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## Endochondral bone in an Early Devonian 'placoderm' from Mongolia

Brazeau, Martin D.; Giles, Sam; Dearden, Richard P.; Jerve, Anna; Ariunchimeg, Ya.; Zorig, E.; Sansom, Robert; Guillerme, Thomas; Castiello, Marco

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1	Endochondral bone in an Early Devonian 'placoderm' from Mongolia
2	
3	Martin D. Brazeau <sup>1,2</sup> *, Sam Giles <sup>2,3,4</sup> , Richard P. Dearden <sup>1,5</sup> , Anna Jerve <sup>1,6</sup> , Y.A.
4	Ariunchimeg <sup>7</sup> , E. Zorig <sup>8</sup> , Robert Sansom <sup>9</sup> , Thomas Guillerme <sup>10</sup> , Marco Castiello <sup>1</sup>
5	
6	<sup>1</sup> Department of Life Sciences, Imperial College London, Silwood Park Campus, Buckhurst
7	Rd, Ascot, SL5 7PY, UK;
8	<sup>2</sup> Department of Earth Sciences, Natural History Museum, Cromwell Road, London SW7
9	5BD, UK;
10	<sup>3</sup> School of Geography, Earth and Environmental Sciences, University of Birmingham,
11	Birmingham, UK;
12	<sup>4</sup> Department of Earth Sciences, University of Oxford, South Parks Road, Oxford, OX1 3AN,
13	UK;
14	<sup>5</sup> CR2P Centre de Recherche en Paléontologie – Paris, Muséum national d'Histoire
15	naturelle, Sorbonne Universités, CNRS, CP 38, 57 Rue Cuvier, 75231, Paris, Cedex 05,
16	France
17	<sup>6</sup> Department of Organismal Biology, Subdepartment of Evolution and Development,
18	Uppsala University, Norbyvägen 18A, 752 36 Uppsala, Sweden;
19	<sup>7</sup> Natural History Museum, P.O. Box 46/52, Ulaanbaatar 1420, Mongolia
20	<sup>8</sup> Institute of Paleontology, Mongolian Academy of Science, P.O. Box 46/650, S. Danzan
21	Street 3/1, Chingeltei District. Ulaanbaatar 15160, Mongolia;
22	<sup>9</sup> School of Earth and Environmental Sciences, University of Manchester, Manchester M13
23	9PT, UK;
24	<sup>10</sup> Department of Animal and Plant Sciences, The University of Sheffield, Sheffield S10 2TN,
25	UK;

27 \* Author for correspondence.

28 Endochondral bone is the main internal skeletal tissue of nearly all osteichthyans-the 29 group comprising more than 60,000 living species of bony fishes and tetrapods. 30 Chondrichthyans (sharks and their kin) are the living sister group of osteichthyans and 31 have cartilaginous endoskeletons, long considered the ancestral condition for all jawed 32 vertebrates (gnathostomes). The absence of bone in modern jawless fishes and the 33 absence of endochondral ossification in early fossil gnathostomes appears to lend 34 support to this conclusion. Here we report the discovery of extensive endochondral bone 35 in *Minjinia turgenensis*, a new genus and species of 'placoderm'-like fish from the Early 36 Devonian (Pragian) of western Mongolia described using x-ray computed 37 microtomography (XR-µCT). The fossil consists of a partial skull roof and braincase 38 with anatomical details providing strong evidence of placement in the gnathostome stem 39 group. However, its endochondral space is filled with an extensive network of fine 40 trabeculae resembling the endochondral bone of osteichthyans. Phylogenetic analyses 41 place this new taxon as a proximate sister group of the gnathostome crown. These 42 results provide direct support for theories of generalised bone loss in chondrichthyans. 43 Furthermore, they revive theories of a phylogenetically deeper origin of endochondral 44 bone and its absence in chondrichthyans as a secondary condition.

45

The vertebrate skeleton comprises two main systems: the exoskeleton (external achondral dermal bones) and endoskeleton (internal chondral bones and cartilages, as well as some intramembranous bones)<sup>1</sup>. An ossified exoskeleton evolved at least 450 million years ago in jawless stem gnathostomes<sup>2,3</sup>, but the endoskeleton in these taxa is not endochondrally ossified. Endochondral bone, in which the cartilaginous endoskeletal precursor is invaded by and eventually replaced by bone, is widely considered an osteichthyan apomorphy<sup>3-7</sup> and such a reliable identifying character gives the group its name. Extant chondrichthyans lack dermal bone and possess a mainly cartilaginous endoskeleton enveloped by a structurally diverse range of tessellate calcified cartilage<sup>8</sup>. Outgroups of the gnathostome crown also lack endochondral ossification. Galeaspids surround their cartilaginous skeleton in globular calcified cartilage {NianZhong:2005tj}, while osteostracan and 'placoderm' endoskeletons were sheathed in perichondral bone<sup>3</sup>. Consequently, the last common ancestor of jawed vertebrates was long thought to have been perichondrally ossified, but lacking endochondral ossification<sup>3</sup>.

60 In this paper, we describe a new genus and species of 'placoderm' from the Early 61 Devonian of western Mongolia. Although Mongolia is known for some of the geologically oldest putative gnathostome fossils (isolated chondrichthyan-like scales <sup>9-12</sup>), it remains a 62 poorly sampled region of the world with respect to early vertebrates. 'Placoderms' were until 63 64 now known from only a single fragmentary occurrence<sup>13</sup> in the early Middle Devonian 65 (Eifelian). Our new data highlight the importance of Mongolia as a key region for studies of 66 early gnathostome evolution. We describe a braincase and partial skull roof representing the 67 first substantial body fossil of an early gnathostome from Mongolia and displaying an 68 unexpected occurrence of endochondral bone analysed using XR-µCT. We conducted 69 phylogenetic analyses to reconstruct the evolutionary relationships of this new taxon. To 70 explore the evolutionary history of endochondral bone in light of this new discovery, we used 71 parsimony and maximum likelihood ancestral states reconstruction. Finally, we discuss these 72 results in the context of earlier statements about endochondral bone in non-osteichthyans, 73 new developments in understanding the complexity and diversity of chondrichthyan 74 endoskeletal tissues, and current uncertainties about early gnathostome phylogenetic 75 relationships.

76

## 77 Systematic palaeontology

78	Gnathostomata Gegenbaur, 1874 <sup>14</sup>
79	Minjinia turgenensis gen. et sp. nov.
80	
81	Etymology. Generic name honours the memory of Chuluun Minjin for his extensive
82	contributions to the Palaeozoic stratigraphy of Mongolia, his enthusiastic support of this
83	work, and introducing us to the Yamaat River locality. Specific name recognises the
84	provenance of the fossil from the Turgen region, Uvs aimag of western Mongolia.
85	
86	Holotype. Institute of Paleontology, Mongolian Academy of Sciences MPC-FH100/9.1, a
87	partial braincase and skull roof.
88	
89	Type locality. Turgen Strictly Protected Area, Uvs province, western Mongolia; near the top
90	of the stratigraphic sequence that occurs between the Tsagaan-Salaat and Yamaat Rivers.
91	
92	Formation and age. Upper part of Tsagaansalaat Formation, Pragian (Early Devonian) <sup>15,16</sup> .
93	
94	Diagnosis. 'Placoderm'-grade stem gnathostome with endochondral bone, deep epaxial
95	muscle cavities flanking a slender occipital ridge, and the following possible autapomorphies:
96	dermal bones covered in sparsely placed tubercles, penultimate spino-occipital nerve canal
97	substantially larger in diameter than others.
98	
99	Description
100	MPC-FH100/9.1 consists of a partial braincase and skull roof (Fig. 1). The skull roof is
101	ornamented with sparsely distributed finely ridged tubercles resembling those of the Siberian
102	'placoderm' <i>Dolganosteus</i> <sup>17</sup> ; the tubercles become more broadly separated towards the

103 midline of the skull. They are distinct from those of *Dolganosteus* in that towards the midline 104 of the skull roof, the tubercles are larger and more pointed. The specimen shows signs of 105 extensive post-mortem transport, with angles of the braincase worn off and much of the skull 106 roof and some of the braincase preserved as a mould. Individual skull roof ossifications 107 cannot be identified, although this may be due to the dominantly mouldic preservation. There 108 appears to have been a prominent nuchal plate eminence comparable to certain 109 acanthothoracids such as *Romundina*<sup>18</sup> and *Arabosteus*<sup>19</sup>.

110

111 Endoskeletal tissue. The braincase of MPC-FH100/9.1 is well ossified, comprising an 112 external bony sheath filled with an extensive matrix of spongy tissue (Fig. 2a-b; Extended 113 Data Fig. 1; Supplementary Video 1). The trabecles forming this tissue are irregular and 114 branching, less than 1 mm thick and often curved, and resemble most closely the endochondral tissue of osteichthyans (Fig. 2c-d; Supplementary Video 2). As such, we 115 116 interpret this as endochondral bone. Notably, this is found in all preserved regions of the 117 braincase, in contrast to the isolated trabeculae previously identified as endochondral bone in Boreaspis<sup>20</sup> and Bothriolepis<sup>21</sup>. The margins of the braincase, the endocranial walls, and the 118 boundaries of nerve and blood canals, are formed from a thicker tissue which we interpret as 119 120 perichondral bone. This suggests that the endoskeleton of Minjinia comprises osteichthyanlike endochondral bone, with an ossified perichondrium. To address the possible alternative 121 122 explanation that it is an aberrant instance of calcified cartilage, we compared the structure of this tissue with rarely-preserved mineralized cartilage in the stem chondrichthyan 123 Diplacanthus crassismus (National Museums of Scotland specimen NMS 1891.92.334; Fig. 124 125 2e-f) observed using synchrotron tomography. The cancellae within the endochondral tissue of Minjinia are irregular, with a diameter of approximately 1-2 mm. This tissue is distinctly 126

unlike the calcified cartilage of *Diplacanthus* in appearance, which consists of a densely
packed matrix of irregularly stacked chondrons between 20-60 µm in diameter.

129

130 **Neurocranium.** The braincase is preserved from the level of the right posterior orbital wall 131 to the posterior end of the occipital ridge. Occipital glenoid condyles are not preserved, but 132 much of the rest of the broad, flat parachordal region is present, separated by a midline 133 groove that accommodated a relatively narrow notochordal tunnel. An asymmetric transverse 134 fissure spans the basic anial surface at about mid-length of the preserved portion. It appears 135 to demarcate the anterior margin of the parachordal plates and may correspond to the ventral 136 cranial fissure of crown-group gnathostomes. However, unlike in crown gnathostomes, it is 137 traversed by a substantial anterior extension of the cranial notochord. The courses of the 138 lateral dorsal aortae are marked by a pair of sulci on the lateral margins of the parachordal plates, though only a short part of the canal is preserved on the right side of the specimen. A 139 140 narrow, shallow sulcus for the efferent hyoid artery is present on the preserved right side of 141 the specimen, immediately behind the level of the orbit (Fig. 1a). 142 The lateral surface of the braincase is preserved on the right side as a mouldic

impression in the matrix (Fig. 1). A sharply demarcated hyoid fossa is present on the lateral wall of the otic region (Fig. 1). Posterior to this, a stout but pronounced vagal process with a pair of rounded eminences likely corresponds to the branchial arch articulations. There is no evidence for a pair of anterior and posterior divisions to the vagal process, which are typically seen in other 'placoderms'. A well-developed 'placoderm'-like craniospinal process is absent; its homologous position is instead covered in perichondral bone and marked by a low ridge (Fig. 1).

In posterior view, a tall, narrow median occipital ridge is evident and resembles the
 morphology of *Romundina*<sup>22</sup> and *Arabosteus*<sup>19</sup>. Similar to these taxa, the median otic ridge is

152 flanked by two large occipital fossae for the epaxial musculature. The notochordal tunnel is 153 approximately the same size as or smaller than the foramen magnum, as in 'placoderms' and 154 in contrast with crown-group gnathostomes. A metotic fissure is absent.

155

156 Endocast. A partial cranial endocast is preserved, consisting of the hindbrain cavity, partial 157 midbrain cavity, labyrinth cavities, and posteromedial corner of the orbital region. The two 158 primary trunk canals of the trigeminal nerve (N.V<sub>1</sub> and N.V<sub>2,3</sub>) are preserved (Fig. 3). The 159 acoustic (N.VIII) and facial nerve (N.VII) canals share a common trunk canal behind the 160 trigeminal nerves, as in many other 'placoderms' <sup>22-25</sup>. The facial nerve canal branches into 161 palatal and hyomandibular branches between the saccular chamber and rear orbit wall (Fig. 3; Extended Data Fig. 2), indicating this division was internal (deep) to the otic process. The 162 163 supraophthalmic branch opens into the rear wall of the orbit and part of its supraorbital 164 course is preserved (Fig. 3; Extended Data Fig. 2). A slender branch extends below the 165 labyrinth and divides into palatine and hyomandibular branches (Fig 3; Extended Data Fig. 166 2). As in other 'placoderm'-grade taxa, the vagus nerve (N. X) trunk canal is very large in 167 diameter and exits from immediately behind the labyrinth cavity (Fig. 3). The spino-occipital 168 region resembles other 'placoderms' in being extended. At least four spino-occipital nerve 169 canals are present in a linear series, and the penultimate canal is largest in diameter (Fig. 3). 170 Intercalating these is a network of occipital artery canals branching from the dorsal aortae. 171 The skeletal labyrinth is not complete on either side of the specimen, but can mostly 172 be reconstructed according to the assumption of bilateral symmetry. The most significant 173 feature is that the labyrinth and endolymphatic cavity are joined to the main endocavity 174 chamber (Fig. 3). This is a striking contrast to other 'placoderms' and closely resembles crown-group gnathostomes<sup>26</sup>. The endolymphatic canals are elongate and tubular, extending 175 176 posterolaterally to reach the skull roof, though external openings cannot be clearly identified.

The anterior semi-circular canal follows the saccular cavity closely as in petalichthyids<sup>27</sup>(Fig.
3). However, the horizontal and posterior canals appear to extend well away from the

179 saccular chamber (Fig. 3). The dorsal junctions of the anterior and posterior canals are joined

180 in a crus commune, as in *Romundina*<sup>22</sup> and *Jagorina*<sup>23</sup>. A sinus superior is absent.

181

### 182 Phylogenetic analyses

183 We conducted phylogenetic analyses under four different protocols: equal weights 184 parsimony, implied weights parsimony, an unpartitioned Bayesian analysis, and a Bayesian analysis with characters partitioned by fit determined under implied weights parsimony<sup>28</sup> (see 185 186 Extended Data Figs. 3-6). All phylogenetic analyses consistently place Minjinia as a stem-187 group gnathostome, proximate to the gnathostome crown (Fig. 4, Extended Data Figs 3, 4). 188 Minjinia is recovered in a position crownward of arthrodires but outside of a grade consisting 189 of Entelognathus, Ramirosuarezia, and Janusiscus. Under implied weights parsimony, these 190 three taxa move onto the osteichthyan stem and Minjinia is placed as the immediate sister 191 taxon of the gnathostome crown. Under parsimony, the crownward position of *Minjinia* is 192 unambiguously supported by the skeletal labyrinth and endolymphatic duct being confluent with the main cranial cavity<sup>26</sup> (Supplementary Information). In common with arthrodires and 193 the gnathostome crown<sup>29</sup>, *Minjinia* possesses a division of the facial nerve(Fig. 3; Extended 194 195 Data Fig. 2) deep to the transverse otic process. However, Minjinia is excluded from the 196 gnathostome crown group due to the absences of a metotic fissure and a posterior dorsal 197 fontanelle, and presence of broad, flat parachordal plates expanded behind the saccular cavity 198 (Fig. 3, Supplementary Information).

We undertook ancestral states reconstructions to assess the evolutionary history of
endochondral bone (Fig. 4; Extended Data Figs. 5 & 6; Supplementary Information).
Interestingly, parsimony analysis fails to recover secondary homology of this trait between

202 Minjinia and osteichthyans. The crownward placement of Minjinia is, in fact, based on 203 independent evidence relating to anatomical features of the braincase and endocast. However, 204 the resolution becomes ambiguous if missing data in either Entelognathus or Ramirosuarezia 205 are resolved as having endochondral bone. The reconstruction becomes similarly ambiguous 206 if Janusiscus is moved a single branch (requiring only two additional steps) onto the 207 chondrichthyan stem. The strict precision of parsimony reconstructions makes it insensitive 208 to this underlying uncertainty. To explore this, we used likelihood reconstructions and 209 compared the ancestral state reconstructions under equal rates (ER) and all rates different 210 (ARD) variants of the Mkv model on branch-length-rescaled parsimony trees and Bayesian 211 trees. Both models show substantial non-zero marginal likelihoods if endochondral bone is 212 assumed present in the common node of *Minjinia* and Osteichthyes, with ARD strongly 213 favouring its presence (0.33 for ER; 0.81 for ARD; Fig. 3, Table 1, Extended Data Figs. 5, 6, 214 Supplementary Table 1). Under the ARD model, there is nearly equivocal support for 215 presence or absence of endochondral bone at the gnathostome crown node (Table 1). The 216 ARD model shows the best fit for endochondral bone (likelihood ratios 4.75 for parsimony [p 217 = 0.029] and 5.26 for Bayesian, [p = 0.022]) (Table 1, Supplementary Table 1), favouring 218 repeated losses of this tissue over multiple gains (see Discussion).

219

### 220 **Discussion**

*Minjinia turgenensis* presents an unexpected discovery of extensive endochondral bone in a 'placoderm'-grade fish, with repercussions for the phylogenetic origin of this tissue and the problem of early gnathostome relationships more generally. The prevailing hypothesis has been that endochondral bone is an osteichthyan apomorphy<sup>3,7,29</sup>. However, recent discoveries have cast doubt on this assertion. The recognition that dermal bone is secondarily lost in chondrichthyans<sup>30,31</sup> (Fig. 4) is consonant with prior knowledge of the loss of perichondral bone in this same lineage<sup>32</sup>. Taken together, this has revived uncertainty about the true
phylogenetic timing of the origin of endochondral ossification<sup>33</sup>. *Minjinia* provides direct
corroboration for a more ancient origin.

230 Minjinia does not represent the first report of endochondral bone outside of Osteichthyes. However, it is by far the most extensive and unequivocal example and raises 231 explicit questions in light of the proximity of *Minjinia* to the gnathostome crown (Fig. 4; 232 233 Extended Data Figs. 3, 4). Isolated examples of trabecular endoskeletal bone have historically been reported in boreaspid osteostracans<sup>20,34</sup>, a rhenanid<sup>35</sup>, arthrodires<sup>36</sup>, a ptyctodont<sup>37</sup>, and a 234 235 petalichthyid<sup>38,39</sup>. However, these reports are nearly all unillustrated statements; they have all been considered tenuous<sup>3</sup> or dismissed as misidentifications<sup>5</sup>. In line with these assessments, 236 237 we found no evidence of endochondral bone in material of *Buchanosteus* held in the Natural 238 History Museum, London, or indeed in any other 'placoderms' we have examined. The Epipetalichthys holotype (Museum für Naturkunde, Berlin specimen MB.f.132.1-3) shows an 239 apparently spongiose infilling in the anterior region of the braincase, but the identity of this 240 241 structure, or even whether it is biological, cannot be determined. The *Epipetalichthys* tissue figured by Stensiö<sup>38</sup> was very superficial, and possibly represents the retreat of perichondral 242 bone deposited during cartilage growth<sup>39</sup>. Most recently, trabeculae in supposed endoskeletal 243 pelvic bones of *Bothriolepis* have been termed endochondral bone<sup>21</sup>, although the small scale 244 of these is in line with 'superficial' perichondral trabeculae seen elsewhere<sup>38</sup>. The reported 245 246 examples in boreaspid osteostracans have also been dismissed by later authors<sup>3,5</sup>. Although they warrant further study, their tissue structures are unlikely to be homologous to 247 248 osteichthyans owing to their phylogenetic remoteness and nested position in the Osteostraci<sup>40</sup>. 249

Among chondrichthyans, endochondral bone has been mentioned in 'acanthodians'<sup>3,41</sup>
and superficial bone-like tissues have been reported in the skeletons of extant

252 chondrichthyans. We are unable to substantiate statements about acanthodians: no authors have cited primary sources or specimens. One possible source is Watson's<sup>42</sup> description of 253 254 "massive ossification" of the endoskeleton of Diplacanthus. However, our synchrotron data 255 of this same specimen (Fig. 2) shows that this tissue is undoubtably calcified cartilage. Some 256 authors have speculated that the superficial mineralised tissue in the jaws of acanthodians or chondrichthyans may have developed in an endochondral position<sup>39</sup>. Histological studies 257 258 show that endoskeletal mineralization in the jaws of acanthodians is globular calcification and occasionally 'sub-tessellate'<sup>8,43</sup>. Recent comparative histology and development in extant 259 260 chondrichthyans has shown the presence of an extensive canalicular network in the tesserae<sup>44</sup> and a trabecular tesseral network in some vertebral elements<sup>45</sup>, both resembling bone. 261 Whether these represent homologues of osteichthyan examples remains open to debate; 262

future works could employ synchrotron microtomography of stem-chondrichthyan cartilagesto address these questions.

265 Does endochondral bone have a deep origin within the gnathostome stem group? This 266 would imply repeated losses of this tissue. We do find statistical support for this hypothesis (Fig. 4, Table 1, Extended Data Figs. 5, 6, Supplementary Table 1), and the model is well 267 justified on prior phylogenetic and biological grounds. Endochondral bone has long been 268 269 known to be inconsistently developed across 'primitive' bony fishes: incomplete, 270 polymorphic, or entirely absent ossification of the endoskeleton is known in both Palaeozoic actinopterygians<sup>41,46,47</sup> and sarcopterygians{Cloutier:wm}, as well as more recent taxa<sup>48</sup>. The 271 272 frequent absence of endochondral bone in osteichthyans is considered secondary, and other 273 controlling factors such as body size, maturity, mechanical stress, and buoyancy can 274 determine its degree of development<sup>1</sup>. Our findings are also in agreement with studies establishing a genetic basis for secondary loss of all bone types within chondrichthyans<sup>49-51</sup>, 275

with the failure to produce endochondral bone likely representing arrested development of
chondrocytes as opposed to a primary lack of ability<sup>52</sup>.

278 Another confounding factor in this question is the problem of 'placoderm' 279 relationships. Although currently resolved in most analyses as a deeply pectinate grade along 280 the gnathostome stem (Fig. 4), the backbone of this arrangement has poor statistical support, 281 even in the present analysis (Extended Data Figs. 3). There is a lack of consistency in the 282 arrangement of plesia and Bayesian tip-dating methods have even recovered a monophyletic Placodermi<sup>53</sup>. *Minjinia* itself highlights this uncertainty, given its highly unexpected character 283 284 combinations. Notwithstanding its endochondral bone and crown-gnathostome-like inner ear structure, it resembles 'acanthothoracids'---the 'placoderms' widely considered among the 285 286 most removed from the gnathostome crown (i.e. most 'primitive'): it possesses deep epaxial 287 fossae either side of a prominent occipital ridge and a nuchal eminence otherwise seen only in acanthothoracids such as *Romundina*<sup>18</sup> and *Arabosteus*<sup>19</sup>. This apparent character conflict 288 289 could perhaps be more easily reconciled with a more coherent (though not necessarily 290 monophyletic) 'placoderm' assemblage. Indeed, the highly pectinate structure of the 291 'placoderm' grade seems symptomatic of an overemphasis on characters and taxa resembling 292 the crown group, thereby undersampling characters that could stabilise a clear picture of 293 'placoderm' interrelationships.

*Minjinia turgenensis* reveals new data on 'placoderm' endoskeleton and tissue diversity recorded from Mongolia—an otherwise extremely poorly known biogeographic realm for early gnathostomes. The phylogenetic placement of this 'acanthothoracid'-like taxon crownward of all non-maxillate 'placoderms', in conjunction with possession of extensive endochondral bone, highlights the importance of material from traditionally undersampled geographic areas. The presence of endochondral bone renews the hypothesis that this tissue is evolutionarily ancient and was lost secondarily in chondrichthyans<sup>6,33</sup>. This

301	view is overall consistent with evidence of generalised bone loss in chondrichthyans,
302	potentially as a result of the suppression of bone-generating molecular genetic pathways <sup>51,52</sup> .
303	Continued work in Mongolia and re-evaluation of phylogenetic datasets will be necessary to
304	address this, with the results likely to lead to substantial re-evaluation of gnathostome
305	phylogeny.
306	
307	
308	Methods
309	
310	X-ray computed microtomography. We scanned MPC-FH100/9.1 using the Nikon XT
311	225s at the Museum of Paleontology, University of Michigan with the following parameters:
312	200kV, 140 $\mu$ A, over 3123 projections and a voxel size of 32.92 $\mu$ m. We conducted
313	segmentation using Mimics 19.0 (http://biomedical.materialise.com/mimics; Materialise,
314	Leuven, Belgium) and we imaged models for publication using Blender
315	(https://www.blender.org).
316	Synchrotron light propagation phase contrast tomography. We imaged Diplacanthus
317	crassismus specimen NMS 1891.92.334 on Beamline 19 of the European Synchrotron
318	Radiation Facility, using propagation phase-contrast synchrotron microtomography. We
319	performed a spot scan with an energy of 116keV, achieving a voxel size of 0.55 $\mu$ m. We
320	processed the resulting tomograms using VG StudioMax 2.2 (Volume Graphics, Germany),
321	and prepared images in Blender.
322	<b>Phylogenetic analysis.</b> We conducted a parsimony analysis using TNT 1.5 <sup>54</sup> and Bayesian
323	analysis using MrBayes v 3.2.755. The dataset consisted of 95 taxa and 284 discrete
324	characters based on a pre-existing dataset <sup>56</sup> . We employed Osteostraci and Galeaspida as
325	composite outgroups. We conducted parsimony analysis using both equal weights and

326 implied weights methods. Global settings were 1000 search replicates and a hold of up to 1 million trees. Equal weights parsimony analyses were conducted using the ratchet with 327 328 default settings. Implied weights parsimony used a concavity parameter of 3 and the search 329 was without the ratchet. Command lists are included in Supplementary Information. We 330 conducted Bayesian analysis using both a partitioned and unpartitioned dataset. We used the Mkv model<sup>57</sup> and gamma rate distribution. We ran the analyses for 5 million generations with 331 332 a relative burn-in fraction of 0.25. Runs were checked for convergence using Tracer<sup>58</sup>. We partitioned the dataset using a newly proposed method<sup>28</sup> that partitions the data according to 333 334 homoplasy levels. Using the results of implied weights parsimony conducted in TNT, we created a text table of character fit values. We wrote an R<sup>59</sup> script to generate a list of 335 336 partition commands for MrBayes.

337 We assessed parsimony ancestral states visually using Mesquite<sup>60</sup>. Likelihood and Bayesian ancestral states were estimated in R using the castor package<sup>61</sup> version 1.5.7. Prior 338 to calculating likelihood ancestral states on parsimony trees, we scaled branch lengths using 339 PAUP\*<sup>62</sup> and calculated the likelihood scores for all of the trees under the Mkv model with 340 gamma rate parameter. The trees were then exported with branch lengths. To account for 341 342 overall uncertainty in tree estimates, we estimated ancestral states on 100 trees randomly 343 selected from the fundamental set of most parsimonious trees and two times 50 trees selected 344 from the 75% last trees of each posterior tree distribution from the Bayesian analysis. We 345 then run an ancestral states estimation Mk model (using the castor R package) using both the Equal Rates (ER) and All Rates Different (ARD) models. This resulted in 400 ancestral states 346 347 estimations. For each estimation we extracted the overlap log likelihood, the AIC (counting 348 one parameter for the ER model and two for the ARD model) and the scaled log likelihood 349 (probability) for the presence and absence of the endochondral bone character (character 4) 350 for the last common node of *Minjinia* and crown-group gnathostomes. We present the median

351	value of these distributions of the estimations overall log likelihoods, AICs and presence or							
352	absence of endochondral bone in Table 1.							
353								
354	Data availability							
355	The holotype specimen of Minjinia turgenensis will be permanently deposited in the							
356	collections of the Institute of Paleontology, Mongolian Academy of Sciences. Original							
357	tomograms are available at (doi:10.6084/m9.figshare.12301229) and rendered models are							
358	available at (doi:10.6084/m9.figshare.12301223). The phylogenetic character list and dataset							
359	are available as Supplementary Information S1 and S2. The LifeScience Identifier for							
360	Minjinia turgenensis is urn:lsid:zoobank.org:act:82A1CEEC-B990-47FF-927A-							
361	D2F0	B59AEA87						
362								
363	Code availability							
364	R code for generating partitions based on character fits and code for likelihood ancestral							
365	states reconstructions and plots are available in the Supplementary Information.							
366								
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513

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531

532 Author Contributions: MDB conceived and designed the study. MDB, AJ, YAA, and EZ 533 participated in all field seasons. RPD and AJ undertook preliminary CT scanning and 534 segmentation that revealed the fossil was a 'placoderm' and had endochondral bone. RS 535 discovered the first vertebrate remains in the first field season at Yamaat Gol in 2010. SG 536 undertook the segmentation of *Minjinia* with input from MDB. AJ performed segmentation 537 of Diplacanthus tissue. MC provided input on occipital comparative morphology of 538 'placoderms'. RPD provided data and comparative analyses and data for endoskeletal tissue. 539 YAA provided background on the geology, palaeontology, and stratigraphy of the type 540 location; EZ and YAA organized field logistics and permitting. MDB, SG, MC, RPD, and AJ 541 undertook the anatomical interpretation and prepared the figures. MDB and SG conducted the 542 phylogenetic analyses. RS conducted the parsimony branch support analyses. TG wrote the 543 script for generating MrBayes partitions from TNT's character fits table and conducted the 544 likelihood and model-fitting analyses. The manuscript was written by MDB, RPD, and SG.

## 546 **Competing interests statement:**

547 The authors declare no competing interests.





549 Fig. 1 | MPC-FH100/9.1 a 'placoderm' skull roof and braincase from the Early

Devonian of Mongolia. a, Ventral view. b, Dorsal view. c, Left lateral view. d, Posterior
view. e, Braincase endocavity in dorsal view. Taupe: endoskeleton; grey: mould; blue:
exoskeleton. crsp.ri, craniospinal ridge; e.hy.a, sulcus for the efferent hyoid artery; f.m.ep,
epaxial muscle fossa; fo.mag., foramen magnum; hy.fo, hyodean fossa; l.d.ao, sulcus for the
lateral dorsal aorta; N.V, trigeminal nerve canal; N.VII, facial nerve canal; N.VIII, acoustic
nerve canal; nch, notochordal canal; occ.ri, occipital ridge; orb, orbit; pr.pv, paravagal
process. Scale bar, 10 mm.



559 Fig. 2 | Endoskeletal mineralisation in fossil gnathostomes. a, Transverse tomographic 560 slice through MPC-FH100/9.1. b, Three-dimensional rendering of trabecular bone structure. c, Transverse tomographic section through the braincase of the osteichthyan Ligulalepis. d, 561 Three-dimensional rendering of the trabecular bone in *Ligulalepis* ( $\mathbf{c}$  and  $\mathbf{d}$  use data from<sup>56</sup>). 562 e, Synchrotron tomography image of the calcified cartilage of the certatohyal of the stem-563 564 group chondrichthyan Diplacanthus crassisimus specimen NMS 1891.92.334. f, Semi-565 transparent three-dimensional structure of calcified cartilage of NMS 1891.92.334. Scale bars, **a** and **b**, 10 mm; **c** and **d**, 1 mm; **e** and **f**, 150 μm. 566 567





Fig. 3 | Braincase endocavity of *Minjinia*. a, Semi-transparent rendering of skull roof and
braincase (grey and blue) showing extent of endocavity (pink). b, Ventral view. c, Dorsal
view. a.scc, anterior semicircular canal; cav.end, endolymphatic cavity; d.end, endolymphatic
duct; h.scc, horizontal semicicular canal; N.V, trigeminal nerve canal; N.VIIhm,

573 hyomandibular branch of facial nerve canal; N.VIIpal, palatine branch of facial nerve canal;

574 N.VIII, acoustic nerve canal; N.X, vagus nerve canal, N.X<sub>a</sub>, anterior branch of vagus nerve

575 canal; N.X<sub>p</sub>, posterior branch of vagus nerve canal; occ.a, occipital artery canals; p.scc,

576 posterior semicircular canal; sac, sacculus; soc, spino-occipital nerve canals; sup.opth, canal

577 for supra-ophtalmic nerve. Scale bars, 10 mm (upper scale bar associates with **a**, lower scale

578 bar associates with **b** and **c**).





endochondral bone maps closely with generalised loss of bone in chondrichthyans whereexoskeletal armour and perichondral bone are also absent.

588

Table 1 | Tree distribution (n=100) ancestral states estimation results. ER = Equal rates
model; ARD = All Rates Different model. The columns AIC and log.like represent the
median AIC and log.lik across the 100 parsimony and Bayesian trees (for both models). The
like.ratio column is the likelihood ratio for the models compared on these trees. The columns
Absent and Present represent the median scaled likelihood for the endochondral bone state.

trees	model	log.like.	like.ratio	AIC	node	Absent	Present
Parsimony	ER	-28.91	4.74	59.82	Minjinia:Gnathostomes	0.67	0.33
(equal weights)	ARD	-26.54		57.09		0.19	0.81
	ER				Crown Gnathostomes	0.91	0.09
	ARD					0.46	0.54
Bayesian	ER	-29.66	5.26	61.32	Minjinia:Gnathostomes	0.73	0.27
(unpartitioned)	ARD	-27.03		58.06		0.17	0.83
	ER				Crown Gnathostomes	0.79	0.21
	ARD					0.22	0.78



596 Extended Data Fig. 1 | Tomograms of endoskeletal ossification in *Minjinia*. Top row:

597 semi-coronal sections through braincase. Double-headed arrows indicate anterior-posterior

- 598 (a-p) dorsal-ventral (d-v) axes. Bottom row: semi-transverse sections through posterior part
- 599 of endocranium. Voids of black space represent mouldic preservation. Scale bars, 10 mm and
- 600 apply across each row of panels.



Extended Data Fig. 2 | Right orbital wall and innervation pattern of *Minjinia*. a, orbit in anterolateral view showing disposition of nerve openings (pink infill). b, endocast in the same perspective showing the relationship between nerve canals and endocast. a.scc., anterior semicircular canal; N.V<sub>2,3</sub> trunk of the trigeminal nerve canal for branches 2 and 3; N.VIIhm, hyomandibular branch of facial nerve canal; N.VIIpal, palatine branch of facial nerve canal; sac., sacculus; sup.opth, canal for supra-ophtalmic nerve. Scale bars, 20 mm (upper scale bar associates with a, lower scale bar associates with b and c).





Extended Data Fig. 3 | Results of phylogenetic parsimony analysis. Dataset consists of 95 taxa and 284 characters. Both trees are strict consensus topologies. Equal weights parsimony analysis using the ratchet resulted in 240 trees with a length of 831 steps. Implied weights parsimony analysis using random addition sequence + branch-swapping resulted in 8 optimal trees with score 85.20513. Double-digit figures above internal branches are bootstrap values of 50% and over; single-digit figures below branches are Bremer decay index values. Blue

- 618 shading: osteichthyan total group (dark blue: crown group); orange shading: chondrichthyan
- 619 total group (dark orange: crown group).
- 620



Bayesian: partitioned by character fits

621

622 Extended Data Fig. 4 | Results of Bayesian phylogenetic analysis using both partitioned

and unpartitioned data. Majority-rules consensus trees with posterior probabilities shown 623

- 624 along branches. Blue shading: osteichthyan total group (dark blue: crown group); orange
- 625 shading: chondrichthyan total group (dark orange: crown group).



628 Extended Data Fig. 5 | Likelihood ancestral state mapping of endochondral bone on

- 629 equal weights parsimony results. a, ARD, all rates different model; b, ER, equal rates
- 630 model.



633 Extended Data Fig. 6 | Likelihood ancestral state mapping of endochondral bone on

- 634 unpartitioned Bayesian analysis results. a, ARD, all rates different model; b, ER, equal
- 635 rates model.