

# Enigmatic dinosaur precursors bridge the gap to the origin of Pterosauria

Martín D. Ezcurra<sup>1,2✉</sup>, Sterling J. Nesbitt<sup>3</sup>, Mario Bronzati<sup>4</sup>, Fabio Marco Dalla Vecchia<sup>5,6</sup>, Federico L. Agnolin<sup>7,8</sup>, Roger B. J. Benson<sup>9</sup>, Federico Brissón Egli<sup>7</sup>, Sergio F. Cabreira<sup>10</sup>, Serjoscha W. Evers<sup>9,11</sup>, Adriel R. Gentil<sup>7</sup>, Randall B. Irmis<sup>12,13</sup>, Agustín G. Martinelli<sup>1</sup>, Fernando E. Novas<sup>7</sup>, Lúcio Roberto da Silva<sup>10</sup>, Nathan D. Smith<sup>14</sup>, Michelle R. Stocker<sup>3</sup>, Alan H. Turner<sup>15</sup> & Max C. Langer<sup>4</sup>

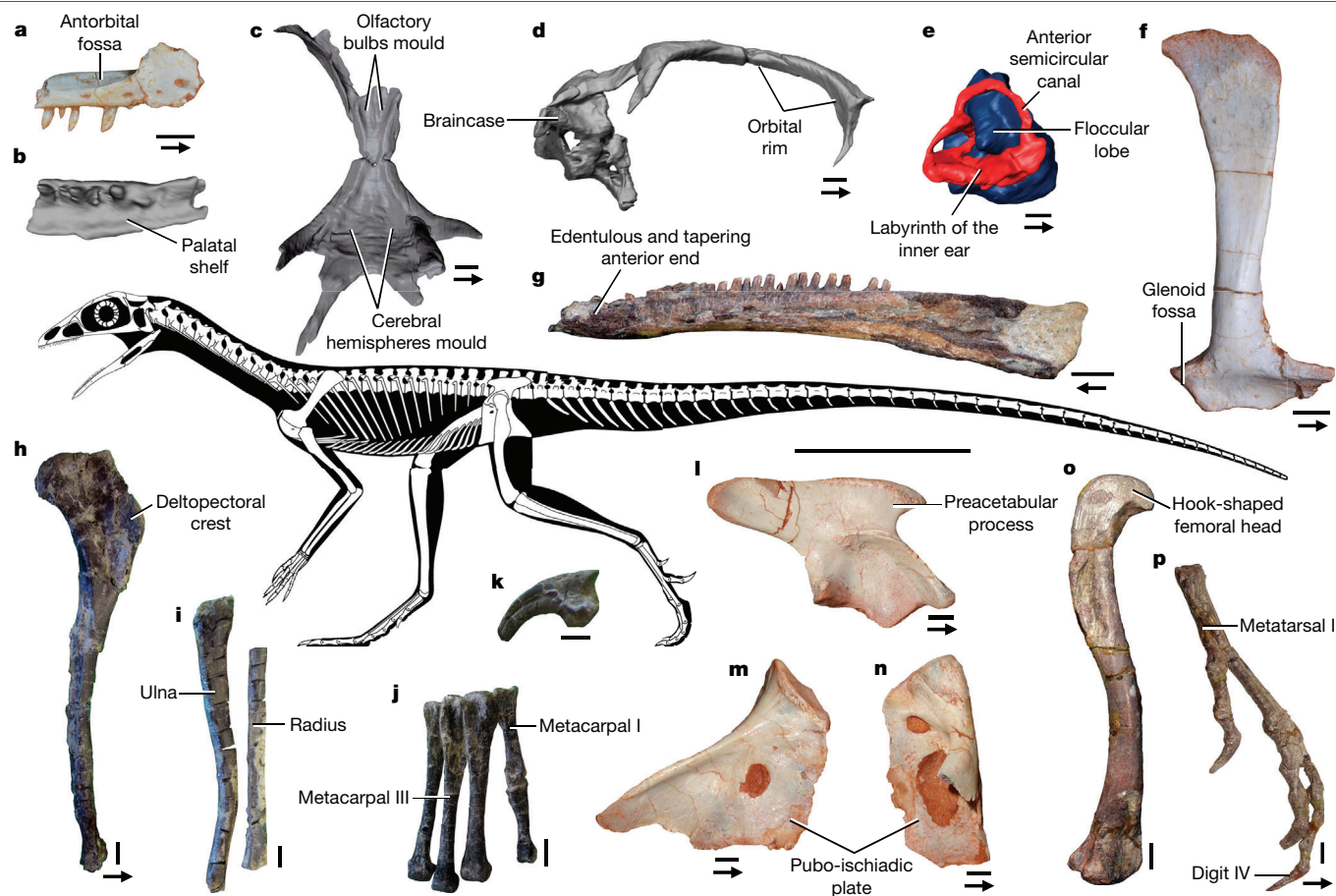
Pterosaurs were the first vertebrates to evolve powered flight<sup>1</sup> and comprised one of the main evolutionary radiations in terrestrial ecosystems of the Mesozoic era (approximately 252–66 million years ago), but their origin has remained an unresolved enigma in palaeontology since the nineteenth century<sup>2–4</sup>. These flying reptiles have been hypothesized to be the close relatives of a wide variety of reptilian clades, including dinosaur relatives<sup>2–8</sup>, and there is still a major morphological gap between those forms and the oldest, unambiguous pterosaurs from the Upper Triassic series. Here, using recent discoveries of well-preserved cranial remains, microcomputed tomography scans of fragile skull bones (jaws, skull roofs and braincases) and reliably associated postcrania, we demonstrate that lagerpetids—a group of cursorial, non-volant dinosaur precursors—are the sister group of pterosaurs, sharing numerous synapomorphies across the entire skeleton. This finding substantially shortens the temporal and morphological gap between the oldest pterosaurs and their closest relatives and simultaneously strengthens the evidence that pterosaurs belong to the avian line of archosaurs. Neuroanatomical features related to the enhanced sensory abilities of pterosaurs<sup>9</sup> are already present in lagerpetids, which indicates that these features evolved before flight. Our evidence illuminates the first steps of the assembly of the pterosaur body plan, whose conquest of aerial space represents a remarkable morphofunctional innovation in vertebrate evolution.

Pterosaurs are deeply rooted in popular culture, are frequently portrayed in books and films<sup>10</sup>, and include the largest flying animals ever known<sup>11</sup>. Their evolutionary history spans more than 150 million years, ending at the close of the Mesozoic era<sup>10,11</sup>. The oldest pterosaurs are from the Upper Triassic series (219–201.3 million years ago) of Europe and North America<sup>12,13</sup>, and the clade diversified into multiple ecomorphologically disparate groups by the Middle–Late Jurassic epochs<sup>11,14</sup>. The oldest recognized pterosaurs already had a highly specialized body plan linked to their ability to fly<sup>12,15</sup>, which was conserved in all pterosaurs: shoulder girdle with strongly posteroventrally enlarged coracoid braced with the sternum and laterally facing glenoid fossa; forelimb with pteroid bone and hypertrophied fourth digit supporting a membranous wing; and pelvic girdle with prepubic bone and strongly developed preacetabular process<sup>1</sup>. This highly modified anatomy results in a large morphological gap between pterosaurs and all other known

Mesozoic reptiles. To complicate matters, early pterosaur specimens are small, scarce and generally represented by taphonomically compressed, almost bidimensional partial skeletons<sup>12</sup>. These preservational problems and the absence of fossils with transitional morphologies have made the origin of pterosaurs one of the most elusive questions in vertebrate evolution for more than 200 years.

Quantitative phylogenetic analyses have generally agreed that pterosaurs fall within the lineage of Archosauria (that is, the least inclusive clade that contains birds and crocodylians) leading to dinosaurs (that is, Pan-Aves = Avemetatarsalia)<sup>2,3,5–8</sup>. However, some studies have alternatively recovered pterosaurs as the sister group to all other pan-archosaurs<sup>16</sup>, among tanystropheid archosauromorphs<sup>17</sup>, among non-archosaurian archosauriforms<sup>18</sup> or sister to the probable stem-diapsid drepanosauromorphs<sup>19</sup>. New fossil discoveries over the past few years have greatly increased the understanding of the

<sup>1</sup>Sección Paleontología de Vertebrados CONICET–Museo Argentino de Ciencias Naturales ‘Bernardino Rivadavia’, Buenos Aires, Argentina. <sup>2</sup>School of Geography, Earth and Environmental Sciences, University of Birmingham, Birmingham, UK. <sup>3</sup>Department of Geosciences, Virginia Tech, Blacksburg, VA, USA. <sup>4</sup>Departamento de Biología, Universidade de São Paulo, Ribeirão Preto, Brazil. <sup>5</sup>Research Group of Mesozoic Faunas, Institut Català de Paleontologia Miquel Crusafont (ICP), Sabadell, Spain. <sup>6</sup>Museo Friulano di Storia Naturale, Udine, Italy. <sup>7</sup>Laboratorio de Anatomia Comparada y Evolución de los Vertebrados CONICET–Museo Argentino de Ciencias Naturales ‘Bernardino Rivadavia’, Buenos Aires, Argentina. <sup>8</sup>Fundación de Historia Natural ‘Félix de Azara’, Departamento de Ciencias Naturales y Antropología, Universidad Maimónides, Buenos Aires, Argentina. <sup>9</sup>Department of Earth Sciences, University of Oxford, Oxford, UK. <sup>10</sup>Associação Sul Brasileira de Paleontologia, Faxinal do Soturno, Brazil. <sup>11</sup>Department of Geosciences, University of Fribourg, Fribourg, Switzerland. <sup>12</sup>Natural History Museum of Utah, University of Utah, Salt Lake City, UT, USA. <sup>13</sup>Department of Geology and Geophysics, University of Utah, Salt Lake City, UT, USA. <sup>14</sup>The Dinosaur Institute, Natural History Museum of Los Angeles County, Los Angeles, CA, USA. <sup>15</sup>Department of Anatomical Sciences, Stony Brook University, Stony Brook, NY, USA. ✉e-mail: martindezcurra@yahoo.com.ar



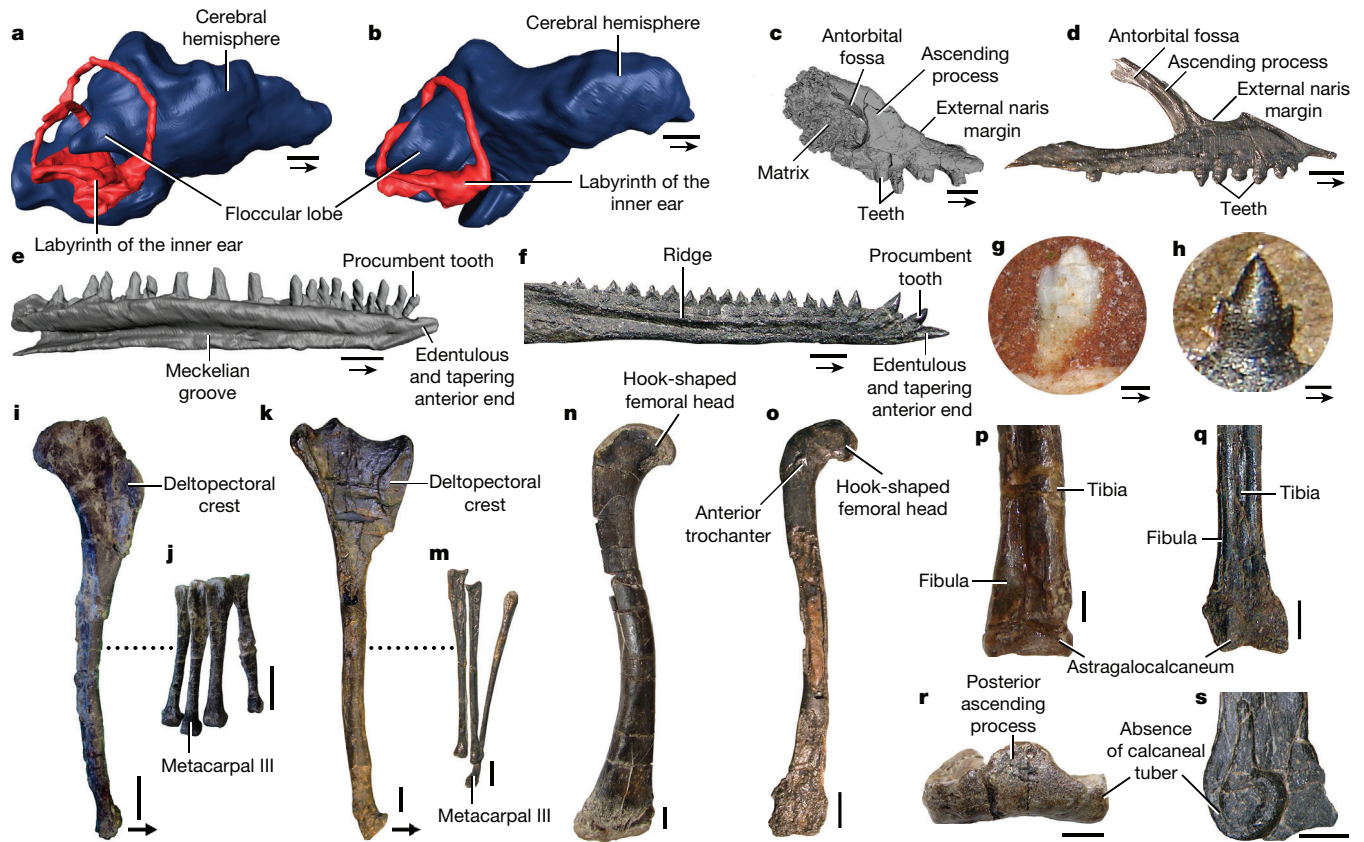
**Fig. 1 | Newly discovered and selected bones characterizing the lagerpetid body plan.** a, b, Partial right maxilla. c, Skull roof. d, Skull roof and braincase. e, Cranial endocast. f, Right scapula. g, Left dentary. h, Left humerus. i, Partial ulna and radius. j, metacarpals I–IV and phalanx I of digit I. k, Ungual of manual digit IV. l, Right ilium. m, Right ischium. n, Left pubis. o, Femur. p, Pes. Images show lateral (a, f, g, k–m), ventral (b, c), right lateral (d, e), medial (h, i, n, p),

dorsal (j) and posteromedial (o) views. a–f, l–n, *Ixalerpeton* (ULBRA-PVT059). g, o, p, *Lagerpeton* (PVL 4625 (g) and PVL 4619 (o, p)). h–k, *Dromomeron romeri* (GR 238). Arrows indicate the anterior direction. Scale bars, 2 mm (a, b, e), 3 mm (c, d, f, g, k–n) and 5 mm (h–j, o, p). Skeletal reconstruction by S. Hartman in collaboration with the authors, based on *Lagerpeton*, *Ixalerpeton* and *D. romeri*.

early evolution of Pan-Aves and of the assembly of the dinosaur body plan<sup>8,20,21</sup>. Nevertheless, a clear morphological gap still remains between pterosaurs and other pan-avians. Here, using new and existing specimens of the enigmatic pan-avian clade Lagerpetidae<sup>21</sup>, we report on a previously undocumented combination of features that reduces the morphological gap between pterosaurs and other reptiles, clarifying the phylogenetic placement of Pterosauria within Pan-Aves. We present anatomical information from across the entire skeleton that demonstrates that lagerpetids are the closest-known evolutionary relatives of pterosaurs. This information derives from detailed first-hand observation of lagerpetid specimens, enhanced by three-dimensional reconstructions from microcomputed tomography scans.

Lagerpetids are small to medium-sized (usually less than 1 m long), gracile and cursorial reptiles from Middle–Upper Triassic rocks of South and North America and Madagascar<sup>21,22</sup>. Previous knowledge of lagerpetid anatomy was mostly limited to vertebrae, hindlimbs and a few cranial bones. Our data are based on improved observations across

*Lagerpeton* and *Ixalerpeton* have an edentulous anterior end that tapers to a point (Figs. 1g, 2e), resembling the condition in the early pterosaurs *Seazzadactylus*, *Carniadactylus* and *Raeticodactylus*<sup>12,23,24</sup> (Fig. 2f), and most silesaurids<sup>20</sup>. The anterior region of the lagerpetid dentary is ventrally curved (Fig. 1g), similar to those of the early pterosaurs *Austriadraco* and *Peteinosaurus*<sup>12,23,24</sup>. The lower jaw of *Lagerpeton* preserves articulated dentaries and lack splenials (Extended Data Fig. 2); the latter bone is fused to the dentary or restricted to the mid-point of the medial surface of the mandible in pterosaurs<sup>13</sup>. *Lagerpeton* and *Ixalerpeton* have 26–27 dentary tooth positions, sharing the high tooth count (more than 20 teeth) of several early pterosaurs<sup>13,23,24</sup> (Figs. 1g, 2e, f and Extended Data Figs. 1, 2). The dentary tooth crowns of these lagerpetids have convex mesial and distal margins, and middle–distal crowns possess a large and tall central cusp flanked by mesiodistally aligned, small accessory cusps (Fig. 2g and Extended Data Fig. 2). Multicusped tooth crowns are rare among archosauriforms, but also occur in several early pterosaurs (for example, *Austriadraco*,



**Fig. 2 | Key comparisons between pterosaur and lagerpetid cranial endocasts and skeletal elements.** **a, b,** Cranial endocasts. **c,** Partial right maxilla. **d,** Left maxilla (reversed). **e,** Left dentary. **f,** Right dentary. **g, h,** Dentary tooth crowns. **i, k,** Left humeri. **j,** Partial manus. **m,** Metacarpals I–III. **n, o,** Femora (**n**, left, reversed; **o**, right). **p, q, s,** Right distal portion of tibia, fibula and astragalocalcaneum. **r,** Right astragalocalcaneum. Images show right lateral (**a, b**), lateral (**c, d, f**), medial (**e, i, k**), labial (**g, h**), dorsal (**j**), dorsal or ventral (**m**), anterolateral (**n, o, s**), anterior (**p, q**) and posterior (**r**) views. Arrows indicate the anterior direction. Scale bars, 3 mm (**a, b, f, q**), 2 mm (**c, e, s**), 5 mm (**d, k–p, r**), 0.3 mm (**g, h**) and 1 cm (**i, j**).

**a,** Lagerpetid *D. gregorii* (TMM 31100-1334). **b,** Pterosaur *Allkaruen* (MPEF-PV 3613). **c,** Lagerpetid *Kongonaphon* (UA 10618). **d, k, m, o,** Pterosaur *Raeticodactylus* (BNM 14524). **e, g,** Lagerpetid *Ixalerpeton* (ULBRA-PVT059). **h, q,** Pterosaur *Austriadraco* (SNSB-BSPG 1994 I 51). **f,** Pterosaur *Seazzadactylus* (MFSN 21545). **i, j, n, r,** Lagerpetid *D. romeri* (GR 238 (**i, j**), GR 218 (**n**) and GR 223 (**r**)). **p,** Lagerpetid *Lagerpeton* (PULR 06). **s,** Pterosaur *Peteinosaurus* (MCSNB 3496). Arrows indicate the anterior direction. Scale bars, 3 mm (**a, b, f, q**), 2 mm (**c, e, s**), 5 mm (**d, k–p, r**), 0.3 mm (**g, h**) and 1 cm (**i, j**).

other archosauromorphs<sup>6,7</sup>. The cranial endocasts of *D. gregorii* and *Ixalerpeton* show strongly developed and posterolaterally tapering cerebellar floccular lobes, which resemble the even more developed floccular lobes of pterosaurs<sup>9,25</sup> (Figs. 1e, 2a, b). The olfactory tract of *Ixalerpeton* is long and ends in broad olfactory bulbs (Fig. 1c), retaining the ancestral condition of Archosauromorpha<sup>7</sup>, whereas these structures are strongly reduced in pterosaurs<sup>9,25</sup>. In the inner ear, the portion of the semicircular canals of the endosseous labyrinth of *D. gregorii*, *Ixalerpeton*, pterosaurs and some early eusaurischian dinosaurs are taller than anteroposteriorly long, with an anterior semicircular canal that forms a considerably longer arc than the posterior semicircular canal (Fig. 2a, b, Supplementary Information).

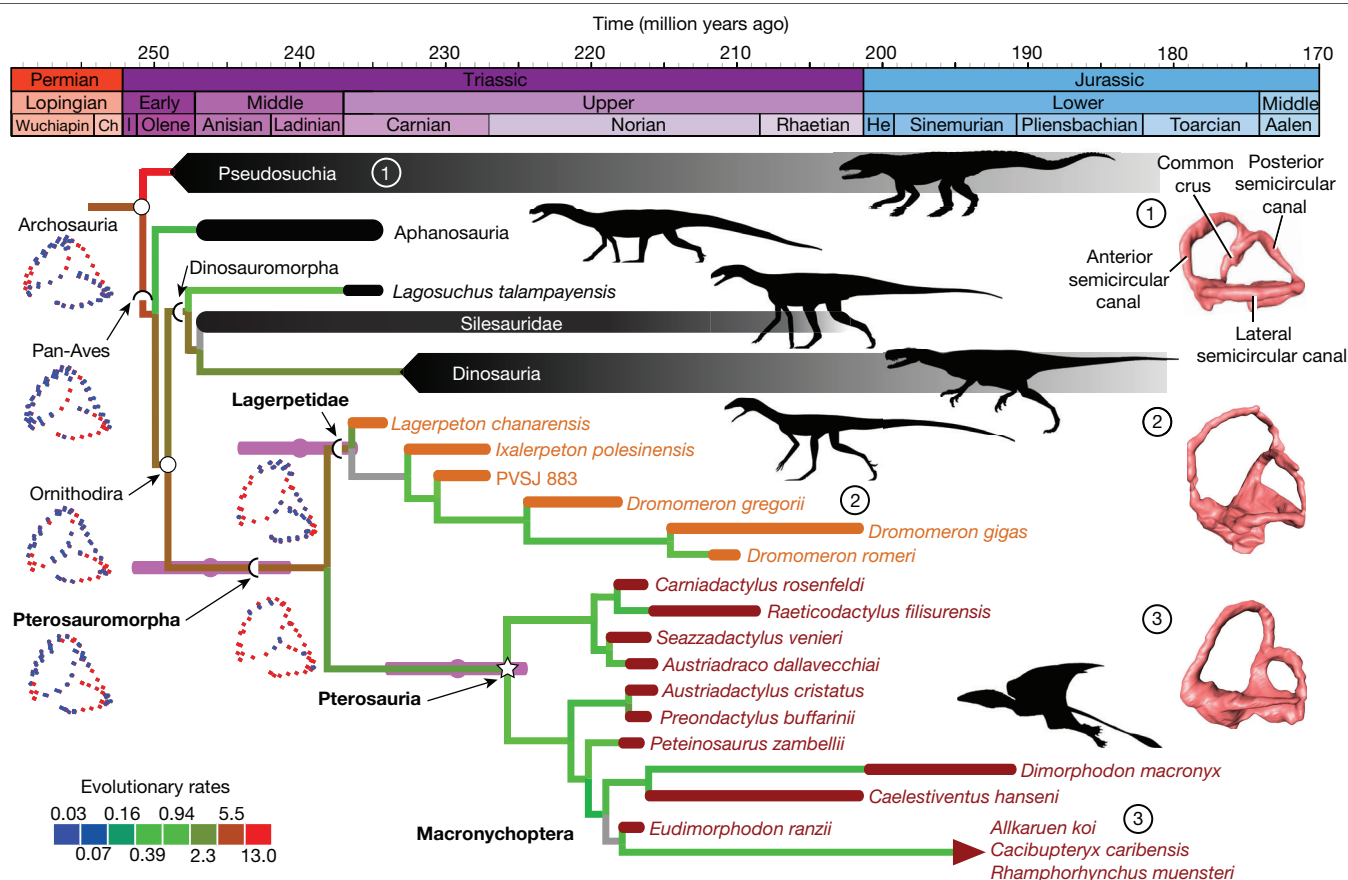
The forelimbs of *Dromomeron romeri* and pterosaurs share a forearm that is longer than the humerus, including a proportionally elongated metacarpus (metacarpal III–humerus length ratio > 0.35) (Fig. 2i–m). The manual digits of *D. romeri* are longer than their respective metacarpal and at least one digit has a trenchant claw (Fig. 1k),

*Dimorphodon*<sup>12,23</sup> (Fig. 1l–n and Extended Data Fig. 3b, c), but not in most other archosaurs<sup>6,7</sup>.

The femora of lagerpetids and early pterosaurs (for example, *Raeticodactylus*, *Peteinosaurus* and *Dimorphodon*) share a hook-shaped proximal head (Figs. 1o, 2n, o and Extended Data Fig. 3c, d). A co-ossified astragalus and calcaneum is present in both lagerpetids and pterosaurs (Fig. 2p–s), but also in heterodontosaurid ornithischians and early neotheropods<sup>6,7</sup>. The absence of both a posterior groove on the astragalus and a calcaneal tuber is shared by pterosaurs and lagerpetids (Fig. 2r, s) and independently arose in some silesaurids and early dinosaurs<sup>6</sup>.

The new anatomical information available for lagerpetids was scored in an expanded version of a comprehensive phylogenetic data matrix focused on Permo-Triassic pan-archosaurs<sup>7</sup>. Our data matrix comprises 157 species (or diagnostic specimens) scored across 822 characters, including all currently valid lagerpetid species, 9 Triassic and 4 Jurassic pterosaurs, 13 early dinosauriforms, and most non-archosaurian





**Fig. 3 | Time-calibrated reduced strict consensus tree (after a posteriori pruning of Kongonaphon) focused on Pterosauria and Lagerpetidae.** Red dotted structures are ancestral optimizations of the labyrinth of the inner ear (blue lines are apomorphic displacements). Violet bars indicate node temporal calibration after Bayesian analysis. White-fill circles indicate node-based

clades, half-circles stem-based clades and the star an apomorphy-based clade. Labyrinths of the inner ear are in lateral view and belong to (1) *Arizonasaurus*, (2) *D. gregorii* and (3) *Allkaruen*. Further information is provided in the Supplementary Information. Ch, Changhsingian; I, Induan; He, Hettangian.

model that incorporates taxon ages using a fossilized birth–death process (posterior probability = 0.99 for the lagerpetid–pterosaur clade) (Extended Data Fig. 6). A minimum of 33 synapomorphies distributed across the skeleton provide strong support for the Lagerpetidae and Pterosauria clade (= Pterosauriomorpha). Some of these synapomorphies are unique to pterosauriormorphs among early archosaurs, including a subtriangular and dorsoventrally tall floccular fossa of the braincase, a height–anteroposterior length ratio of the semicircular canals of the inner ear  $>0.90$ , a reduced to absent splenial, strongly ventrally extended pubo–ischial plate, and hook-shaped femoral head (a complete list of synapomorphies can be found in the Supplementary Information). Bayesian inference analyses indicate high rates of morphological change during deep pan-avian divergences, including on the branches leading to Pterosauriomorpha and Lagerpetidae. These high rates contrast with the more typical ‘background’ rates found on the branch leading to Pterosauria and its internal branches (Fig. 3). By contrast, higher evolutionary rates occur on the pterosaur branch if we force the more traditional position of lagerpetids closer to dinosaurs (Extended Data Fig. 7). These results strengthen the idea that

23 additional steps are needed to force the more traditional position of lagerpetids as the sister taxon to Dinosauriformes. We also tested previous alternative hypotheses for pterosaur relationships, finding that 44 extra steps are required to place them as non-archosaurian archosauriforms and 64 steps to force their placement as sister taxon to Tanystropheidae (but the Pterosauria and Tanystropheidae clade is sister to Lagerpetidae within Pan-Aves). These results agree with a poorly supported placement of Pterosauria outside Archosauria<sup>6,7</sup>. The possible pterosaur-relative<sup>2,3</sup>, but highly problematic taxon *Scle-romochlus* (Upper Triassic of Scotland), was included in a secondary phylogenetic analysis (Methods) and was recovered as the sister taxon to the Pterosauria and Lagerpetidae clade within Pan-Aves (Extended Data Fig. 4).

Additional cladistic analyses included the geometric morphometric configuration of the labyrinth of the inner ear as a continuous three-dimensional character in an attempt to avoid subjective discrete character states on such a complex structure<sup>26</sup> (Supplementary Information). The analyses suggest that the inner ear has a strong phylogenetic signal that supports a pterosaur–lagerpetid affinity (Fig. 3



and highly rounded semicircular canals as a trait combination that is uniquely shared by pterosaurs and lagerpetids (Extended Data Fig. 10c). Lagerpetids and pterosaurs exhibit a strong curvature of the anterior semicircular canal (arc-length versus straight-line ratio >1.5) that results in an increased radius, which convergently evolved in a few early saurischian dinosaurs and is present in birds. The semicircular canals detect head movements (angular acceleration) and a larger radius increases the sense of equilibrium in primates and birds, and therefore is inferred to be related to arboreal, aerial or other agile forms of terrestrial locomotion and rapid movements<sup>27</sup>. Lagerpetids and pterosaurs are the only archosauriforms with a greatly enlarged floccular fossa, representing more than 40% the height of the endocranial cavity (Fig. 2a, b). A large flocculus might have been important for lagerpetid locomotion or predation, given the connection of this structure with coordination of eye, head and neck movements<sup>28</sup>. In pterosaurs, floccular enlargement has been hypothesized to be important for information processing related to flight<sup>9</sup>. Although not as developed as in pterosaurs, the relatively large lagerpetid flocculus indicates that initial enlargement of that structure occurred along the Pterosauriiform branch. Thus, the neuroanatomy of lagerpetids has intermediate features between those of pterosaurs and other Triassic archosauriforms and may have paved the way towards the origin of active flight in pterosaurs.

The evolution of other traits associated with the acquisition of flight remains difficult to trace at the base of Pterosauriiform with the currently available fossil record. Lagerpetid girdles and limbs lack features correlated with the flying behaviour of pterosaurs, for example, a hypertrophied deltopectoral crest of the humerus and wing digit of the manus<sup>1</sup>. However, the strongly recurved manual claws, with an inner curvature of more than 150° in *D. romeri* (Fig. 1k), suggests that the forelimbs had functions other than ground-dwelling locomotion, such as climbing or acquisition of prey (the inner claw curvature of ground-dwelling birds and squamates ranges from 21.5° to 125.5° and in perchers, climbers and predators ranges from 87° to 170.3°)<sup>29</sup>. Non-archosaurian archosauriforms, early crocodylian-line archosaurs and very probably aphanosaurians (the earliest pan-avians) were all quadrupedal, ground-dwelling animals<sup>30</sup>. Our observations suggest that functional forelimb versatility became widespread in ornithomirans (Fig. 3), allowing the evolution of disparate behaviours such as manual processing of food resources in dinosaurs and active flight in pterosaurs.

The recognition of lagerpetids as the sister taxon to pterosaurs provides a new framework to study the origin of Pterosauria, its specialized body plan and flying abilities. Previous phylogenetic hypotheses implied a long ghost lineage, with a minimum of 28 million years, for Pterosauria<sup>8</sup>. Our results shorten this to about 18 million years, because the oldest lagerpetids come from 237-million-year-old rocks, near the Ladinian–Carnian boundary<sup>22</sup> (Fig. 3). Along those millions of years, the enhancement of features that were already present in the common ancestor of lagerpetids and pterosaurs allowed the latter group to explore a new adaptive landscape and conquer aerial space, which probably prompted their impressive adaptive radiation.

## Online content

Any methods, additional references, Nature Research reporting summaries, source data, extended data, supplementary information, acknowledgements, peer review information; details of author contributions

and competing interests; and statements of data and code availability are available at <https://doi.org/10.1038/s41586-020-3011-4>.

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## Methods

### New phylogenetic definitions

**Pan-Aves Gauthier and de Queiroz, 2001** (this study), converted clade name

**Registration number:** 404.

**Phylogenetic definition.** The most inclusive clade containing *Vultur gryphus* Linnaeus, 1758 (Aves), but not *Crocodylus niloticus* Laurenti, 1768 (Crocodylia). This is a crown-based total-clade definition.

**Reference phylogeny.** Phylogenetic hypothesis shown in Fig. 3 and Extended Data Fig. 4. *Vultur gryphus* nests within Dinosauria and *C. niloticus* within Pseudosuchia.

**Composition.** The composition is based on the reference phylogeny, Pan-Aves includes Aphanosauria<sup>8</sup> and Ornithodira (see 'Ornithodira').

**Ornithodira Gauthier, 1986** (this study), converted clade name

**Registration number:** 405.

**Phylogenetic definition.** The least inclusive clade containing *Compsognathus longipes* Wagner, 1859 (Dinosauria), and *Pterodactylus* (originally *Ornithocephalus*) *antiquus* Sömmerring, 1812 (Pterosauria), but not *Alligator* (originally *Crocodylus*) *mississippiensis* Daudin, 1802 (Crocodylia). This is a minimum clade definition.

**Reference phylogeny.** Phylogenetic hypothesis shown in Fig. 3 and Extended Data Fig. 4. *Compsognathus longipes* nests within Dinosauria, *P. antiquus* within Pterosauria and *A. mississippiensis* within Pseudosuchia.

**Composition.** The composition is based on the reference phylogeny, Ornithodira includes the main groups Lagerpetidae (see below), Pterosauria<sup>31</sup>, Dinosauromorpha (see 'Dinosauromorpha') and possibly *Scleromochlus taylori*.

**Lagerpetidae Arcucci, 1986** (this study), converted clade name

**Registration number:** 406.

**Phylogenetic definition.** The most inclusive clade containing *Lagerpeton chanarensis* Romer, 1971, but not *Eudimorphodon ranzii* Zambelli, 1973, *Silesaurus opolensis* Dzik, 2003, and *V. gryphus* Linnaeus, 1758. This is a maximum clade definition.

**Reference phylogeny.** Phylogenetic hypothesis shown in Fig. 3 and Extended Data Fig. 4.

**Composition.** The composition is based on the reference phylogeny, Lagerpetidae includes *L. chanarensis*, *Ixalerpeton polesinensis*, *Kongonaphon kely* and *Dromomeron* spp.

**Dinosauromorpha Benton, 1985** (this study), converted clade name

**Registration number:** 407.

**Phylogenetic definition.** The most inclusive clade containing *C. longipes* Wagner, 1859 (Dinosauria), but not *P. antiquus* Sömmerring, 1812 (Pterosauria), or *A. mississippiensis* Daudin, 1802 (Crocodylia). This is a maximum clade definition.

**Reference phylogeny.** Phylogenetic hypothesis shown in Fig. 3 and Extended Data Fig. 4. *Compsognathus longipes* nests within Dinosauria, *P. antiquus* within Pterosauria and *A. mississippiensis* within Pseudosuchia.

**Composition.** The composition is based on the reference phylogeny, Dinosauromorpha includes the main groups Silesauridae<sup>32,33</sup> and Dinosauria<sup>34</sup>, including Aves<sup>35</sup>, plus some species level taxa such as *Lagosuchus talampayensis*.

**Dinosauriformes Novas, 1992** (this study), converted clade name

**Registration number:** 408.

**Phylogenetic definition.** The least inclusive clade containing *C. lon-*

### Microcomputed tomography scans and digital processing

The partial lower jaw of *Lagerpeton* (PVL 4625) was scanned using X-ray microcomputed tomography at YPF TECNOLOGÍA (Y-TEC) using a Bruker Skyscan, the bones of *Ixalerpeton* (ULBRA-PVT059) were scanned at Centro para Documentação da Biodiversidade, Universidade de São Paulo using a Nanotom Scan machine (GE Sensing & Inspection Technologies) and the braincase of *D. gregorii* (TMM 31100-1334) was scanned at the University of Texas High-Resolution X-ray CT Facility using a custom-built BIR scanner using a Feinfocus X-ray source and an Image Intensifier detector (further information about the scans is provided in the Supplementary Information). The images of *Lagerpeton* were processed using the software 3D Slicer version 4.7<sup>36</sup> and the images of *Ixalerpeton* and *Dromomeron* were processed using the software Amira (version 5.3.3, Visage Imaging).

### Morphogeometric sampling of the endosseous labyrinth of the inner ear

The course of each semicircular canal of the inner ear was quantitatively sampled in available taxa using sliding three-dimensional semilandmarks (on left labyrinths or reflected right labyrinths). A midline skeleton of each canal was generated using the 'autoskeleton' function of Avizo 9 (<https://www.fei.com/software/amira-avizo/>), which represents the mean endolymph flow path through a semicircular canal<sup>37</sup>. The landmarking of midline skeletons captures the relative lengths, orientations and morphology of the canals. Open semilandmark curves started at the intersection of the canal with its ampulla, ending at its intersection with the common crus. These start and end points represent six fixed, single-point landmarks. The posterior ampulla was estimated to be the ventralmost point on the trajectory of the posterior semicircular canal (psc), and the intersection of the lateral semicircular canal (lsc) with the common crus was estimated to be directly ventral to the intersection of the psc with the common crus<sup>38</sup>. In addition, a closed loop of semilandmarks was placed around the inner surface of the anterior semicircular canal (asc) to capture variation in relative canal thickness<sup>39</sup>. This landmarking procedure involved placing arbitrary numbers of points in each semilandmark series. Thus, these series were resampled to equal numbers of points in each specimen using the digit.curves function of the package Geomorph (version 3.2.1)<sup>40</sup> written for R (version 3.6.0)<sup>41</sup>; asc midline skeleton (9 points), psc midline skeleton (8 points), lsc midline skeleton (10 points) and asc inner loop (13 points). Landmark configurations were transformed through a generalized Procrustes superimposition using the gpgen function of Geomorph in R. This procedure removes differences in orientation and position of specimens and separates overall size information (centroid size) from shape information. Semilandmarks were allowed to slide along their curves during superimposition to minimize bending energy difference from the mean shape.

### Principal component analysis

We conducted principal component analyses of our geometric morphometric dataset using the plotTangentSpace() function of the R package Geomorph (version 3.2.1)<sup>40</sup>.

### Phylogenetic analyses

**Maximum parsimony analysis.** The phylogenetic relationships of pterosaurs and lagerpetids were analysed using the data matrix of

In addition, the formulation or wording was modified or additional states were added for 76 characters, and several scorings were changed with respect to previous versions of the matrix (Supplementary Information). Character 119 was excluded before the searches following a previously published study<sup>43</sup>. Because they represent nested sets of homologies, the following characters were considered additive: 1, 2, 7, 10, 17, 19–21, 28, 29, 36, 40, 42, 46, 50, 54, 66, 71, 74–76, 122, 127, 146, 153, 156, 157, 171, 176, 177, 187, 202, 221, 227, 263, 266, 278, 279, 283, 324, 327, 331, 337, 345, 351, 352, 354, 361, 365, 370, 377, 379, 386, 387, 398, 410, 414, 424, 430, 435, 446, 448, 454, 455, 458, 460, 463, 470, 472, 478, 482, 483, 485, 489, 490, 502, 504, 510, 516, 521, 529, 537, 546, 552, 556, 557, 567, 569, 571, 574, 581, 582, 588, 636, 648, 652, 662, 701, 731, 735, 737, 738, 743, 749, 766, 784 and 816. Several terminal taxa were also excluded because they were originally scored only with the purpose of conducting morphological disparity analyses and were not intended to be included, yet, in phylogenetic analyses<sup>44</sup>. The final data matrix is composed of 822 active characters and 157 active taxa. A second analysis was conducted including *S. taylori*, resulting in a total of 158 taxa. Interpretation of detailed anatomical features is extremely difficult for this species, owing to the preservation of all its specimens as natural moulds of very small-sized bones in a coarse sandstone<sup>4</sup>. Thus, we decided to not include this taxon in the first analysis, because this could introduce a substantial amount of scoring errors.

The data matrix was analysed under equally weighted maximum parsimony using TNT 1.5<sup>26</sup>. The search strategies started using a combination of the tree-search algorithms Wagner trees, tree bisection and reconnection (TBR) branch swapping, sectorial searches, Ratchet and tree fusing, until 100 hits of the same minimum tree length were achieved. The best trees obtained were subjected to a final round of TBR branch swapping. Zero-length branches in any of the recovered most-parsimonious trees were collapsed. Branch support was quantified using Bremer support values and a bootstrap resampling analysis, using 1,000 technical pseudo-replicates and reporting both absolute and GC (group present/contradicted) frequencies. The minimum number of additional steps necessary to generate alternative, suboptimal topologies was calculated when constraining the position of pterosaurs and lagerpetids in different parts of the tree or constraining the topology of pterosaur interrelationships found by previous studies and rerunning the analyses.

An alternative analysis was conducted using the three-dimensional morphogeometric configuration of the endosseous labyrinth of the inner ear. The aligned (Procrustes) coordinates were exported to TNT 1.5 and they were used as a single morphogeometric continuous character<sup>26</sup>. The configurations were realigned in TNT by applying the minimum distances criterion<sup>26</sup> and using *Trilophosaurus buettneri* as the reference taxon because it has been recovered as the earliest branching terminal taxon, among those with three-dimensionally sampled endosseous labyrinths, in previous analyses of this dataset<sup>7,42,43</sup>. As a secondary analysis, we used another early branching archosauromorph, *Mesosuchus browni*, as the reference taxon to test changes in the topologies. The discrete characters of the inner ear (characters 729–743) were excluded during the searches using the three-dimensional morphogeometric character because of their non-independence. The search strategy started using 10 technical replicates of Wagner trees followed by the TBR branch-swapping algorithm (holding 10 trees per replication). The best trees obtained were subjected to a final round of TBR branch swapping. Zero-length branches and additive characters

appearance dates for all tips of the tree. We implemented a node age calibration for Archosauria with a uniform prior of 249.2–257.3 million years ago, in which the minimum is informed by the ages of the oldest archosaurian specimens<sup>44</sup>, and the maximum by age estimates for the crocodile–lizard split<sup>46</sup>. We specified a fossilized birth–death process as the tree model using standard parameterizations and values. Fossils were specified to be tips. The deepest split within the tree was parameterized with a uniform (303.4–318.0 million years ago) tree age prior, for which the maximum is based on the age of the Joggins Formation, which documents the earliest crown-amniotes<sup>47</sup> and the minimum is based on the chronostratigraphic uncertainty of *P. kansensis*, the outgroup and oldest taxon of our sample. We used metropolis-coupling Markov chain Monte Carlo algorithms with two independent runs of four chains, using a heating coefficient of 0.05 and 3 swap attempts per generation. Topological convergence, indicated by average standard deviation of split frequencies decreasing below 0.01, was achieved after 23,496,000 generations. Potential scale reduction factors of 1.0, visual inspection of trace plot with Tracer (version 1.7.1)<sup>48</sup> and estimated sample sizes (ESS) for all parameters >200 further indicated convergence. An additional analysis was performed with the same settings, except that the topology was constrained to investigate topological effects on evolutionary rates. The topology is constrained to follow the topology of one of the most-parsimonious trees of the maximum-parsimony analysis after constraining lagerpetids to be dinosauromorphs. This analysis was specified to run for 24,000,000 generations and estimated sample sizes >200 indicated convergence. Additional phylogenetic analysis details are provided in the Supplementary Information.

## Reporting summary

Further information on research design is available in the Nature Research Reporting Summary linked to this paper.

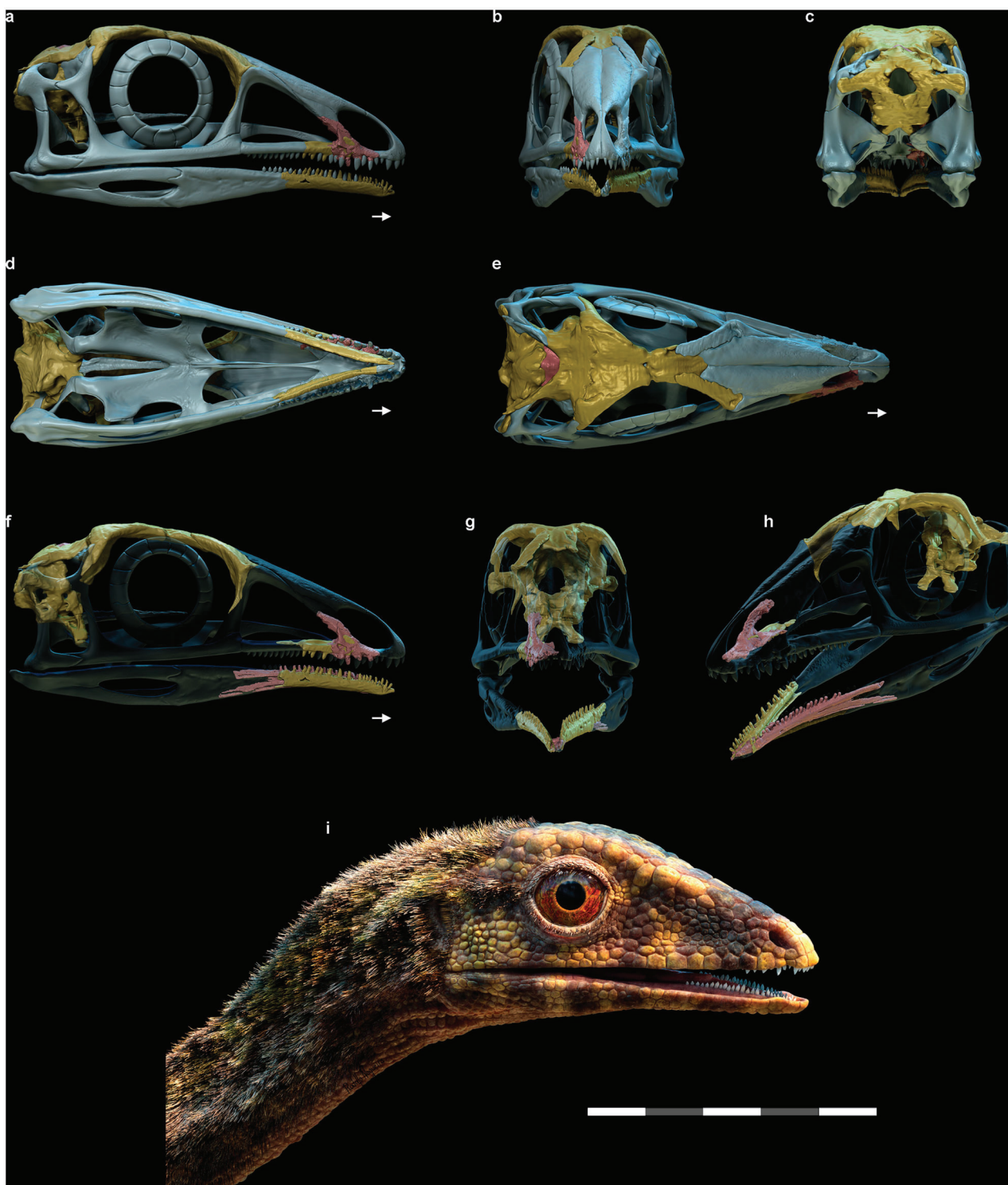
## Data availability

The data matrices for the phylogenetic analyses in NEXUS and/or TNT formats have been deposited in MorphoBank at <http://morphobank.org/permalink/?P3773>. Three-dimensional models of lagerpetid bones in STL format are available in MorphoSource at [http://www.morpho-source.org/Detail/ProjectDetail/Show/project\\_id/1095](http://www.morpho-source.org/Detail/ProjectDetail/Show/project_id/1095) under the following DOIs: <https://doi.org/10.17602/M2/M157269>, <https://doi.org/10.17602/M2/M157271>, <https://doi.org/10.17602/M2/M157273>, <https://doi.org/10.17602/M2/M157275>, <https://doi.org/10.17602/M2/M157280>, <https://doi.org/10.17602/M2/M157282>, <https://doi.org/10.17602/M2/M157283> and <https://doi.org/10.17602/M2/M157284>. Source data are provided with this paper.

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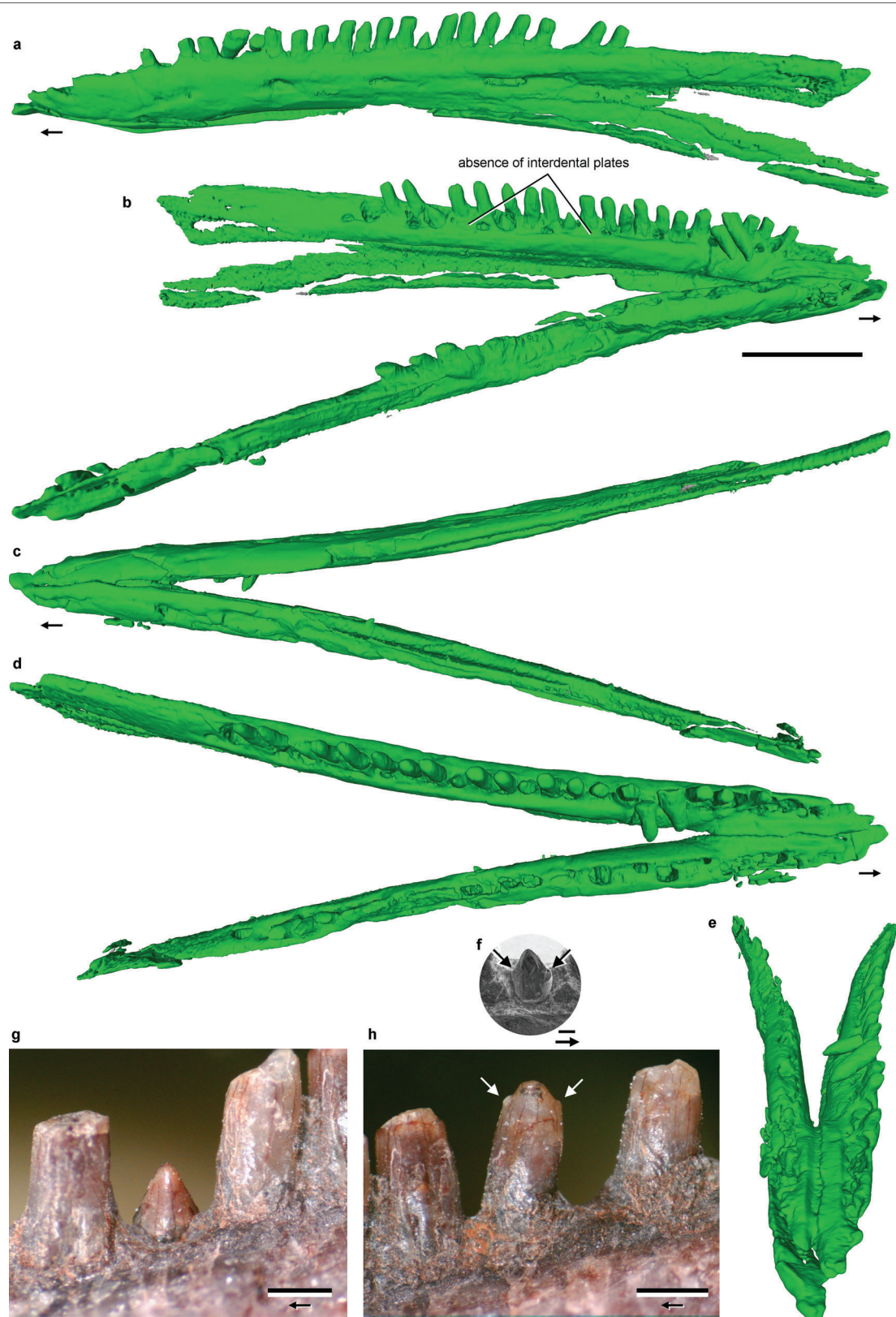




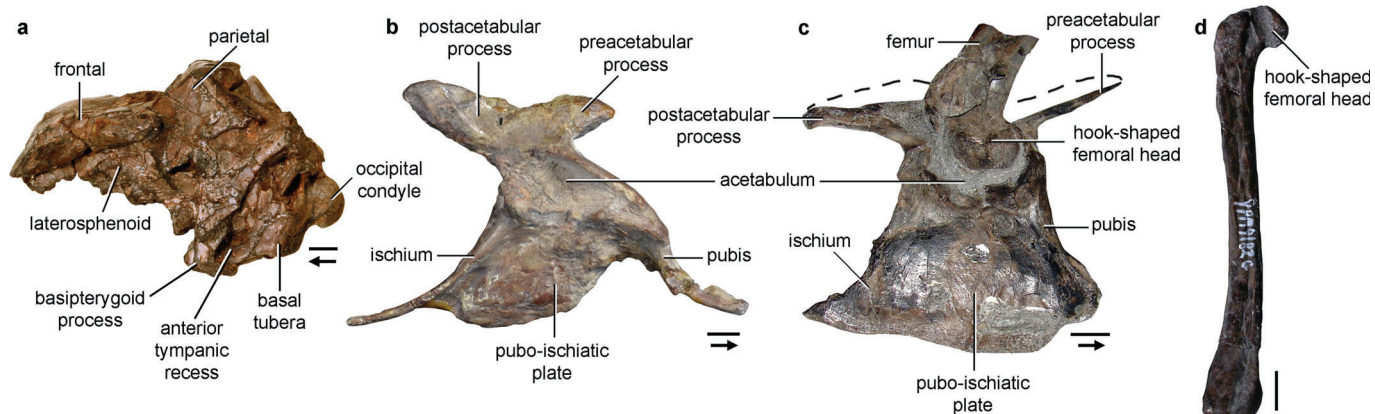


**Extended Data Fig. 1 | Life reconstruction and three-dimensional reconstruction of the skull of the lagerpetid *I. polesinensis* (ULBRA-PVT059, holotype) with additions of cranial bones of other lagerpetids. a–i, Images are shown in right lateral (a, f, i), anterior (b, g), posterior (c), ventral (d), dorsal (e) and anterodorsolateral (h) views.**

roof bones through them. Bones of *I. polesinensis* are indicated in yellow and those of *K. kely* (UA10618, holotype), *D. gregorii* (TMM31100-1334, referred specimen) and *L. chanarensis* (PVL4625, referred specimen) in red, and inferred bones in light blue. Arrows indicate the anterior direction. Complete scale bar, 5 cm. Life and skull reconstruction by R. Nogueira.

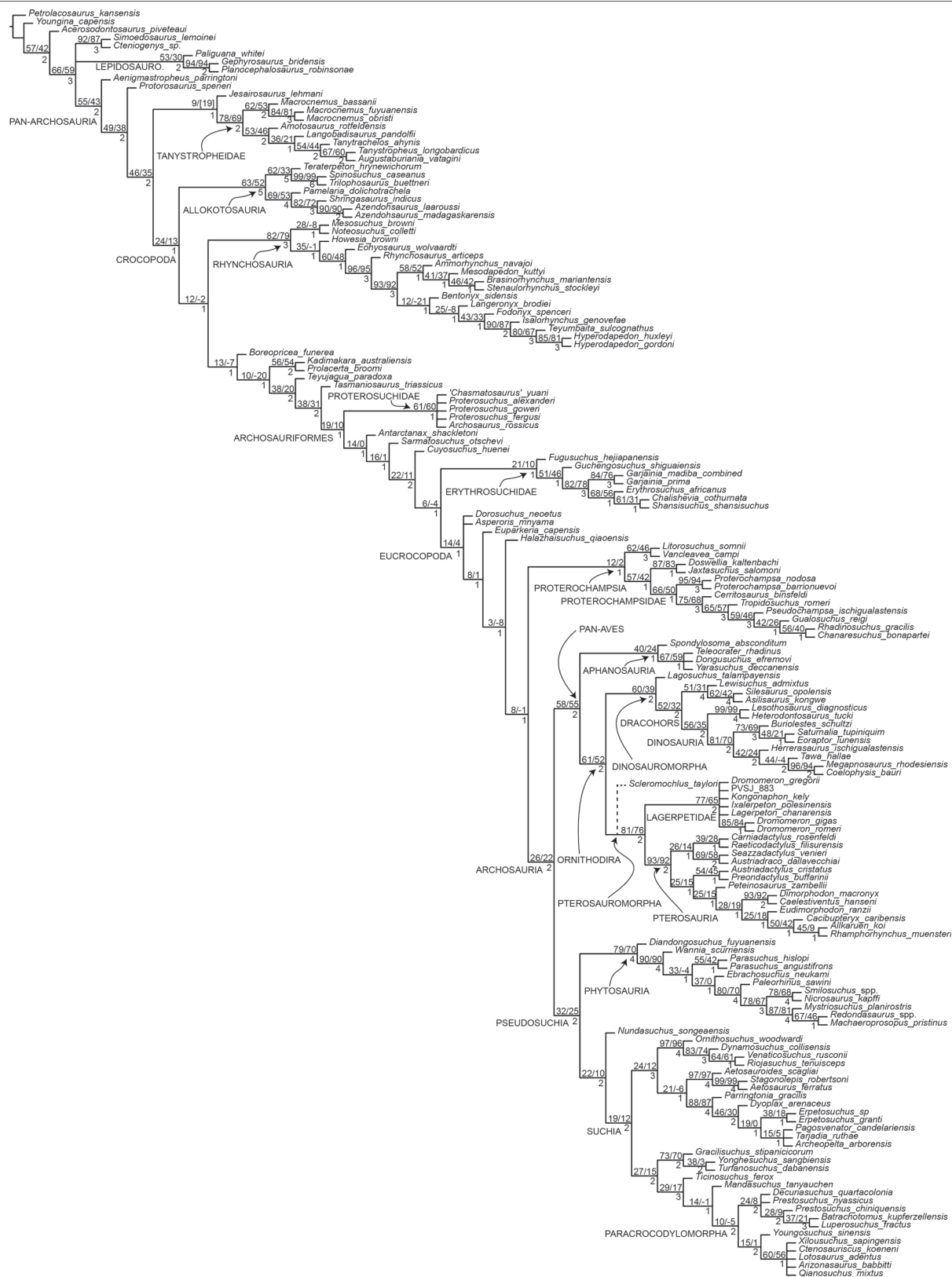


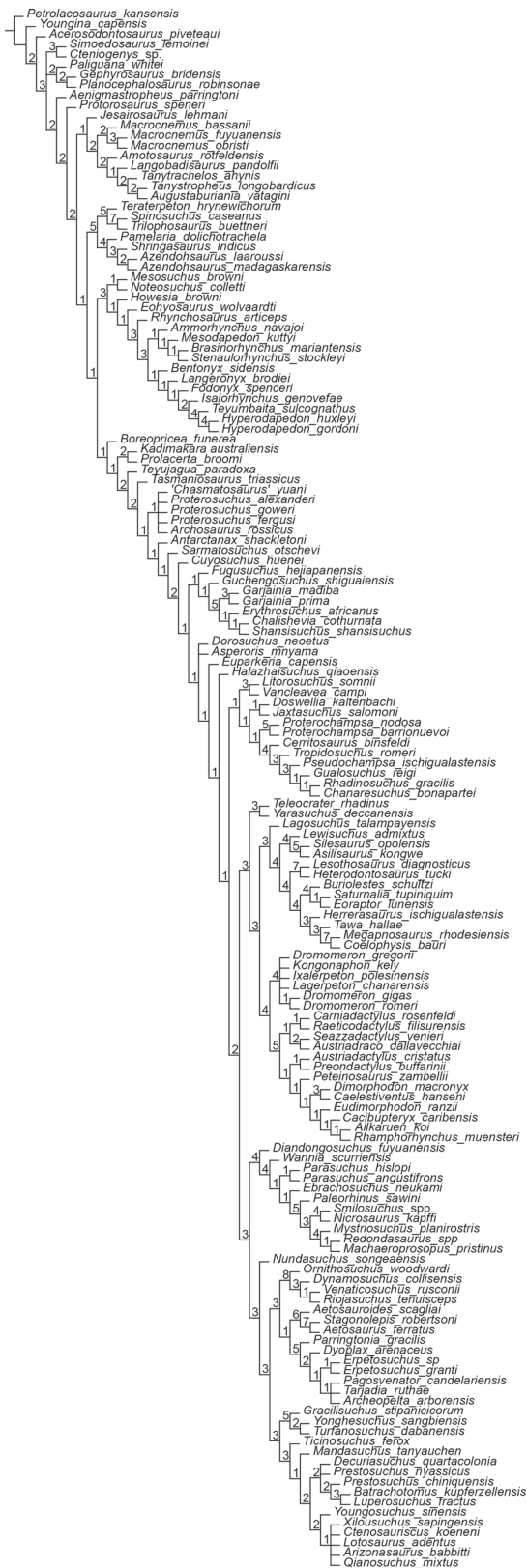




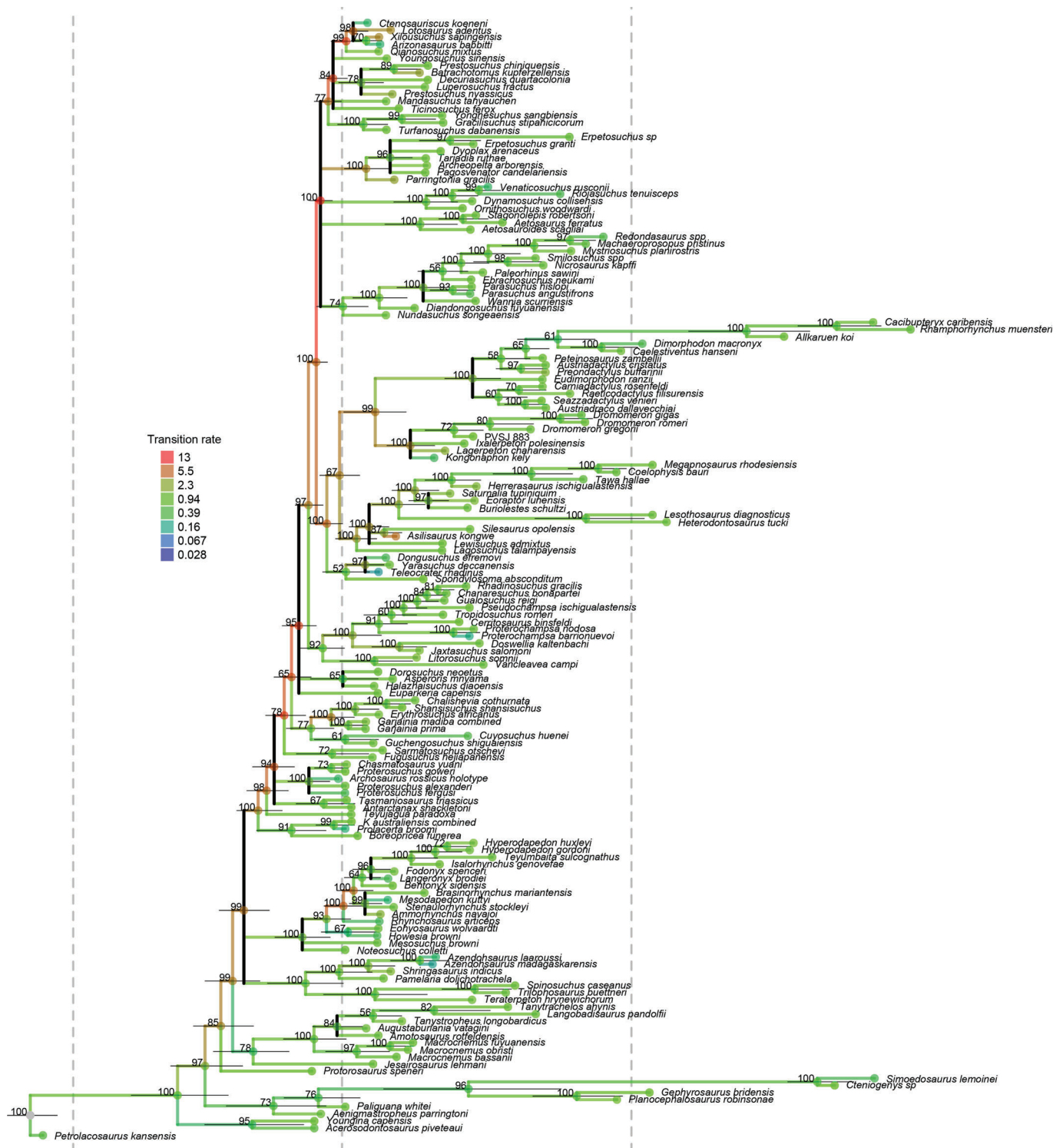
**Extended Data Fig. 3 | Additional images and comparisons between lagerpetids and pterosaurs.** **a**, Partial skull roof and braincase of the lagerpetid *D. gregorii* (TMM 31100-1334) in left lateral view. **b**, Right hemipelvis of the lagerpetid *L. chanarensis* (PVL 4619) in lateral view. **c**, Left hemipelvis and

articulated proximal end of femur of the pterosaur *Dimorphodon macronyx* (NHMUK PV OR 41212, reversed) in lateral view. **d**, Right femur of the pterosaur *D. macronyx* (YPM 9182) in anterolateral view. Arrows indicate the anterior direction. Scale bars, 3 mm (**a**) and 5 mm (**b-d**).









**Extended Data Fig. 6 | Majority rule tree recovered from the unconstrained Bayesian phylogenetic analysis.** Branch colours indicate character state transition rates (that is, the evolutionary rates), numbers at the nodes indicate posterior probabilities, the thin black horizontal line segments indicate the

95% probability distribution of node ages, and dotted vertical lines indicate the boundaries between the Carboniferous, Permian, Triassic and Jurassic geological periods. Thick black vertical bars indicate polytomies and, as a result, transition rates could not be calculated.

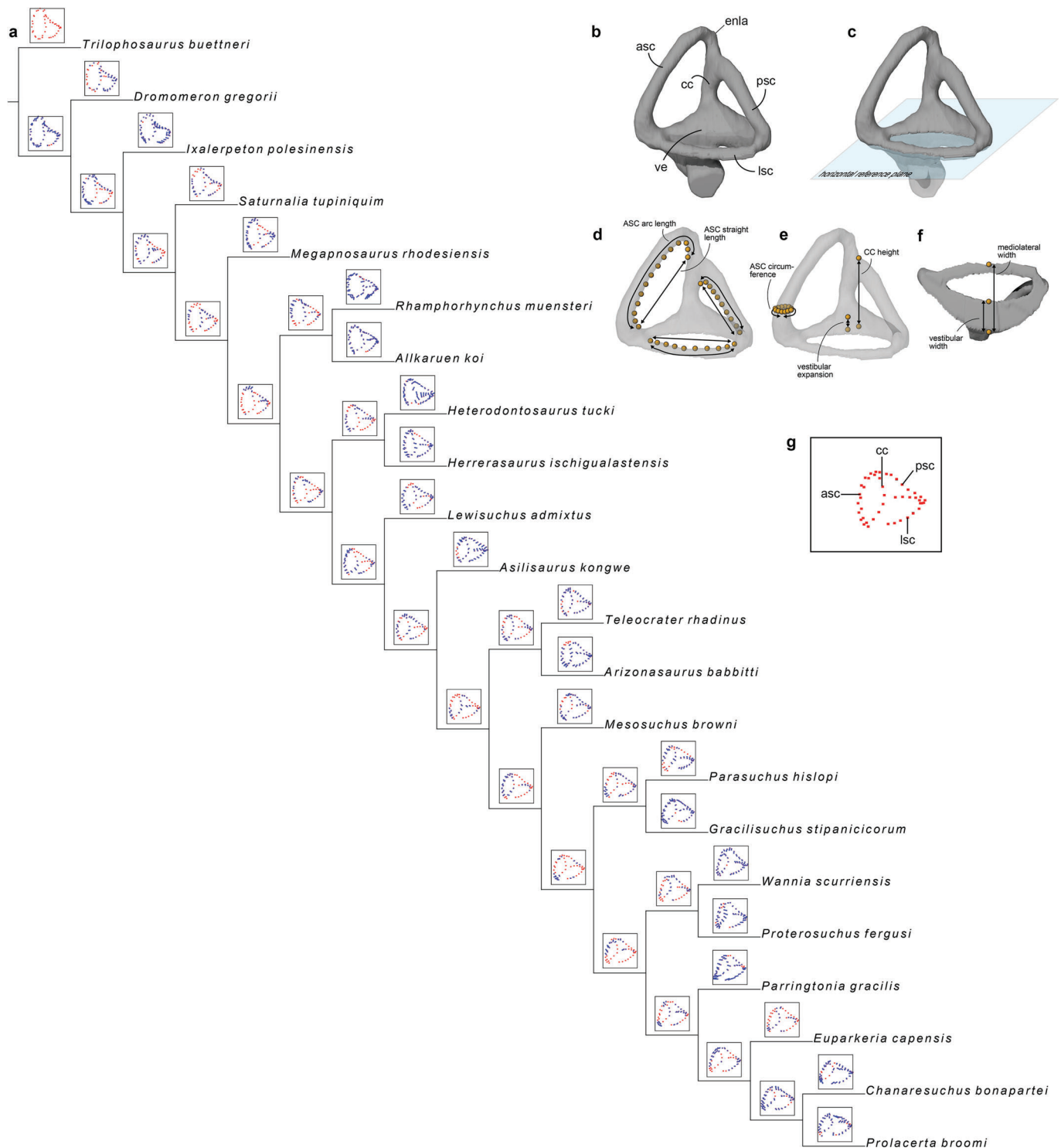


**Extended Data Fig. 7 | Majority rule tree recovered from the constrained Bayesian phylogenetic analysis.** The topology of this tree has been constrained a priori after selecting randomly one of the most-parsimonious trees recovered after forcing the position of lagerpetids as the earliest branching dinosauriforms in the maximum parsimony analysis. Branch

colours indicate character state transition rates, the black horizontal line segments indicate the 95% probability distribution of node ages and the dotted vertical lines indicate the boundaries between the Carboniferous, Permian, Triassic and Jurassic geological periods. Posterior probabilities at the nodes are not shown because the topology is fully constrained.



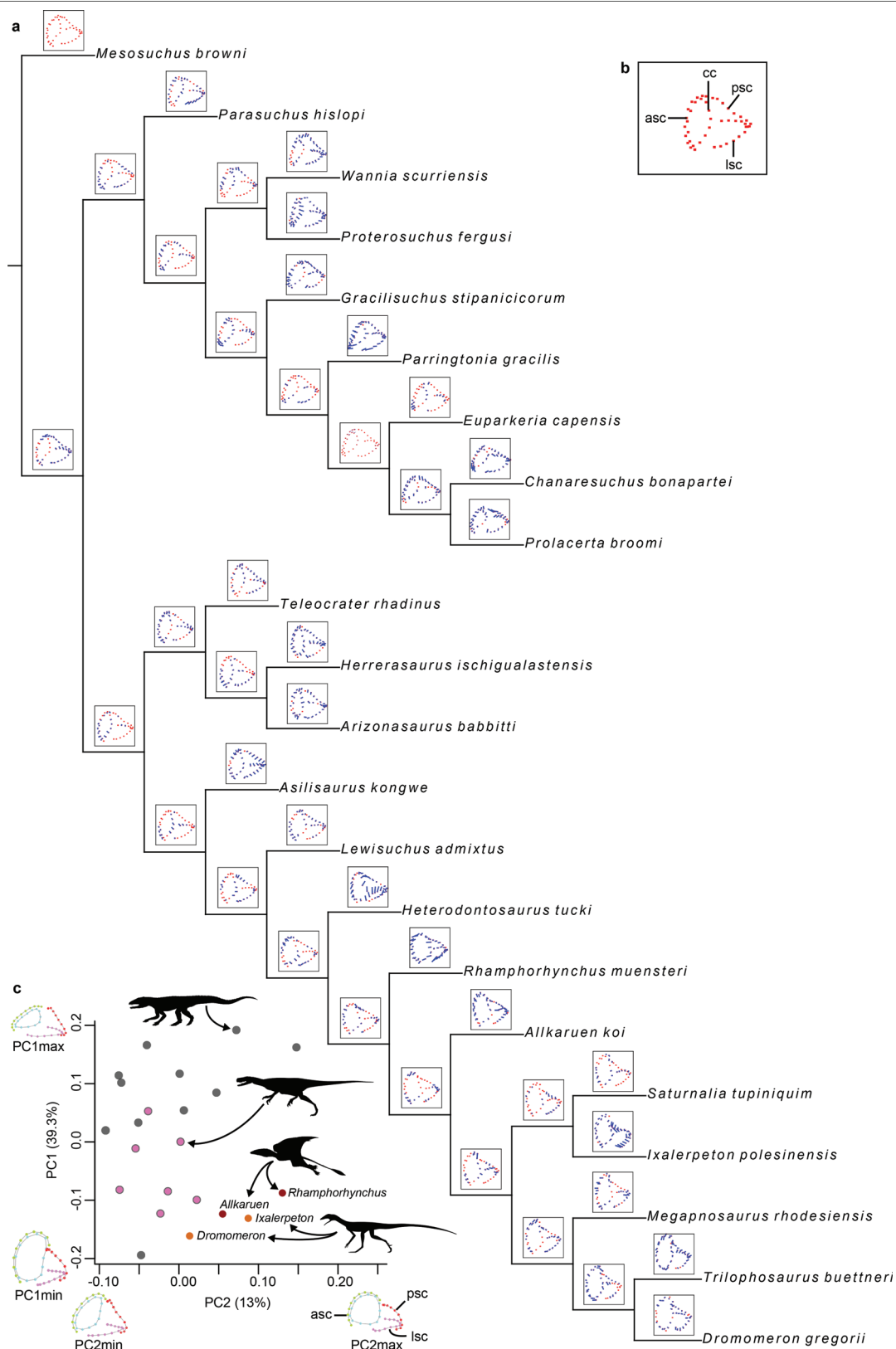




**Extended Data Fig. 9 | Single most-parsimonious tree found when analysing only the three-dimensional morphogeometric configuration of the inner ear and three-dimensional examples of how distances and angles were measured in the three-dimensional endosseous labyrinth models.**

**a**, Tree rooted with *T. buettneri*. **b**, Three-dimensional model of the left endosseous labyrinth of *Platanosaurus* (UMNH 91927) in dorsolateral view.

scheme for semicircular canal length measurements. **e**, Landmark scheme for asc circumference and labyrinth height measurements. **f**, Landmark scheme for labyrinth width measurements in ventral view on the reference plane. **g**, Landmark constellation explanation. asc, anterior semicircular canal; cc, common crus; enla, endosseous labyrinth; lsc, lateral semicircular canal; psc, posterior semicircular canal; ve, vestibule.



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### Software and code

For information about [availability of computer code](#)

For data collection: 3D Slicer version 4.7, Amira version 5.3.3; Avizo lite 9.2 (autokeleton function); Geomorph version 3.2.1 package for R.

For data analysis: TNT version 1.5; MrBayes 3.2.6; Tracer 1.7.1; Geomorph version 3.2.1 package for R.

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A list of figures that have associated raw data

A description of any restrictions on data availability

For fossil specimens used for this study are deposited in public institutions available for qualified researchers upon request. The data matrices for the phylogenetic analyses have been deposited in MorphoBank (<http://morphobank.org/permalink/?P3773>) and the mCT-scan based 3D models of the lagerpetid cranial bones have been deposited in MorphoSource ([http://www.morphosource.org/Detail/ProjectDetail/Show/project\\_id/1095](http://www.morphosource.org/Detail/ProjectDetail/Show/project_id/1095)).



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## Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size	Our study considers all available specimens of the studied species and the phylogenetic analyses include the so far most comprehensive sample of Permo-Triassic archosauromorph species. Terminals of the phylogenetic data matrices were scored based on the type specimens and all available, unambiguously referred specimens.
Taxa exclusions	The taxonomic and character samples of the phylogenetic analyses were intended to be as broad as possible, without exclusions.
Publication	Our taxon-character matrix is the latest modification of that originally published by Ezcurra (2016 – PeerJ), all modifications are described in detail in the supplementary information. This documentation and accessibility of the data matrices allow the replication of the analyses. The matrix of Ezcurra (2016) has been reanalyzed independently several times (>10 analyses) in the last four years and all replication attempts were successful (with exception of minor changes as a result of slight modifications and/or expansions of the dataset).
Randomization	Our study has used all unambiguously referred specimens and in most cases not more than five individuals are available. Thus, no randomizations were required.
Blinding	The taxon-character matrices for the phylogenetic analyses were built on the basis of independent observation of each taxon. Blinding was not necessary, neither possible, because we have used all available specimens for the scoring of characters in each sampled species.

## Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

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## Paleontology and Archaeology

Specimen provenance	The most relevant specimens for our study have been collected during previous research efforts and fieldworks in the Upper Triassic Chañares Formation, Ischigualasto-Villa Unión Basin, Argentina (Lagerpeton chanarensis); Upper Triassic Santa Maria Sequence, Paraná Basin, Brazil (Ixalerpeton polesinensis); Upper Triassic Chinle Formation and Dockum Group, USA (Dromomeron romeri and Dromomeron gregorii), and Middle-Upper Triassic Isalo II beds, Madagascar (Kongonaphon kelyi). No field or other official permits have been required for the work.
Specimen deposition	The most relevant specimens for our study are deposited in Paleontología de Vertebrados, Instituto Miguel Lillo, San Salvador de Tucumán, Argentina (Lagerpeton chanarensis); Museu de Ciências Naturais, Universidade Luterana do Brasil, Canoas, Brazil (Ixalerpeton polesinensis); Ruth Hall Museum of Paleontology, Ghost Ranch, Abiquiu, New Mexico, USA (Dromomeron romeri); Texas Vertebrate Paleontology Collections, The University of Texas at Austin, Austin, Texas (Dromomeron gregorii); and University of Antananarivo, Antananarivo, Madagascar (Kongonaphon kelyi).
Dating methods	No new dates were obtained for this study. Dates used for the Bayesian inference analysis are provided in the supplementary file of

that full information on the approval of the study protocol must also be provided in the manuscript.