Crab in amber reveals an early colonization of nonmarine environments during the Cretaceous

Javier Luque1,2,3,4,†‡, Lida Xing5,6,†, Derek E. G. Briggs2, Elizabeth G. Clark2,†, Alex Duque8, Junbo Hui9, Huijuan Mai10,11, Ryan C. McKellar12,13

Amber fossils provide snapshots of the anatomy, biology, and ecology of extinct organisms that are otherwise inaccessible. The best-known fossils in amber are terrestrial arthropods—principally insects—whereas aquatic organisms are rarely represented. Here, we present the first record of true crabs (Brachyura) in amber—from the Cretaceous of Myanmar (~110 to 99 million years (Ma)). The new fossil preserves large compound eyes, delicate mouthparts, and even gills. This modern-looking crab is nested within crown Eubrachyura, or “higher” true crabs, which includes the majority of brachyuran species living today. The fossil appears to have been trapped in a brackish or freshwater setting near a coastal to fluvio-estuarine environment, bridging the gap between the predicted molecular divergence of nonmarine crabs (~130 Ma) and their younger fossil record (latest Cretaceous and Paleogene, ~75 to 50 Ma) while providing a reliable calibration point for molecular divergence time estimates for higher crown eubrachyurans.

INTRODUCTION

Transitions from marine to nonmarine habitats are infrequent in most metazoan groups, largely due to different physical and physiological requirements in saltwater and freshwater competition with previously established residents, and exposure to new predators (1). True crabs, or Eubrachyura, are among the animal groups that have conquered land, brackish water, and freshwater multiple times independently, although much more recently than other arthropod groups [e.g., (2, 3)].

Eubrachyurans are the most diverse group of crabs today in terms of anatomy, ecology, and species richness. They fall into two main categories based on the position of their sexual openings: the heterotrems (including primarily and secondarily freshwater crabs), with openings on the legs in males and thorax in females, and thoracotremes (including semiterrestrial and terrestrial crabs), with openings on the thorax in both sexes (4–6). The consensus is that heterotrems and thoracotremes together form a monophyletic clade Eubrachyura (7–10). However, two questions, (i) the phylogenetic position of the different freshwater crabs, and (ii) whether thoracotremes are a sister group to heterotrems or derived from within them, are still debated (7).

Molecular phylogenies suggest that the first freshwater and terrestrial crabs diverged from their closest marine relatives during the Early Cretaceous or before ~125 million years (Ma) ago (8). Direct evidence of their colonization of nonmarine environments is sparse, however, because the few fossils known from such environments largely correspond to isolated carapaces or claws of latest Cretaceous to Quaternary age (~73 Ma to Holocene) (11, 12). This gap of ~50 Ma between the fossil record of freshwater and terrestrial crabs and their predicted molecular divergence time obscures our understanding of the transition to nonmarine habitats and the number of times such transitions have occurred across families.

Here, we describe a previously unknown eubrachyuran crab, Cretapsara athanata, preserved in Cretaceous amber (~99 Ma, Cenomanian) from Myanmar, Southeast Asia (Figs. 1 and 2). This is the oldest occurrence of a true crab in amber and one of the oldest crown group eubrachyurans known. Micro–computed tomography (CT) digital reconstructions reveal that antennae, large compound eyes, mouthparts with multiple fine hairs, and even gills are preserved (Figs. 3 and 4). Our phylogenetic analysis supports the establishment of a new family with a unique mixture of primitive and advanced characters. The exceptional preservation of C. athanata in Cretaceous amber shows that highly carcinized brachyurans (13) were already established in nonmarine environments by the earliest Late Cretaceous.

RESULTS

Systematic paleontology

Euarthropoda Lankester, 1904
Decapoda Latreille, 1802
Brachyura Latreille, 1802
Eubrachyura Saint Laurent, 1980
Cretapsaridae Luque, fam. nov.

LSID. Family group: http://zoobank.org/urn:lsid:zoobank.org:act:81705335-CAED-400E-B1EC-161B56B298AE.

Included genus: Cretapsara Luque, gen. nov., by monotypy.
Diagnosis: As for type genus and species.

Remarks: Cretapsara athanata n. gen. et sp. differs from any known crab family in its combination of plesiomorphic and apomorphic...
characters. Superficially, it shares some resemblance to some marine eu-
brachyuran families such as Eogeryonidae, from the Upper Cretaceous
of Spain [Eogeryon elegius, Cenomanian (14)] (Fig. 5H) and the
Lower Cretaceous of Brazil [Romualdocarcinus salesi, Albian (15)], and
Marocarcinidae, from the Upper Cretaceous of Morocco [Marocarcinus
pasinii, Cenomanian (16)] (Fig. 5E). Cretapsara differs from Eo-
geryonidae and Marocarcinidae in the presence of a bilobate rostrum
and lack of orbital fissures, compared to the bifid and acute rostrum
and two well-developed orbital fissures diagnostic of these two families
[Marocarcinidae was initially interpreted as having no orbital fissures
due to the erroneous interpretation of the eyestalks as part of the
orbit (16) but is here recognized as having two orbital fissures and
scored as such in our analysis].

The overall carapace outline, the broad front, long legs, and the
lack of orbital fissures that characterize C. athanata (Figs. 1 and 2)
resemble those of some modern Grapsoida—a group of highly ter-
restrial thoracotreme crabs common in subtidal and supratidal settings
worldwide. Despite this, C. athanata differs from grapsoids in im-
portant respects. In most grapsoids, there is a conspicuous V-shaped
notch between the merus and ischium of the third maxillipeds (mxp3),
so that the mandibles are visible through a large, rhomboidal gap when
the mxp3 are closed, a feature absent in Cretapsara (Figs. 2, C and F,
and 3, B, E, and G). Moreover, the podomeres of the walking legs in
grapsoids, as in most thoracotreme crabs, are subtriangular to flattened
in cross section, whereas those in Cretapsara are circular in cross
section and more similar to those of other heterotremes (Fig. 3, movie S1,
and data file S1). Such similarities between Cretapsara and some
grapsoids are likely convergent adaptations to similar ecologies.

Cretapsara athanata Luque, gen. et sp. nov.

LSID. Species group: http://zoobank.org/urn:lsid:zoobank.org:act:
B6E29B76-C37E-47D8-9FDE-228DEB3DB9BA.

Etymology: The genus name combines Creta (Latin), for chalk as
in Cretaceous, and Apsara, a spirit of the clouds and waters; gender
is feminine. The specific epithet is based on athanatos (Greek),
immortal, referring to its lifelike preservation. The family name is
based on the only known genus, erected herein.

Holotype: Specimen LYAM-9 (Figs. 1 and 2) deposited in the
Longyin Amber Museum, Xishan District, Kunming 650228, Yunnan
Province, China. Specimen collected by local miners in 2015.

Type locality and horizon: Angbamo site, Tanai Township,
Myitkyina District, Kachin Province, Myanmar [~98.8 ± 0.6 Ma;
lowermost Cenomanian (17–19)].

Diagnosis: The carapace is subhexagonal, nearly as long as it is
wide; the orbitofrontal margin is nearly as wide as the body and bears
wide, shallow orbits that lack supraorbital fissures or intraorbital
spines. The rostrum is bilobate, short, and wide, measuring about
half the width of the orbitofrontal margin. The anterolateral margin
are circular in cross section, whereas those in Cretapsara are circular in cross
section and more similar to those of other heterotremes (Fig. 3, movie S1,
and data file S1). Such similarities between Cretapsara and some
grapsoids are likely convergent adaptations to similar ecologies.

Cretapsara athanata Luque, gen. et sp. nov.

LSID. Genus group: http://zoobank.org/urn:lsid:zoobank.org:act:
E6F68190-D2A7-4D6E-8CE8-EDD319E8DA13.
reduce in size posteriorly. The sterno-abdominal cavity is shallow, and there is no evidence of a linea media traversing the sternites. The pleon has a small subtriangular telson and, based on the stereo-microscope images and the micro-CT scanning, apparently has six free and unfused pleonites, from which at least the first two are fully visible dorsally; there is no evidence of uropods or uropod remains. The eyes are as large as the orbits; the corneal eye is globular, as wide as it is long, and apparently covered by small hexagonal facets; the eyestalk is short and cylindrical. The antennulae and antennae are reduced.

The claws or chelipeds are equal in shape and size, and the four pairs of walking legs are well developed, slender, and similar in size and shape; they lack chelate, subchelate, or flattened distal podomeres (e.g., propodi and dactyli); the podomeres are semicircular in cross section. Excurrent openings are well developed, small, and circular.

Description: Dorsal carapace: The carapace is subhexagonal in outline and about as long as wide (~2 mm) (Figs. 1D and 2B). The orbitofrontal margin is nearly as wide as the maximum width of the carapace, and each orbit is wide, concave, and shallow, with a sinuous supraorbital margin that lacks supraorbital fissures or spines (Figs. 2, A to C, and 3), apart from a small, anterolaterally diverging outer orbital spine (Fig. 3, F and G, thick arrow). The rostrum is short and half the width of the orbitofrontal margin, downturned, apparently with a shallow axial sulcus and raised subparallel lateral margins grading into the inner orbital margins (Figs. 2B and 3, A and F, and movie S1). The anterolateral and posterolateral carapace margins are distinct and separated by a shallow lateral expression of the cervical groove (Fig. 3, F and G, dotted line). The anterolateral margin is slightly convex, about one-third the length...
of the posterolateral margin, and it bears two anterolateral spines that are short and subtriangular (Figs. 2 and 3F). The edge of the posterolateral margin is straight to slightly convex, and it bears at least four small tubercles that are somewhat equidistant from each other (Fig. 3F, movie S1, and data file S1). The posterior margin of the carapace is about two-thirds of the maximum carapace width, lacks reentrants, and, as evidenced by the micro-CT scans, appears to be rimmed and bearing a row of small tubercles (movie S1 and data file S1).

The cervical groove becomes faint axially, deepens laterally reaching the anterolateral margin, and is expressed ventrally (Fig. 3, C, F, and G); the groove is concave posterior to the mesobranchial region and sinusoid toward the lateral margin. The branchiocardiac groove is well developed, somewhat deep, flanking the cardiac region and, apparently, the urogastric region mesially and the mesometabranchial region distally (Fig. 3F).

The epigastric region is wide, lacking spines or tubercles, laterally inflated, and axially sulcate; the lateral lobes reach the frontal region and are nearly continuous posteriorly with the protogastric lobes (Fig. 3, A and F). The protogastric region is wide, lacking spines or tubercles, laterally tumid, and forming an almost ovate longitudinal lobe flanked by the epigastric region; axially, the protogastric depression separates the protogastric lobes and opens to the narrow anterior mesogastric region. The mesogastric region is relatively small, lacking spines or tubercles, defined anteriorly by two grooves that separate the protogastric lobes and posteriorly by a concave
cervical groove. The metagastric region is well developed, inflated, and about three times as wide as long; it is bounded anteriorly by the cervical groove, laterally by grooves that separate it from the epibranchial regions, and posteriorly by a narrow and depressed urogastric region. The urogastric region is considerably shorter and narrower than the metagastric region, about twice as wide as it is long, depressed relative to the metagastric and cardiac regions, and separated from them by faint grooves, delimited laterally by the branchiocardiac groove. The cardiac region is almost subtrapezoidal, lacking spines or tubercles, is wider posteriorly, and is delimited laterally by the branchiocardiac groove. The intestinal region appears to be wide, shorter mesially, and slightly wider laterally, with two lateral swellings parallel to the posterior margin (Fig. 3, A and F, and movie S1).

Ventral carapace: The buccal cavern is wide and covered by a pair of operculiform mxp3. The coxae of the mxp3 separate the sternum from the pterygostome. The pterygostome is curved, bulged, finely granulated, with a row of coarser tubercles parallel to the linea brachyura or molting plane (Figs. 2, C and F, and 3, and movie S1).

The thoracic sternum has fused sternites 1 to 3 that are triangular, about as long as wide, and are visible ventrally; sternite 4 appears to be subtrapezoidal in outline; sternite 5 is the widest and measures approximately two-thirds of the maximum carapace width; sternite 6 is narrower than sternite 5 but nearly as wide as sternite 4; sternites
7 and 8 are small and barely evident in the micro-CT scan. The sterno-abdominal cavity is shallow, and there is no evidence of a linea media traversing the sternites. The pleon, based on the micro-CT scan reconstruction, appears to comprise six free, unfused pleonites and a small telson. The pleonites are subrectangular in outline. The first pleonite is smallest to comprise six free, unfused pleonites and a small telson. The linea media traversing the sternites.

The eyes and eyestalk are large, with each eye approximately one-third the maximum carapace width. The corneal eye is globular, as wide as long, measuring about one-fifth the width of uropods.

Eyes and antennae: The eyes and eyestalk are large, with each eye approximately one-third the maximum carapace width. The corneal eye is globular, as wide as long, measuring about one-fifth the width of the carapace, and seems to be covered with small hexagonal facets;
the antennula and antenna are very small (Figs. 2, A to C, and 3, B, E, and G). The coxae of the mxp3 are small and do not meet axially; the ischiomerus has straight occlusal margins lined with fine setae (Fig. 2, A, C, and F) and does not form a rhomboid gap with the other mxp3 that reveals the other mouthparts, neither does it bear a crista dentata (Figs. 2, C and F, and 3, B, E, and G). The palp, which is formed by the propodus, carpus, and merus, is long, thin, and located in an inner-mesial position.

The first of the five pairs of pereopods (P1) constitutes the claws (or chelipeds), which are symmetrical or homochelous (Figs. 1 and 3). Their mobile fingers (or dactyli) are long and slender, barely curving downward, and parallel to the fixed finger or pollex of the propodus, which is also long and slender and apparently bears six to eight very small teeth on the occlusal margin (Fig. 2D, movie S1, and data file S1). The other four pairs of thoracic appendages are slender walking legs (P2 to P5), similar in size and shape, with slender, acute dactyl and lacking chelate or subchelate terminations (Figs. 1 and 3). The podomeres are semicircular in transverse section rather than flattened (Figs. 1 to 4, movie S1, and data file S1). The claws and the legs have several small, fine setae scattered over their surface (Fig. 2D and E).

Gills and excurrent openings: The micro-CT scan of the holotype of *C. athanata* revealed the presence of at least six pairs of phyllobranchiate gills, bearing apparently up to 20 series of well-developed, flattened branchial lamellae that are perpendicular to the main gill shaft (Fig. 4, D to F, and movies S2 to S4). A pair of conspicuous, small, and subcircular excurrent openings is evident beneath the antennal sockets adjacent to the epistomial region (Figs. 2, A, C, and F, and 3, B and G; movie S1; and data file S1).

**DISCUSSION**

**Taphonomy**

*C. athanata* is the first crab discovered in Mesozoic amber. The holotype is complete and articulated and preserves delicate features such as antennae, compound eyes, and mouthparts covered in minute hairs or setae (Fig. 2), which are rarely preserved in fossil crabs. The overall posture of the legs and claws (Figs. 1 to 4), the fully articulated posterior carapace and pleon (Fig. 3, A, C, D, and F), the delicate mouthparts barely open (Figs. 1 to 3), and the intact molting line (linea brachyura) and pterygostome in place (Figs. 2C and 3, B, C, E, and G; and movie S1) indicate that the holotype was entombed in resin while still alive. Although there are no struggle marks evident within the surrounding amber (e.g., drag bubbles), the pereopod 5 on the left side of the body has detached at the base of the coxa (Fig. 1, white arrows), possibly indicating limb autotomy as the crab was engulfed before the resin polymerized.

Milky amber is present along the ventral surface of the carapace adjacent to the sternites and, to a lesser extent, around the maxillipeds (Fig. 2, F and G). It is unclear whether this might be the result of decay products or moisture from the gill cavity interacting with the surrounding resin to form microscopic bubbles within the amber (20). The presence of small euhedral pyrite crystals inside the limbs and carapace (Fig. 4A, white masses) indicates the availability of iron and the activity of sulfate-reducing bacteria within the decaying carcass or surrounding sediment. Sulfate is present in elevated concentrations within seawater, and pyrite may indicate proximity to coastal or brackish settings (21). However, sulfate may also be sourced by weathering or exposure to marine water brought inland due to tropical storms or flooding events. The growth of these crystals in voids within the exoskeleton and also in vacuoles within the surrounding amber suggests that pyrite formation occurred late in the taphonomic process and may provide more information about diageneric conditions and porewater chemistry than about the biostratinomy or habitat of the crab (22).

Although there are no other arthropod syninclusions associated with the crab to guide interpretation of the preservational setting and nearby habitats, the amber appears to be a single flow of resin that contains numerous particulate inclusions and bubbles. Among the particulates are numerous carbonized wood fibers (Fig. 2): Some of these clumps of dark brown or black plant tissue show cell structure, whereas others appear to represent masses of insect frass. Such features are associated with litter amber that is formed on or near the forest floor (23). Similar organic particulates have been found alongside inclusions from marine habitats in Burmese amber, such as an ammonite shell and marine snails (24) and a marine ostracod (25). They have also been found alongside microfossils that were blown into Cretaceous resin from coastal habitats (26, 27) and shrimps that were trapped in a predominantly freshwater setting (28). The amber containing the crab differs from previously recovered aquatic samples, however, in that it does not contain beach sand [e.g., (24)], it lacks distinct layers within the amber that appear to alternate between terrestrial and aquatic resin flow conditions [e.g., (28)], and it contains a carcass instead of an empty shell or molt (e.g., 27). Thus, it is likely that *C. athanata* was entombed in a freshwater or brackish setting near the amber-producing forest—not in a fully marine setting where resin is exposed to aquatic organisms through variations in water levels or winds. On the basis of the current sample size and information available, the possibility that *C. athanata* was fully terrestrial cannot be ruled out, although an amphibious lifestyle is more likely given that several extant species of primarily freshwater crabs are amphibious.

**Ontogeny and life stage**

The holotype of *C. athanata* (carapace 2 mm wide, leg span 5 mm) is considerably smaller than most other known crab species—fossil or extant, marine or nonmarine (Figs. 1 to 4). Body sizes of just a few millimeters long are typical of early ontogenetic stages in crabs [i.e., megalopae larvae and early postlarval juveniles (29)] but tend to become much larger in subsequent molts. However, some species with miniature adults are known among extant spider crabs and pea crabs (mostly marine) (30,31), while extinct, pedomorphic (7), and “dwarfed” (32–34) species are known from the early Late Cretaceous and the Cretaceous-Paleogene interval, respectively. A number of diagnostic characters indicate that the specimens of *Cretapsara* described here are postlarval stages: (i) the pleon is folded beneath the cephalothorax; (ii) the sterno-abdominal cavity is shallow; (iii) setal tufts at the tip of the dactyl of the last leg pair (P5), which are typical of crab megalopae, are absent; and (iv) there is no evidence of uropods. However, it is not clear whether the holotype of *Cretapsara* represents a minute adult or the juvenile of a species for which adults have yet to be discovered.
Gill anatomy
Micro-CT scan images of the holotype of *C. athanata* revealed the preservation of complete gill lamellae (Fig. 3 and movies S2 to S4). To date, fossilized gills in crabs have only been reported in *Romaleon franciscae*, a marine cancroid from the Miocene of the Kerguelen Islands (35). Those specimens, preserved in concretions a few centimeters long, have phosphatized gill tissues interpreted as the result of early diagenesis acting on the crab carcasses (35). Although the delicate cuticle of the gills can be present in crab exuviae before degradation, the completeness, preservation, and posture of *C. athanata* indicate that the specimen is a carcass and not a molt (see the “Taphonomy” section).

Gills play a crucial role in a number of physiological processes including osmoregulation, ion mobilization, excretion, and respiration [(36) and references therein]. As such, the shape, size, and number of gills may vary across crabs with different lifestyles. Unfortunately, the nature of the gills among most groups of living crabs is incompletely known (4), and comparisons across a wide range of taxa and habitats are unavailable. Brachyuran crabs have phylloblanchiate gills that are characterized by a flattened gill shaft with one dorsally flowing afferent branchial sinus and one ventrally flowing efferent sinus; these two vessels are connected anteriorly and posteriorly by the gill lamellae (4). The six pairs of gills clearly recognizable in the holotype of *C. athanata* are phylloblanchiate, bearing apparently up to 20 well-developed, flattened branchial lamellae perpendicular to the main gill shaft (Fig. 4, D to F, and movie S4).

In fully marine and freshwater crabs, the gills tend to be well developed and occupy most of the branchial chamber, whereas the gills of terrestrial and semiterrestrial crabs may be reduced in size and number, leaving room for the secondary development of a vascularized “lung”-like spongy tissue to facilitate respiration (4). As a consequence, terrestrial and semiterrestrial crabs may have wider, sometimes inflated, branchial chambers to accommodate this spongy tissue (37). The gills of *Cretapsara*, however, are not reduced, and the branchial regions are too narrow to house complementary lung-like structures (Fig. 3, movies S2 to S4, and data file S1). Furthermore, the efferent openings in *Cretapsara* (Fig. 2, A, C, and F) are unlike those of modern land crabs. They are small subcircular openings beneath the antennal sockets adjacent to the epistomial region, similar to those in freshwater crabs. Thus, there is no evidence that *Cretapsara* was fully land-dwelling, but most likely water-dwelling or facultatively amphibious, as are several extant freshwater crabs (Fig. 5).

Amber fossil assemblage
Land-dwelling insects are the most common syninclusions in Cretaceous Burmese amber, but rarer terrestrial arthropods such as oniscidean isopods *Myanmararces deboiseae* (38) and *Palaearmariosedillo microsoma* (39), acari, spiders, and millipedes [e.g., (40, 41) and references therein] also occur, many of them associated with humid forest floor habitats and “litter amber” (23). Aquatic invertebrates have also been reported in Burmese amber, but predictably, these are much rarer. They include marine myopod and brackish-to-freshwater podocopid ostracods [e.g., (25, 42)], marine and land snails (43, 44), and even fragmented and transported empty ammonite shells (24). Fragments of crinoid column ossicles, corals, and oysters have been found attached to the outer surface of pieces of amber (45, 46), and borings made by pholadid bivalves are common within this amber deposit (21). Adding to the list of aquatic fauna trapped in Burmese amber, freshwater penaeoid shrimp associated with freshwater penny larvae (Psephenidae) have been reported from the area (28). These occurrences suggest that the resin-producing forest where *C. athanata* lived, presumably rich in Araucariaceae (47) or Cupressaceae (48), was close to fluvio-estuarine (28) and coastal (24) settings.

Paleoecology
All previous crabs reported from amber are representatives of the family Sesarmidae from the early Miocene of Chiapas, Mexico (~15 Ma) (49–52). Extant sesarmids are land- and tree-dwelling thoracoctreme crabs from tropical to subtropical coastal systems worldwide, some of which live in phytotelm or small water bodies in rainforests (Fig. 5R) (53, 54). Such a lifestyle facilitates the inclusion of sesarmid crabs in tree resins (49–51).

The available evidence suggests that *C. athanata* was neither marine nor fully terrestrial but a fully brackish to freshwater crab, even facultatively amphibious (movie S5), that lived either on the forest floor or in shallow bodies of water on the forest floor. Most living eubrachyuran families are fully marine (Fig. 5, white squares), but just over a dozen have representatives that have invaded freshwater or moved onto land—several of them independently (55).

Extant members of primarily or true freshwater crabs are restricted to the superfamilies Trichodactylidea, Pseudothelphusioidea, Potamoidea, Potamonautoidea, and Gecarcinucoidea (Fig. 5, light gray squares). They are exclusively freshwater, and their larvae undergo direct development. Some of these species are facultatively terrestrial or even arboreal, reaching small water bodies formed in rainforest trees and epiphytes (54, 56), as do some sesarmids crabs. Secondarily freshwater crabs, such as some varunids, live in freshwater as adults but, unlike primarily freshwater crabs, their free-living larval stages rely on the ocean (55). Land-dwelling eubrachyurans, either intertidal or fully terrestrial, are largely restricted to thoracocretine families within Grapsoidea and Ocydoodoeida (Fig. 5, Q to T, dark gray and black squares). Obligate air-breathing land crabs such as the Christmas Island red crab, *Gecarcoidea natalis*, undergo massive migrations to the coast to release their planktonic zoeae larvae in the ocean. After 3 to 4 weeks of development, they return to the shoreline by the thousands as new megalopae recruits, which metamorphose into early postlarval juveniles before migrating inland where the adults dwell (57). Similar patterns of megalopae and juvenile postlarvae recruitment occur in other true land crabs, such as the gecarcinids *Gecarcinus* and *Johngarthia* [e.g., (58)], and in semiterrestrial crabs, such as the sesarmid *Chiromantes* spp. [e.g., (59)].

Although the available evidence suggests that *C. athanata* was neither fully terrestrial nor semiterrestrial, unlike the groups mentioned above, it could provide an analog to modern primarily freshwater crabs or even have been a catadromous freshwater crab with marine larval development, whose early postlarvae were trapped in fluvio-estuarine to coastal settings.

Phylogenetic and evolutionary implications
The origins of Eubrachyura
The discovery of *Cretapsara* in amber of early Cenomanian age (~99 Ma), together with the occurrence of several other Cenomanian heterotreme eubrachyurans such as *Eogeryon* from Spain (14), *Lithophylax* from France (60), and *Marocarcinus* from Morocco (Fig. 5) (16), shows that crown eubrachyurans were already diverse worldwide and occupied novel habitats, including freshwater, by
this time. Molecular phylogenies indicate that heterotremes and thoracotremes diverged in the latest Jurassic—earliest Cretaceous (175 to 100 Ma) (8, 9). However, the oldest unequivocal heterotremes, which are all marine, are from the latest Early Cretaceous (Aptian–Albian, ~115 to 110 Ma) [e.g., (7, 15, 61–63)]. In contrast, the oldest unequivocal thoracotremes, which are also marine, are from the Eocene (~50 Ma) (Fig. 5) (52, 64, 65). Unfortunately, most of these early occurrences are so incomplete that their systematic affinities remain unclear, and they are therefore unreliable for node calibrations and inferences about the origins of modern eubrachyurans.

Despite the superficial resemblance of C. athanata to some thoracotremes (e.g., sesarmids), the form of its cylindrical limb podomeres, the narrow sternum, the visibility of the first pleonites dorsally, and the marked grooves in the dorsal carapace are similar to those in several crown eubrachyuran body plans. Our phylogenetic results support the heterotreme affinities of Cretapsara, which is recovered well nested within the heterotremes and closer to the clade formed by most crown eubrachyurans (Fig. 5).

Recent phylogenomic studies indicate that Potamidae, a freshwater family of heterotreme crabs, are sister to Thoracotremata—the group that includes the true land crabs and allies—rather than being closer to other freshwater heterotremes (see also Fig. 5) (10, 66, 67). Our results support the paraphyly of “Heterotremata,” with thoracotremes nested within heterotremes (Fig. 5, purple clade). This placement supports previous hypotheses [e.g., (6)], in which Heterotremata is a synonym of Eubrachyura, and includes thoracotremes as a highly derived heterotreme group. On the other hand, if extinct and extant heterotremes form a monophyletic Heterotremata sister to a monophyletic “Thoracotremata,” then the most recent common ancestor of both clades must have been either a heterotreme or a thoracotreme. This raises the question of the relationships between total-group Eubrachyura and their closest nonbrachyuran relatives, e.g., the crown podotremes Raninoidea and Cyclodorippoida that, together with Eubrachyura, form a clade well supported by morphological, molecular, and phylogenomic studies (Fig. 5) [e.g., (7, 8, 66–68)]. As such, the earliest eubrachyurans may have diverged from a common podotreme-like ancestor shared with raninoids and cyclodorippoids but not with other podotremes such as homolodromioids, dromioids, and homolodromioids.

The colonization of nonmarine habitats by true crabs

Exceptionally preserved fossils such as C. athanata from early Cenomanian (99 Ma) Burmese amber are crucial to better constrain the timing of incursions into nonmarine habitats by true crabs during the Cretaceous (7, 69), as well as the origination of modern eubrachyurans and fully carinized forms, i.e., crab-like body forms with a wide and flat carapace, well-developed legs, a reduced muscular pleon folded under the body, and reduced to absent uropods (13).

Molecular phylogenetic studies suggest that crown Thoracotremata—the group containing semiterrestrial and terrestrial crabs—diverged from other eubrachyurans during the Early Cretaceous (8). Unfortunately, the oldest known putative thoracotremes are all marine and Cenozoic in age and consist of carapaces from the lower Eocene (~50 Ma) of Italy [Potamidae, a dorsal carapace (11)] and Peru [Trichodactylidae, isolated claw remains (12)] and from the Oligocene (~25 Ma) of Tanzania [Potamonautidae, a dorsal carapace and chelipeds (71)] and Brazil [Trichodactylidae, isolated claw remains (12)]. The occurrence of Cretapsara (~100 Ma in Burmese amber bridges the gap between the estimated molecular divergence time of nonmarine crabs and their Cretaceous fossil record. Molecular studies show that crown groups of primarily freshwater crabs may have evolved at least three times independently: (i) in the African and Eurasian heterotreme clade formed by Gecarcinidae + (Potamidae + Potamonautidae) (Fig. 5, N to P) and likely sister to Thoracotremata (8), (ii) in the Neotropical family Pseudothelphusidae (Fig. 5M), and (iii) in the Neotropical family Trichodactylidae, which might be closer to portunoids than to other Neotropical freshwater crabs (8). Thus, C. athanata may represent a fourth independent incursion of brachyuran crabs into nonmarine habitats.

Isolated claws of fluvio-lacustrine crabs showing possible affinities with xanthoids or portunoids occur in association with dinosaur bones in the Upper Cretaceous (late Campanian, 74 to 72 Ma) of France (72) and the United States (Milner and Luque, personal observation) (Fig. 5L). Unfortunately, the taxonomic affinities of these specimens are unclear as claw morphology tends to converge with similar dietary habits across groups, and associated carapaces are unknown. These claw remains cannot be assigned with confidence to any crown freshwater group, and thus, they may represent at least one additional incursion into freshwater habitats by crabs during the Mesozoic. Other secondary nonmarine incursions include freshwater and cave-dwelling crabs within otherwise mainly marine families such as Chasmocarcinidae, Hymenosomatidae, Hexapodidae, and Portunidae [e.g., (55, 73–75)]. Brackish waters, estuarine systems, and the intertidal zone, which are transitional between open marine and fully land and freshwater habitats, must have played a crucial role in the evolution of terrestrial, semiterrestrial, and freshwater crabs during the Cretaceous Crab Revolution and the onset of modern clades and body plans (7, 69).

MATERIALS AND METHODS

Experimental design

The main objective of this study was to fully characterize LYAM-9 and place it in a phylogenetic context as a basis for interpreting the habitat and preservational history of the specimen. To fully describe LYAM-9 and provide characters for phylogenetic analysis, we used approaches ranging from standard macro- and microscopic observations of external structures, to x-ray micro-CT scanning and observations of 3D models.

Materials and imaging

The amber sample studied (Longyin Amber Museum, Xishan District, Kunming, Yunnan Province, China) measures 31.5 mm by 19.8 mm by 7.0 mm, weighs 3.05 g, and contains one complete, small brachyuran crab specimen, holotype LYAM-9. The specimen was tested by the National Gemological Training Center in China, and all of its features are consistent with Burmese amber from the Hukawng Valley in the Kachin State of Myanmar. This published work and the nomenclatural acts have been registered in ZooBank, the official registry of Zoological Nomenclature. The LSID for this publication is
The age of the Burmese amber coming from the region where holotype LYAM-9 originates has been estimated as late Albian–early Cenomanian (latest Early Cretaceous to earliest Late Cretaceous, ~105 to 95 Ma), on the basis of biostratigraphic evidence (aminationites and pollen) from the Angbamo site (17, 18). This estimate is consistent with radiogenic isotope data (U-Pb) from volcanioclastic zircons attached to amber clasts, which indicate a maximum age of 98.8 ± 0.6 Ma (19) for the matrix surrounding amber in the region, but not necessarily the amber it contains, which could be either of the same age as the matrix that contains it or slightly older if they have been eroded and redeposited in younger strata.

Specimen microphotography and observations were conducted with a Leica SZ 12.5 stereomicroscope equipped with a Canon EOS Rebel T6i camera and a Z-stacking imaging system, which consists of a computer-driven stage with a Canon EOS 5D DSLR camera equipped with a Canon MP-E 65-mm lens and studio lighting. Submersion in glycerin was used to reduce refraction and optical distortions due to the curvature of amber surfaces.

The amber inclusion was scanned with a Micro-X-ray–CT Xradia 520 Versa (Carl Zeiss X-ray Microscopy Inc., Pleasanton, USA) at the Yunnan Key Laboratory for Palaeobiology, Yunnan University, Kunming, China. The entire amber piece was scanned with a beam power of 50 kV/4 W, with the LE3 filter for 390 min. The exposure time of each projection was 5.5 s. The voxel size was 8.35 μm. The sample was rotated 360° during the scan, and a total of 3201 projections were acquired. All of the projections were used for reconstruction through the automatic reconstruction function of the instrument’s own software (Zeiss Scout-and-Scan Control System 11.1.7859). The center shift (pixels) was 2.216, beam hardening correction constant was 0.05, and smooth constant was 0.5. From the reconstruction, a total of 1012 radiographs were registered and saved as TIFF images for further processing. On the basis of the obtained image stacks, structures of the specimen were rendered using Dristhi 2.6.4 and Amira 5.4 (Visage Imaging, San Diego, USA). The subsequent volume rendering and animations were performed using VGSTUDIO MAX 3.0 (Volume Graphics, Heidelberg, Germany). The surface morphology of the specimen was extracted as a watertight polygon mesh and imported into MeshLab where small isolated polygonal faces were removed. We digitally removed the objects (bubbles and particulates) obscuring the dorsal view of the specimen using MeshLab. Subsurface polygons of the mesh of the left eye and the mesh of the tip of the left claw were removed, and the remaining faces were remeshed via the Screened Poisson surface reconstruction technique to fill holes and smooth the surface. The original and the new meshes of the eye and the claw were imported into Maya (Autodesk, San Rafael, USA), where they were superimposed. Copies of the eye, claw, and right pereopod 5 mesh were reflected across the body plane in Maya and repositioned to create the right eye, the right claw, and the detached left leg. The surface morphology of the gills was extracted in the same manner as the rest of the body; small isolated polygonal faces were removed using MeshLab.

**Phylogenetic analysis**

The dataset, containing 30 taxa and 93 adult morphological characters, was built in MorphoBank (data available at http://morphobank.org/permalink/?P4044), modified from (7). Underdetermined characters and those not preserved were scored as “?,” and inapplicable characters were scored as “—.” Multiple character states present in a given terminal were scored as polymorphisms.

Bayesian inference was used on the dataset, as implemented in MrBayes v. 3.2.6 (76). The dataset was analyzed under the traditional Mk model (77) with an ascertainment bias correction to account for scoring only variable morphological characters. Each analysis was performed with two independent runs of 3 × 10⁶ generations each. We used the default settings of four chains (one cold and three heated) per independent run. The relative burn-in fraction was set to 50%, and the chains were sampled every 1000 generations. We set the temperature parameter to 0.01 as determined by preliminary runs to achieve chain mixing values in the optimal range (0.4 to 0.8). Convergence of independent runs was assessed through the average standard deviation of split frequencies (ASDSF << 0.01) and potential scale reduction factors [PSRF ≈ 1 for all parameters (78)]. We used Tracer v.1.6 (79) to determine whether the runs reached stationary phase and to ensure that the effective sample size for each parameter was greater than 200. Results of the Bayesian runs were summarized as a majority-rule consensus tree of the post–burn-in sample with a node support threshold of 75% (nodes with posterior probability support <75% were collapsed).

**Supplementary materials**

Supplementary material for this article is available at https://science.org/doi/10.1126/sciadv.abj5689

View/request a protocol for this paper from Bio-protocol.

**References and notes**

1. G. J. Vermeij, R. Dudley, Why are there so few evolutionary transitions between aquatic and terrestrial ecosystems? Biol. J. Linn. Soc. 79, 541–554 (2000).
2. J. Lozano-Fernandez, R. Carter, A. R. Tanner, M. N. Puttick, M. Blaxter, J. Vinther, J. Olesen, G. Giribet, G. D. Edgecombe, D. Pisani, A molecular palaeobiological exploration of arthropod terrestrialization. Philos. Trans. R. Soc. B Biol. Sci. 371, 20150133 (2016).
3. G. D. Edgecombe, C. Strullu-Derrien, T. Góral, A. J. Hetherington, C. Thompson, M. Koch, Aquatic stem group myriapods close a gap between molecular divergence dates and the terrestrial fossil record. Proc. Natl. Acad. Sci. U.S.A. 117, 8966–8972 (2020).
4. P. J. F. Davie, D. Guinot, P. K. L. Ng. Treatise on Zoology-Anatomy, Taxonomy, Biology, The Crustacea, Volume 9 Part 3 (2 vols) (Brill, 2015), pp. 11–163.
5. J. Luque, W. T. Allison, H. D. Bracken-Grissom, K. M. Jenkins, A. R. Palmer, M. L. Porter, J. M. Wolfe, Evolution of crab eye structures and the utility of ommatidia morphology in resolving phylogeny. bioRxiv, preprint (2019).https://doi.org/10.1101/786010.781101/786087.
6. R. Von Sternberg, N. Cumberlidge, On the heterotreme-thoracotreme distinction in the Eubrachyura de Saint Laurent, 1980 (Decapoda, Brachyura). Crustaceana 74, 321–338 (2001).
7. J. Luque, R. M. Feldmann, W. Vernygora, C. E. Schweitzer, C. B. Cameron, K. A. Kerr, F. J. Vega, A. Duque, M. Strange, A. R. Palmer, C. Jaramillo, Exceptional preservation of mid-Cretaceous marine arthropods and the evolution of novel forms via heterochrony. Sci. Adv. 5, eaav3875 (2019).
8. L. M. Tsang, D. C. Schubart, S. T. Ahyong, J. C. Y. Lai, E. Y. C. Au, T. Y. Chan, P. K. L. Ng, K. H. Chu, Evolutionary history of true crabs (Crustacea: Decapoda: Brachyura) and the origin of freshwater crabs. Mol. Biol. Evol. 31, 1173–1187 (2014).
9. J. M. Wolfe, J. W. Breinholt, K. A. Crandall, A. R. Lemmon, E. M. Lemmon, L. E. Timm, M. E. Siddall, H. D. Bracken-Grissom, A phylogenomic framework, evolutionary timeline and genomic resources for comparative studies of decapod crustaceans. Proc. R. Soc. B 286, 20190079 (2019).
10. K. Y. Ma, J. Qin, C.-W. Lin, T.-Y. Chan, P. K. L. Ng, K. H. Chu, L. M. Tsang, Phylogenetic analyses of brachyuran crabs support early divergence of primary freshwater crabs. Mol. Phylog. Evol. 135, 62–66 (2019).
11. A. De Angeli, F. Caporiondo, Alontecarcinus buratoi n. gen., n. sp. (Decapoda, Brachyura, Potamonidae) un nuovo crostaceo d’acqua dolce dell’Eocene (Bartoniano) di Alonte (Monti Berici, Vicenza, Italia settentrionale). Bollettino del Museo Civico di Storia Naturale di Verona 43, 5–16 (2019).
12. S. Klaus, C. Magalhães, R. Salas-Gismondi, M. Gross, P.-O. Antoine, Paleogene and Neogene brachyurans of the Amazon Basin: A revised first appearance date for primary freshwater crabs (Crustacea, Brachyura, Trichodactylidae). Crustaceana 90, 953–967 (2017).
17. R. D. Cruickshank, K. Ko, Geology of an amber locality in the Hukawng Valley, Northern Myanmar. Asian Earth Sci. 21, 441–455 (2003).

18. A. J. Ross, M. C. Shearman, J. W. Jagt, R. H. Fraaije, New Cretaceous crabs (Crustacea, Brachyura) from the Kerguelen Islands (Miocene). J. Crust. Biol. 26, 127–132 (2006).

19. V. Perrichot, V. Girard, A. R. Schmidt, S. Saint Martin, S. Struwe, V. Perrichot, J. P. Saint Martin, J. Asian Earth Sci. 109, 155–163 (2012).

20. A. J. Ross, D. A. Grimaldi, J. W. Jagt, A. J. Ross, Extraordinary Lagerstätten in Amber, with particular reference to the Cretaceous of Burma, in Terrestrial Conservation Lagerstätten: Windows into the Evolution of Life on Land, N. C. Fraser, H. D. Sues, Eds. (Dunedin Acad. Press Ltd., 2017), pp. 287–342.

21. F. J. Vega, L. Xing, Y. Su, H. Zhang, F. Xia, Y. Fu, C. Sai, D. Huang, Various ambergard marine animals on Burmese amber with discussions on its age. Palaeontologia 1, 91–103 (2010).

22. A. J. Poinar Jr., J. B. Lambert, A. Shi, A catalogue of Burmite fossil inclusions from Burma in a recent study. Palaeoentomology 1, 91–103 (2010).

23. W. Burggren, A. Pinder, B. McMahon, M. Wheatley, M. Doyle, Ventilation, circulation and their interactions in the land crab, Cardiodes guanhumi. J. Exp. Biol. 117, 133–154 (1985).

24. B. W. van Bakel, E. V. Mychko, A. Spiridonov, J. W. Jagt, R. H. Fraaije, New Cretaceous crabs (Crustacea, Brachyura) from Moscow Oblast and Dagestan (Russia): Patterns in phylogeny and morphospace of the oldest eubrachyurans (Dorippoidea). Cretac. Res. 117, 104675 (2021).

25. D. Guinot, G. Breton, I. decapodi dell’area di Bolca: Verona e Vicenza Italia nordorientale. Geodiversitas 39, 2100020 (2021).

26. C. Beschin, A. Busulini, G. Tessier, R. Zorzini, I crustacei associati a coralli nell’Eocene inferiore dell’area di Bolca: Verona e Vicenza, Italia nordorientale (Memorie del Museo Civico di Storia Naturale di Verona. 2016). vol. 2. serie. Sezione Scienze della Terra – N. 9, pp. 189.

27. I. S. Wehrtmann, C. Magalhães, O. C. Bello-González, First confirmed report of a primary invertebrate in mid-Cretaceous amber from Myanmar suggests a cyclophoroidean (Gastropoda) ancestry. Cretac. Res. 93, 114–119 (2019).

28. J. Luque, The oldest higher true crabs (Crustacea: Decapoda: Brachyura): Insights of toxic metals. Plankton Benthos Res. 11, 306–316 (2020).

29. C. Beschin, A. Busulini, G. Tessier, R. Zorzini, I crustacei associati a coralli nell’Eocene inferiore dell’area di Bolca: Verona e Vicenza, Italia nordorientale (Memorie del Museo Civico di Storia Naturale di Verona. 2016). vol. 2. serie. Sezione Scienze della Terra – N. 9, pp. 189.

30. M. de Lourdes Serrano-Sánchez, G. Guerao, E. Centeno-García, F. J. Vega, Crabs (Brachyura: Grapsoidae: Sesarmidae) as inclusions in Lower Miocene amber from Chiapas, Mexico. Bol. Soc. Geol. Mex. 68, 37–43 (2016).

31. D. A. Grimaldi, M. A. Coutiño, Eds. (Dunedin Acad. Press Ltd., 2017), vol. 117, 104675 (2021).

32. C. Beschin, A. Busulini, G. Tessier, R. Zorzini, I crustacei associati a coralli nell’Eocene inferiore dell’area di Bolca: Verona e Vicenza, Italia nordorientale (Memorie del Museo Civico di Storia Naturale di Verona. 2016). vol. 2. serie. Sezione Scienze della Terra – N. 9, pp. 189.
66. J. Chen, Y. Xing, W. Yao, X. Xu, C. Zhang, Z. Zhang, Q. Liu, Phylomitoigenomics reconfirm the phylogenetic position of the genus Metaphix inferred from the two grapsid crabs (Decapoda: Brachyura: Grapsoidae). PLOS ONE 14, e0210763 (2019).

67. B. P. Tang, Y. Liu, Z. Z. Xin, D. Z. Zhang, Z. F. Wang, X. Y. Zhu, Y. Wang, H. B. Zhang, C. L. Zhou, X. Y. Choi, Q. N. Liu, Characterisation of the complete mitochondrial genome of Helice wuana (Grapsoidae: Varunidae) and comparison with other Brachyuran crabs. Genomics 110, 221–230 (2018).

68. S. T. Ahyong, J. C. Y. Lai, D. Sharkey, D. J. Colgan, P. K. L. Ng, Phylogenetics of the brachyuran crabs (Crustacea: Decapoda): The status of Podotremata based on small subunit nuclear ribosomal RNA. Mol. Phylogenet. Evol. 45, 576–586 (2007).

69. J. Luque, L. Xing, R. C. McKellar, A. Osso, S. Charbonnier, G. R$. Oliveira, W. Santana, A. P. Pinheiro, A. Saria, E. G. Clark, D. Briggs, in 7th Symposium on Cretaceous and Cenozoic Decapod Crustaceans (I, jubilana, Slovenia, 2019).

70. D. Lima, M. Tavares, R. T. Lopes, O. M. O. de Araújo, A. Aguilera, Uca maracoani (Crustacea, Decapoda, Ocypodidae) from a Miocene paleomangrove in Brazil: A case of evolutionary stasis among tropical American fiddler crabs. J. South Am. Earth Sci. 99, 102517 (2020).

71. R. M. Feldmann, P. M. O’Connor, N. J. Stevens, M. D. Gottfried, E. M. Roberts, S. Ngasala, L. Xing, D. E. G. Briggs, E. G. Clark, A. Duque, J. Hui, H. Mai, R. C. McKellar, Crab discovery grants 2021-04174 (R.C.M.). Ethics statement: The fossil crab in amber studied here (LYAM-9) is housed in the Longyin Amber Museum (LYAM), Kunming, Yunnan Province, China. It comes from a batch of commercial “raw” amber pieces collected by local miners and sold to a vendor at an amber jewelry market in Myitkyina on 12 May 2015. The polished piece containing specimen LYAM-9 was acquired by LYAM from the vendor's mineral store in Tengchong, China, on 10 August 2015. We acknowledge the existence of a sociopolitical conflict in northern Myanmar and have limited our research to material predating the 2017 resumption of hostilities in the region. We hope that conducting research on specimens collected before the conflict and acknowledging the situation in the Kachin State will serve to raise awareness of the current conflict in Myanmar and the human cost behind it. Author contributions: L.X. and J.H. provided specimen access. J.L. and L.X. conceived and designed the study. H.M. arranged CT scans. L.X. and J.H. prepared the initial description of the specimen studied, and J.L. completed the final description and manuscript. J.L. made the figures and analyzed the data. E.G.C. and A.D. designed the digital reconstructions and animations. L.X., D.E.G.B., J.H., H.M., and R.C.M. contributed laboratory/logistics/materials/analysis tools. All authors edited the manuscript and confirmed the content of the paper. Competing interests: The authors declare that they have no competing interests. Data and materials availability: All data needed to evaluate the conclusions in the paper are present in the paper and/or the Supplementary Materials. Data for this study are available in MorphoBank: http://morphobank.org/permalink/?P4044. This published work and the nomenclatural acts it contains have been registered in ZooBank: http://zoobank.org/urn:lsid:zoobank.org:act:F2CC5291-0DE0-4B21-AB16-9F7B570BAD8E.

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