fossils (e.g., lacking details of mandible and gut contents), and its position as basal to the two genera pollinating cycads leaves its biology somewhat ambiguous. Nevertheless, P. jurassicum demonstrates that Boganiidae were present during that period, originating at least as early as the Middle Jurassic, and were more widespread than initially believed. Biogeographically, Boganiidae may have also occurred across Gondwana during the Middle-Late Jurassic, as similar distribution patterns have been observed in multiple beetle lineages as shown by fossils from the Jurassic of China and Australia [33]. The Boganiidae-Encephalarteae association was probably established during the Early-Middle Jurassic, and the specific beetle-herbivore and cycad-host pairs were probably widespread on Gondwana, at least in what would become southern Africa, southwestern Australia, and Antarctica, the last then connecting the former two continents [34]. Like boganiids, the present global distribution of pollinating erotylids in Africa, Asia, Australia, and North and Central America [3] suggests an early development of an erotylid-cycad relationship before late-Mesozoic continental drift [35]. Collectively, the preliminary cycad-insect associations are probably ancient, extending to the Triassic or earlier [28], whereas insect pollination of cycads may be as early as the Early Jurassic [4, 9, 12], long before the origin and diversification of major groups of angiosperms and the concomitant diversification of angiosperm pollinators such as moths, flies, and bees later in the Cretaceous [5, 6, 36, 37]. Moreover, such an ancient association was probably widespread at least

Figure 4. Geographic Distribution of the Known Entomophilous Cycads of the Tribe Encephalarteae and Their Boganiid Pollinators

The phylogenetic relationships and divergence time of the widely separated lineages within Boganiinae are shown. The arrow indicates the divergence time estimated by separation of Gondwana in the Jurassic. Abbreviations: AU, Australia; MM, Myanmar; ZA, South Africa. See also Figure S3 and Table S1.

across Gondwana in the Mesozoic, as evidenced by Cretaceous cycads of the Encephalarteae from Antarctica, India, and South America [22].

Our discovery of a mid-Mesozoic insect pollinator of cycads provides a window into ancient insect-gymnosperm interactions alongside the diversification of angiosperms. Unlike some beetles that shifted plant-hosts from gymnosperms onto angiosperms during the Aptian-Albian gap [5], the beetle pollination of Zamiaceae cycads has most likely persisted from the early Mesozoic to the present, although confined to particular areas in the Southern Hemisphere. Their austral disjunct occurrence resulting from vicariance during the Jurassic followed by considerable stasis in pollination biology over the intervening epochs all in the face of dramatic global climate change, shifting landmasses, and considerable declining diversity among cycads and other major gymnosperm lineages.

STAR * METHODS

Detailed methods are provided in the online version of this paper and include the following:

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SUPPLEMENTAL INFORMATION

Supplemental Information includes three figures, one table, and one data file and can be found with this article online at https://doi.org/10.1016/j.cub. 2018.06.036.

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AUTHOR CONTRIBUTIONS

C.C. conceived and designed the study. C.C. drafted the manuscript, to which M.S.E., Z.Y., and H.E.E. contributed. C.C., Z.Y., H.E.E., D.H., and M.S.E participated in the morphological studies of the boganiid fossil. L.L. and C.C. participated in the studies of the cycad pollen grains. C.C. conducted the phylogenetic analyses.

DECLARATION OF INTERESTS

The authors declare no competing interests.

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STAR*METHODS

KEY RESOURCES TABLE

REAGENT or RESOURCE	SOURCE	IDENTIFIER	
Software and Algorithms			
Helicon Focus 3.10	Helicon Soft	https://www.heliconsoft.com/	
MrBayes 3.2.3	[38]	http://mrbayes.sourceforge.net	
Tracer version 1.6.0	Tracer Website	http://tree.bio.ed.ac.uk/software/tracer/	
TNT 1.5	[39]	https://cladistics.org/tnt/	

CONTACT FOR REAGENT AND RESOURCE SHARING

Further information and requests for resources and reagents should be directed to and will be fulfilled by the Lead Contact, Chenyang Cai (cycai@nigpas.ac.cn).

EXPERIMENTAL MODEL AND SUBJECT DETAILS

The Burmese amber specimens studied here are derived from the Hukawng Valley in Tanai Township, Myitkyina District of Kachin State, Myanmar. The holotype (NIGP166883) is deposited in the Nanjing Institute of Geology and Palaeontology, Chinese Academy of Sciences, Nanjing, China. The new taxonomic actions established herein are registered in ZooBank LSID urn:lsid:zoobank.org:pub: A05BB478-ECFE-4797-8FC2-86536C84A8E3. The amber piece was polished with sand papers of different grits and diatomite mud.

METHOD DETAILS

Specimen imaging

Photomicrographs were taken using a Zeiss Discovery V20 microscope system, and those with green background (Figures 1A and 1F) used fluorescence as a light source attached to a Zeiss Axio Imager 2 compound microscope. Focus stacking software (Helicon Focus 3.10) was used to increase the depth of field.

Phylogenetic analysis

We scored *Cretoparacucujus cycadophilus* sp.nov. for 63 adult characters and integrated the fossil taxon along with representatives of extant and extinct genera of Boganiidae, their related cucujoid genera, and a cleroid outgroup. Most characters (60 out of 63) of extant genera of Boganiidae, their related cucujoid genera, and an outgroup were derived from Escalona et al. [7] We added three new characters to the matrix that are relevant to the diagnosis of *Cretoparacucujus* (first three characters in the character list in Supplementary Information). We excluded the larval characters used by Escalona et al. [7], as larvae of the fossil taxa remain unknown. The character states and matrix are presented in Table S1. All characters were non-additive, unordered, and equally weighted. Bayesian analysis was carried out using MrBayes 3.2.3 [38] and the discrete Mkv+G model [40]. Two MCMC runs of four chains were run for three million generations. The consensus tree was estimated after a burn-in of 25% of the sampled trees. Convergence was confirmed with Tracer version 1.6.0. Parsimony analysis was conducted using TNT 1.5 [39] using New Technology search. We used two search strategies, including equal weights and implied weights. For the implied weights analyses, we tested a set of concavity (*k*) values from 0.25 to 10 and found no changes to the tree topology as it relates to Boganiinae. Branch support values were estimated using 10,000 bootstrap replicates.

DATA AND SOFTWARE AVAILABILITY

The detailed systematic paleontology for *Cretoparacucujus cycadophilus*, taxon selection and morphological characters used for phylogenetic analyses are available in Data S1.

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Supplemental Information

Beetle Pollination of Cycads in the Mesozoic

Chenyang Cai, Hermes E. Escalona, Liqin Li, Ziwei Yin, Diying Huang, and Michael S. Engel



Figure S1. Photomicrographs of *Cretoparacucujus cycadophilus* (NIGP166883) and Mouthparts of Extant *Paracucujus rostratus*. Related to Figure 1

- (A) General habitus, dorsal view, against black background.
- (B) General habitus, ventral view, against white background.
- (C) Left antenna, under reflected light.
- (D) Detail of head, dorsal view of C. cycadophilus, under epifluorescence.
- (E) Detail of head, dorsal view of *P. rostratus*, under scanning electron microscope (SEM).
- (F) Detail of head, ventral view of C. cycadophilus, under reflected light.
- (G) Detail of head, ventral view of *P. rostratus*, under SEM.
- Abbreviations: a1, antennomere 1; cl, clypeus el, elytron; ey, compound eye; fs, frontoclypeal sulcus;
- hw, hind wing; lp2,3, labial palpomeres 2,3; mp, maxillary palpus; mp3,4, maxillary palpomeres 3,4;
- pro, pronotum. Scale bars: 500 µm in (A) and (B); 200 µm in (C)–(G).



Figure S2. Details of *C. cycadophilus* (under normal transmitted light) and *P. rostratus* (Under SEM). Related to Figure 1

- (A) Left maxillary palpus of *C. cycadophilus*.
- (B) Left maxillary palpus of *P. rostratus* for comparison.
- (C) Left hind wing of *C. cycadophilus*.
- (D) Maxillae of C. cycadophilus, ventral view.
- (E) Adhesive setae on the first mesotaromere of *P. rostratus*.
- (F) Right mesotarsus of C. cycadophilus, lateral view.
- (G) Mesotarsus of *P. rostratus*, lateral view.
- (H) Genitalia of C. cycadophilus, female.
- (I) Female genitalia of *P. rostratus*.

Abbreviations: el, elytron; gs, gonostylus; hw, hind wing; mp2-4, maxillary palpomeres 2-4; mst1-

5, mesotarsomeres 1–5. Scale bars: 100 μ m in (A), (D) and (F); 200 μ m in (C); 50 μ m in (H).



Figure S3. Ecological Reconstruction of the Mid-Cretaceous *Cretoparacucujus cycadophilus*. Related to Figure 4

Characters	1	2	3	4	5	6
Taxa	0	0	0	0	0	0
Eronyxa	11211222132121	1233221211	1122121212121	11212111211	13211111132	211312231
Byturus	11211211132131	1112221311	1122132123	12111111321	121212123	121312122
Ericmodes	11211233132121	1113221111	2122141213	11111112321	1221212113	121212112
Protosphindus	11111233132113	82113221211	1123141213	11211213221	1321221112	121112112
Pharaxonotha	11111211132121	1113222111	23222222122	22111222421	12112121122	211112211
Hobartius	11211123132121	2112222111	2121121312	11212212321	1221121113	111211111
Metacucujus	11212212221133	82231111221	1122121112	12112212321	12211131212	212312112
Paracucujus	21112212221133	82231111221	1122121212121	12112212321	12211131212	212312112
Athertonium	11111212231233	32231111232	1122121212121	11212212321	12211133222	222312112
Dzumacium	11223212231133	82231111231	1122121212	11212212321	122?11312?2	222312122
Boganium	11223212231133	32231221233	1121121212121	11212212321	12211132222	222312122
Afroboganium	11213212231113	32231221233	1121121212121	11212212321	122?1122222	212312122
Cretoparacucujus	22122212211133	32?31111?21	112212?212	12112212321	1?211131??'	???????1?2

Table S1. Character matrix for phylogenetic analyses, including 63 adult morphologicalcharacters. Related to Figure 3