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1 The diversification of lynx lineage during the Plio-Pleistocene——  
2 evidence from a new small *Lynx* from Longdan, Gansu Province,  
3 China  
4

5 Running title: A new small *Lynx* from China

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23       **Abstract:** A new small-sized lynx from Longdan, Gansu Province, *Lynx hei* sp. nov., is  
24 described in this study. The new species displays the characteristic *Lynx* generic traits, e.g.,  
25 distinct buccal grooves in the upper canine, presence of anterior groove in the upper canine,  
26 absence of the P2, and moderately developed mastoid process, but it is markedly smaller than  
27 the previously described *L. issiodorensis* specimens from the same site, and is also overall  
28 smaller than most living species, comparable to *Lynx rufus* in size. The new species has a  
29 relatively wide and deep zygomatic arch, similar to that of living *L. lynx*, *L. pardinus*, *L.*  
30 *canadensis* but wider than that of *L. rufus*. Our phylogenetic analyses suggest that *L. hei* falls  
31 within the crown group *Lynx*, being the sister group to *L. rufus*, or less probably to be a sister  
32 group to *L. issiodorensis* + three other living species of *Lynx*. The Plio-Pleistocene *L.*  
33 *issiodorensis* is supported as the ancestor of *L. lynx*, *L. pardinus* and *L. canadensis*. Our  
34 phylogenetic study suggests that *Lynx* diversification over the Plio-Pleistocene is first  
35 achieved by body size differentiation putatively forced by intraspecific competition with  
36 other carnivorans, and later followed by morphological divergence.

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39   Key Word: *Lynx*, diversification, Pleistocene, eastern Asia

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## 43 **Introduction**

44       The genus *Lynx* includes several medium-sized cat species with Holarctic distribution  
45 (Sunquist and Sunquist, 2002; Bellani, 2019). Four **modern** species are recognized in the  
46 genus, i.e., Eurasian lynx *Lynx lynx*, Iberian lynx *Lynx pardinus*, Canadian lynx *Lynx*  
47 *canadensis*, and bobcat **Lynx** *rufus*. The first three species are adaptive to dense forest in the  
48 temperate and subarctic regions (Tumlison, 1987; Heptner, 1992; Sunquist and Sunquist,  
49 2002; Poole, 2003; Bellani, 2019), whereas *L. rufus* is more like a generalist, although  
50 preferring forest environment when available (Lovallo and Anderson, 1996; Sunquist and  
51 Sunquist, 2002; Chamberlain et al., 2003). *L. rufus* is different from other species in both  
52 morphology and habitat, whereas the rest three species are less well distinguished (Werdelin,  
53 1981; Tumlison, 1987). Recent phylogeny of multilocus data and genomic scale (Johnson et  
54 al., 2006; Li et al., 2016), both supported the first divergence of *L. rufus* during the middle  
55 part of the Pliocene, followed by *L. canadensis* during the middle Early Pleistocene, and  
56 lastly *L. pardinus* and *L. lynx* during the late Early Pleistocene.

57       As a largely temperate lineage, *Lynx* has one of the best fossil records among Felinae  
58 (Werdelin, 1981). The discovery of the fossil material of *Lynx* can be dated back to the early  
59 **19th** century, i.e. *Felis issiodorensis* (Croizet and Jobert, 1828). More specimens assigned to  
60 the *Lynx* genus were later described in Asia (Teilhard de Chardin and Leroy, 1945), North  
61 America (Schultz and Martin, 1972), and even Africa (Hendey, 1974; Geraads, 1980, 2016).

62       *Lynx issiodorensis* represents the best-recorded and widely spread fossil *Lynx* from the  
63 latest Pliocene to Early Pleistocene (Werdelin, 1981; Kurtén and Werdelin, 1984). In Europe,  
64 *L. issiodorensis* gradually evolved to the now endangered Iberian species *Lynx pardinus*

65 (Werdelin, 1981; Boscaini et al., 2016; Mecozzi et al., 2021), whereas *Lynx lynx* probably  
66 evolved in Asia population of *L. issiodorensis*, and **did not appear** in Europe until the Late  
67 Pleistocene (Werdelin, 1981; Mecozzi et al., 2021). However, the records of *Lynx* in Eastern  
68 Asia Early Pleistocene were **found in a few localities, best represented the material from**  
69 **Longdan, Linxia Basin** (Qiu et al., 2004). Materials from Longdan were referred to *Lynx*  
70 *shansius* in previous works (Qiu et al., 2004), here *L. shansius* is considered a junior  
71 subjective synonym of *L. issiodorensis* following the interpretation of Kurtén and Werdelin  
72 (1984).

73 In this study, we study a new cranium from Longdan, Linxia Basin (Fig. 1, for detailed  
74 geological background, see Qiu et al., 2004). This new cranium is distinctly smaller than  
75 those of *L. issiodorensis* from the same locality, and the same taxon from Europe, further  
76 displaying characteristic *Lynx* traits, representing a previously unknown form and provide  
77 key information to understand the early diversification of this iconic Holarctic felid genus.

## 78 **Materials and methods**

79 The fossil material described in this study is housed in Hezheng Paleozoological  
80 Museum (HM). Fossil and living Felidae specimens (including living species of Felidae, as  
81 well as fossil *L. issiodorensis* **from both Europe and Asia**) housed in AMNH, USNM, IGF,  
82 IVPP, IOZ, KIZ have been observed for comparison.

83 The anatomical terms of the cranial anatomical terms of Jayne (1898) on the cat are  
84 followed with additional modification as in Qiu et al. (2014). For the craniodental  
85 measurement elements, see Fig. 2 for detail.

86 Our phylogenetic analysis is based on a new morphological matrix, partially from that of  
87 Salles (1992) and Werdelin and Flink (2018), including 87 characters for 23 taxa. The matrix  
88 is edited using software Mesquite 3.4 (Maddison and Maddison, 2021). The phylogenetic  
89 analyses of Maximum Parsimony using software TNT 1.5 (Goloboff et al., 2008; Goloboff  
90 and Catalano, 2016) and Bayes Inference using software Mrbayes 3.2.7 (Huelsenbeck and  
91 Ronquist, 2001; Ronquist et al., 2012b) were performed. All analyses were set for 4 chains,  
92 including three cold chains and one hot chain. Two independent runs were performed, each  
93 10 million generations. Constraint on the lineage of the living species using the backbone  
94 from phylogeny inferred from genomic data (Johnson et al., 2006; Li et al., 2016) was used.  
95 See the *Comparison and Discussion* section for more detailed settings.

96

97 **Institutional Abbreviations**—**AMNH FM** fossil mammal collection of the American  
98 Museum of Natural History, New York, USA; **AMNH M** mammalian collection of the  
99 American Museum of Natural History, New York, USA; **HMV** Hezheng Paleozoological  
100 Museum, Hezheng, China; **IGF**, Natural History Museum of Firenze, Italy; **IOZ** Institute of  
101 Zoology, Chinese Academy of Science, Beijing, China; **IVPP** Institute of Vertebrate  
102 Paleontology and Paleoanthropology, Chinese Academy of Sciences, Beijing, China; **KIZ**  
103 Kunming Institute of Zoology, Chinese Academy of Sciences, Kunming, China; **USNM**  
104 Smithsonian National Museum of Natural History, Washington DC, USA

105 **Anatomical Abbreviations**—**BW** blade width of the P4 (width across the paracone posterior  
106 to the protocone); **H** height; **L** length; **L1** skull total length; **L2** skull condylobasal length; **L3**  
107 skull basilar length; **LDP** diastema between the C-P3; **LT** toothrow length from C-P4; **M/m**

108 upper/lower molar; **Lme** metacone length; **P/p** upper/lower premolar; **Lpa** parastyle length;  
109 **W** width; **W1** zygomatic width; **W2** rostrum width across P4/M1; **W3** rostrum width across  
110 the canine.

## 111 **Results**

112 Order Carnivora Bowdich (1821)

113 Family Felidae Batsch (1788)

114 Subfamily Felinae Batsch (1788)

115 *Lynx* Kerr (1792)

116 *Lynx hei* sp. nov.

117 (Fig. 3)

118 **Referred specimen**—Type specimen HMV2010, almost complete cranium.

119 **Measurements**—Table 1

120 **Diagnosis** —small-sized lynx with similar overall size to that of living *Lynx rufus*.

121 Rostrum relatively narrow compared with most *Lynx* species. Postorbital process of frontal

122 small. Sagittal crest anteroposteriorly long and weak. Palate narrow. Basisphenoid-

123 basioccipital bone narrow. Upper canine with distinct anterior groove. P2 absent. P3 main

124 cusp high in buccal view. P4 with moderately sized protocone, without buccal groove.

125 **Etymology**—In honor to Mr. Wen He, the former curator of Hezheng Museum, who has

126 contributed to the fossil collection and management in Linxia Basin.

127 **Type locality**—Longdan, Linxia Basin, Gansu Province, China.

128 **Distribution and chronology**—So far known from the early Early Pleistocene in Linxia

129 Basin (ca. 2.4-2.2Ma).

130 **Description**—The cranium is complete, but slightly dorsal-ventrally compressed, so the  
131 ventro-dorsal height of this cranium is not original, but the width of the cranium is **weakly**  
132 affected. Judging from the dental wear (apex of the P3 slightly worn) and development of  
133 sagittal crest, the individual represents a full adult.

134 **Dorsal view (Fig. 3C)** The rostrum is relatively narrower than most other species of *Lynx*.  
135 There is not distinct postcanine constriction, but the rostrum gets slightly narrower backward.  
136 The forehead is slightly wider than the rostrum. The postorbital process of the frontal is  
137 distinct but not strong. The temporal ridges unit into the sagittal crest at the point posterior to  
138 the frontal-parietal suture. The lambdoid crest is well developed. **Lateral view (Fig. 3A)** The  
139 anterior contour of the naris opening is concave. The highest point of the cranium is located  
140 slightly behind the level of the postorbital process of the frontal in lateral view, in the anterior  
141 part of the cranium. The sagittal crest is low. The lateral contour of the C-P3 diastema is  
142 distinctly **dorsally concave**. The infraorbital foramen is located at the level of the mesial  
143 border of the P3. The anterior border of the orbit is located at the level of the P3 main cusp.  
144 The zygomatic arch has a uniform height and is slightly dorsally arched. The postorbital  
145 process of the jugal is distinct. The central ridge in the jugal is located around at the middle  
146 line of the bone. The mastoid process is distinct. **Ventral view (Fig. 3B)** The incisor row is  
147 nearly straight. The palate is relatively narrow for a lynx. The toothrow gets wider backward.  
148 The anterior border of the palatine is located at the level of the P3. The bony choana extends  
149 distinctly posterior to the M1. The preglenoid process is relatively weak. The foramen ovale  
150 is located medially to the glenoid fossa. The auditory **bullae are** large, whereas the



151 basisphenoid-basioccipital between the bullae is narrow. The posterior lacerated foramen is  
152 located in the same fossa as the hypoglossal foramen.

153 **Dentition** The incisors are relatively small. The I1 and I2 are in a similar size, whereas  
154 the I3 is distinctly larger. There are two strong vertical grooves in the buccal side of the  
155 canine. A slightly weaker vertical groove is present in the anterior side of the tooth. There is  
156 no P2 in the left, and a tiny root, which seems to be DP2, is present in the right side. The P3  
157 main cusp is high crowned in buccal view. There is no mesial accessory cusp, whereas the  
158 distal accessory cusp and posterior cingulum are well developed. There is weak distolingual  
159 convexity in the P3. The P4 has a straight buccal border. There is no vertical groove in the  
160 buccal side of the tooth. The ectoparastyle is absent. The parastyle is relatively short. The  
161 protocone is moderate in size, and anteriorly located. The M1 is small and transversely  
162 elongated.

## 163 **Comparison and discussion**

### 164 **The characteristic craniodental traits of genus *Lynx***

165 The most widely known trait for *Lynx* genus is the stable absence of the P2 (Werdelin,  
166 1981). We observed this trait in other felids based on observation of large collection in  
167 AMNH and USNM. The P2 is nearly always present in *Panthera* but is sometimes absent in  
168 *Neofelis* and *Prionailurus bengalensis*. The P2 seems to be frequently absent in some species  
169 of *Leopardus* and in *Pardofelis marmorata*, and *Caracal caracal*. It is absent in all species of  
170 *Lynx* we have observed, and all *L. issiodorensis* from Longdan, Gansu Province, and  
171 Caozhuang, Shouyang, Shanxi Province, but in two crania from the Mazegou Formation of

172 the Yushe Basin (Late Pliocene), it is present, though very small (probably DP2). Therefore,  
173 the absence of the P2, if taken alone, is not diagnostic for the genus *Lynx*, as several other  
174 felids regularly lack this tooth. The presence of the P2, however, suggests the felid is not very  
175 likely to be a *Lynx*.

176 Furthermore, the upper canines of *Lynx* genus display another discriminant character. In  
177 its buccal surface, two very deep vertical grooves are present in all living species of *Lynx*  
178 (Fig. 4). We studied a large sample of living felid species and found that such deep grooves  
179 are not shown by other feline cats, e.g., *Caracal* or *Leptailurus*, except some species of  
180 *Leopardus*, e.g., *Le. pardalis*. More interestingly, there is a distinct groove at the anterior side  
181 of the canine, just buccal to the anterior ridge of the canine. We check through the living felid  
182 species, and again, this groove is absent in all cats except most species of *Leopardus*. The  
183 grooves in the canine, therefore, provide more useful evidence to judge whether a cat could  
184 belong to *Lynx* genus.

185 The extant lynxes, except for *L. canadensis*, seldom develops mesial accessory cusp in  
186 the P3. It is also absent for all *L. issiodorensis* from Longdan we observed and rarely present  
187 in European *L. issiodorensis* and Early Pleistocene *L. pardinus* (Fig. 5). This trait is variable  
188 in many felids, e.g., *Leptailurus serval*, *Caracal caracal*, *Leopardus* spp., *Prionailurus* spp.,  
189 and *Felis* spp. The widest part of the tooth lies at the level around the main cusp/distal  
190 accessory cusp boundary (which means the distal side of the tooth is wider) in all species of  
191 living *Lynx* except *L. rufus*. In the latter species, the widest part is often more anteriorly near  
192 the apex of the main cusp. In *Leopardus* spp. and *Prionailurus viverrinus*, it is more common  
193 to see the widest part lies anteriorly, and this point lies posteriorly in *Prionailurus*

194 *bengalensis*, *Catopuma temminckii*, *Pardofelis marmorata*, but quite variable in *Leptailurus*  
195 *serval*, and *Caracal caracal* et al. Therefore, these two P3 traits only provide limited  
196 evidence for genus judgment but could be used in species inference.

197 The P4 protocone of all living species of *Lynx* is often somewhat reduced in size (Fig.  
198 5). This trait is commonly seen in *L. lynx* and *L. pardinus*, and less often seen in *L.*  
199 *canadensis* and *L. rufus*. In *L. issiodorensis*, the protocone is also often slightly reduced in  
200 size. A small preparastyle is quite common in all three living species, whereas in *L. rufus*, this  
201 small cusp is present in around the half specimens we observed. In *L. issiodorensis*, this cusp  
202 is also often present. Of particular interest, we noticed that in *L. issiodorensis* (both the Late  
203 Pliocene and Early Pleistocene populations from Europe and Asia) there are distinct vertical  
204 buccal grooves in most specimens (Fig. 6). This trait, however, is absent the living species  
205 (occasionally as shallow imprint in *L. lynx*) or fossil *L. pardinus*. We regard this trait as an  
206 important feature for *L. issiodorensis*.

207 Regarding the cranial features, *Lynx* is most characteristic by its broad cranium (wide  
208 zygomatic arch, wide rostrum, and wide forehead, see Fig. 7) for a medium-sized cat (smaller  
209 cats, e.g., *Felis*, could have an even broader cranium). This trait, however, can only be  
210 applied to *L. lynx*, *L. pardinus* and *L. canadensis*, whereas *L. rufus* only has a moderately  
211 wide cranium, comparable to that of the similar sized other cats, e.g., *Catopuma temminckii*,  
212 *Leopardus pardalis*, and *Leptailurus serval*. In lateral view, the ventral border of the palate in  
213 the C-P3 diastema is distinctly dorsally concave in all living species of *Lynx*, as well as in *L.*  
214 *issiodorensis*. This trait is generally absent or weak in *Leopardus* spp., *Felis* spp.,  
215 *Prionailurus* spp., but present in *Caracal caracal*, *Catopuma temminckii*, *Leptailurus serval*,

216 *Prionailurus viverrinus*, though in most of these latter species, the degree of convexity is not  
217 so strong as seen in *Lynx*. The zygomatic arch is rather deep and robust in all three species  
218 *Lynx* except *L. rufus*. None of the living small to medium-sized cats has such a deep  
219 zygomatic arch. The supraorbital process is strong and sharp in all living species of *Lynx* and  
220 being variable *L. issiodorensis*. In most feline cats, this process is sharp or variable. The  
221 mastoid process is moderately developed in all living species of *Lynx* and *L. issiodorensis*.  
222 The mastoid process is weaker in *Felis* spp., and most species of *Prionailurus* spp. but  
223 similarly strong in medium-sized felids, e.g., *Caracal caracal*, *Catopuma temminckii*,  
224 *Leptailurus serval*, *Prionailurus viverrinus*, so this trait is more likely a size-related character  
225 in the feline. The separation of the posterior lactated foramen and hypoglossal foramen is  
226 viewed as one of the most important traits to distinguish *L. lynx* from *L. pardinus*, in which  
227 the two foramina are located in the same fossa (Boscaini et al., 2016; Mecozzi et al., 2021).  
228 Recent observation suggests that the situation in *L. issiodorensis* is the same as in *L. pardinus*  
229 (Mecozzi et al., 2021). We checked this trait across the living species as well as in fossil  
230 Asian species. The confluence the posterior lactated and hypoglossal foramina is present in  
231 most species, in *L. rufus*, *L. hei* as well as *L. pardinus* as previously recognized (Fig. 7).  
232 However, it is clearly separated in *L. canadensis*, as in *L. lynx*. *L. issiodorensis* mostly has  
233 two foramina in the same fossa. In most other felids, these two foramina are in the same  
234 fossa, and this state is likely the primitive state for felids, including *Lynx*.

### 235 ***Lynx* affinity of the small cranium from Longdan**

236 The feline cranium H MV2010 from Longdan has characteristic *Lynx* craniodental

237 features. The cranium is relatively broad, and the upper canine has two deep buccal grooves,  
238 and one distinct anterior groove. The P2 is absent. This trait combination is not seen in any  
239 other genera of Felidae. Of noteworthy importance is the feature of the upper canine. As  
240 mentioned above, the presence of anterior groove is only present in *Lynx* among the Old  
241 World feline, and at least since the Late Pliocene. In the New World, most species of  
242 *Leopardus* also have deep buccal grooves and distinct anterior groove, but this genus  
243 typically retains P2 (except *Leopardus colocolo*, which frequently lack this tooth), and  
244 distinctly lower zygomatic arch, so the Longdan felid is unlikely a member of *Leopardus*.

245         The Longdan *Lynx* is also different from all living and known fossil species of *Lynx*. The  
246 body size of Longdan *Lynx* is distinctly smaller than that of *L. issiodorensis* from the same  
247 locality and other eastern Asian records (e.g., Yushe Basin, Nihewan Basin) and is also  
248 smaller than all living species except *L. rufus* (Fig. 8). The general morphology is similar to  
249 that of *L. rufus*, e.g., relatively narrow rostrum, P4 parastyle not elongated. However, The  
250 Longdan *Lynx* has a wider zygomatic width, and deeper zygomatic arch than those of *L.*  
251 *rufus*. The other three living species of *Lynx* have broader cranium and more elongated P4  
252 parastyle. In Europe, the early and middle Villafranchian lynx generally has a large body size,  
253 but since the late Villafranchian or slightly earlier, some smaller-sized population appeared  
254 (Mecozzi et al., 2021; Cuccu et al., 2022). The Longdan felid is still distinctly smaller than  
255 these lynx population (P4 of the small individual of late Villafranchian *L. issiodorensis* is still  
256 larger than 17 mm) (Werdelin, 1981). *L. issiodorensis* from Longdan shows clear affinity to  
257 *L. lynx* in size, dental robustness (Fig. 8A-B), but also differs significantly in relatively P3  
258 size (Fig. 8-E), rostrum length and width (Fig. 8G-I). A more detailed analysis of different

259 populations of *L. issiodorensis* and their comparison with the living species will be the topic  
260 for future more specific works. Both the type of *L. issiodorensis* and the Longdan population  
261 of this species are distinctly larger than *L. hei* in all lineal measurements, and has larger  
262 postorbital process of the frontal, but the craniodental proportions of the two species are  
263 similar. The type of *L. issiodorensis* also has a deeper zygomatic arch and proportionally  
264 smaller auditory bulla. *L. canadensis* is only slightly larger than *L. rufus*, but it differs  
265 significantly from the latter in several aspects, e.g., narrower cheek teeth, distinctly longer P4  
266 parastyle and wider rostrum. The Longdan felid is therefore represents a new species of *Lynx*,  
267 i.e., *L. hei* sp. nov.

## 268 **The diversification of *Lynx* during the Plio-Pleistocene**

269 To better study the phylogenetic position of *L. hei*, and the diversification of *Lynx* during  
270 the Plio-Pleistocene, we performed phylogenetic analyses for all living species of *Lynx*, *L.*  
271 *issiodorensis*, *L. hei*, and most living genera of Felidae, in fact the first phylogenetic analysis  
272 ever performed in fossil lynx. Note that our matrix coding of *L. issiodorensis* is based on  
273 materials from Shouyang, Yushe, and Longdan of eastern Asia.

274 In addition to Maximum Parsimony analysis, we perform Bayes Inference analysis  
275 under Mkv model (Lewis, 2001), as it outperforms Maximum Parsimony for estimation of  
276 phylogeny from discrete morphological data (Wright and Hillis, 2014), but see some debate  
277 in Goloboff et al. (2018). Our morphological matrix is based on matrix of Salles (1992) and  
278 Werdelin and Flink (2018), and expanded considerably based on our observation of all living  
279 and selected fossil species of Felidae. Those only characteristic of machairodonts (subfamily

280 Machairodontinae) were not included in the analysis. Apomorphy for each species is retained  
281 when available, as it is informative in the tip-dating analysis (Matzke and Irmis, 2018; Zhang  
282 and Wang, 2019). The gray wolf *Canis lupus* and the fossa *Cryptoprocta ferox* were chosen as  
283 the outgroup for Felidae following the choice of Werdelin and Flink (2018). Two Miocene  
284 felids *Proailurus lemanensis* and *Hyperailurictis intrepidus* were chosen to represent the  
285 primitive members of the subfamily. The majority of the living genera (here including 11  
286 genera in addition to *Lynx*) were included in the analyses. In Bayes Inference analyses, we  
287 performed both ordinary and tip-dating methods (Ronquist et al., 2012a). In the latter method,  
288 we did not include molecular data, so the tip-dating mainly acts as a constraint on branch  
289 length. As our fossil sampling of fossil feline is insufficient, we did not analyze the divergent  
290 time inferred from the tip-dating method, which can be more reliable when more fossil  
291 members were included in the analyses.

292 In two of the three analyses (MP, BI, tip-dating BI), the monophyly of *Lynx* is not  
293 supported, and *Otocolobus manul* mostly groups within *Lynx*. Even though molecular  
294 phylogeny clearly suggested that *O. manul* has a closer relationship to *Prionailurus* (Johnson  
295 et al., 2006; Li et al., 2016), our result is expected from a morphological aspect, as there is  
296 considerable convergence. *O. manul* has a broad cranium, wide forehead, absence of the P2,  
297 and reduction of the P4 protocone, elongated m1, that are also present in three living species  
298 of *Lynx* (not in *L. rufus*). In BI analysis, *Acinonyx jubatus* groups into *Lynx*, as there is also  
299 convergence between them, e.g., p4 anterior accessory cuspid enlarged, deep mandibular  
300 corpus. In MP analysis, the two species of *Panthera* are grouped with *Acinonyx*, together in  
301 *Lynx*, in a similar result of convergence. In the final analyses, we use constraint forcing

302 *Otocolobus* grouped with *Prionailurus*, and *Acinonyx* grouped with *Puma*, as we regard that  
303 molecular lineage defined by Johnson et al. (2006) as correct, considering its data size.

304 Nevertheless, all Bayes analyses (Fig. 9 and appendix), *L. hei* and *L. issiodorensis* are  
305 grouped within the crown group of *Lynx*. In MP analysis, *L. issiodorensis* is grouped with *L.*  
306 *lynx*, *L. pardinus* and *L. canadensis*, whereas the position of *L. hei* is not resolved in the strict  
307 consensus tree. We checked the original equal parsimony tree and found that this species is  
308 either sister to *L. rufus*, or sister to the group of *L. issiodorensis* + three living species of  
309 *Lynx*. In the BI analyses after constraining *Otocolobus* and *Acinonyx* out of *Lynx*, both tip-  
310 dating and ordinary analyses support the sister group *L. hei* and *L. rufus*, and *L. issiodorensis*  
311 to the other three species of *Lynx*, but in ordinary BI analysis, the posterior probability for the  
312 internal node within *Lynx* is rather low. In the tip-dating BI analysis without constraint, *L.*  
313 *issiodorensis* is grouped with *L. hei* + *L. rufus*, but with rather low posterior probability for  
314 this node. In most analyses (e.g., ordinary BI and MP analyses without constraint), the Asiatic  
315 golden cat *Catopuma* is supported as a sister group to *Lynx*. This result is congruent with the  
316 recent genomic phylogeny (Li et al., 2016). Another relationship of the recent genomic  
317 phylogeny, which *Felis* and *Prionailurus* are sister groups, is also supported in our analyses.  
318 Some other relationship, e.g., the sister group of *Puma* and *Acinonyx*, is not supported in any  
319 non-constraint analyses. This result is likely due to the lacking in intermedium form between  
320 the much more specialized *Acinonyx* and more conservative *Puma*, leaving a considerable  
321 morphological gap between them. Adding more fossil members will be helpful to break this  
322 long branch.

323 Our analyses provide interesting new information on the diversification of the crown



324 members of *Lynx*. The fossil records of *Lynx* are quite common in Eurasia, but mostly  
325 assigned to *L. issiodorensis*, which was supposed to be the common ancestor for the three  
326 other living species of *Lynx*, suggesting a simple widely distributed common ancestor-  
327 vicariance-speciation model. Our new finding suggested that *Lynx* crown group already  
328 experienced diversification during the Plio-Pleistocene. Molecular phylogenetic dating (Li et  
329 al., 2016) suggests a middle Pliocene divergence bobcat *L. rufus* to the rest of living species.  
330 Our new fossil and phylogeny support such a scenario. *Lynx* began to diverge into two  
331 ecomorphologically different species during the middle Pliocene, differs mainly by their body  
332 size. One population of the small-sized lineage migrated to North America and evolved there  
333 since then, finally evolved to the living *L. rufus*, whereas *L. hei* represents the population of  
334 this small-sized lineage retained in eastern Asia. At this evolutionary level, both large and  
335 small-sized lineage still have primitive craniodental traits, e.g., skull relatively narrow, and  
336 cheek teeth not enlarged. These traits are retained in *L. rufus*, whereas *L. issiodorensis*  
337 experienced to the second wave of diversification since the Pleistocene, and acquired more  
338 derived traits, e.g., broad skull, shorter rostrum.

339       It is interesting to note that after the larger sized lineage acquirement of a broad skull,  
340 shorter rostrum, it then began to occupy the niche of smaller body size: living *L. canadensis*  
341 has a comparative body size similar to *L. rufus*. This pattern suggests the first divergence of  
342 ecomorphology is achieved by evolving into different body size, and during the evolutionary  
343 history, the two lineages gradually their unique morphology, large enough to allow them to  
344 explore different niches without body size differences, as is seen in the living *L. canadensis*  
345 and *L. rufus* (Fig. 10).

346 **Conclusion**

347       The new Early Pleistocene fossil cranium from Longdan described here, represents a  
348 new species of lynx, *L. hei*, phylogenetically closely related to the modern North American *L.*  
349 *rufus*. Our phylogeny and comparative analyses suggest that the crown group of *Lynx*  
350 experienced its initial diversification during the mid-Pliocene in Eurasia into two lineages  
351 i.e., *L. hei-rufus*, and *L. issiodorensis*-**other** living species. The two lineages mainly differ in  
352 their body size during the Plio-Pleistocene. Whereas latter larger sized lineage acquired new  
353 craniodental morphology, and experienced a second wave of diversification in radiation in  
354 body size/morphology during the Pleistocene.

355  
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372

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449

450 **Figure captions**

451

452 Fig. 1. Location of Longdan.

453 Fig. 2. Craniodental measurement of *Lynx* used in this study. For measurements see

454 Abbreviation section.

455 Fig. 3. Holotype of *Lynx hei* sp. nov., H MV2010, from Longdan.

456 Fig. 4. Anterior view of rostrum of various *Lynx* species, marking the presence of the

457 anterior groove in the upper canine. A. *L. hei* sp. nov., H MV2010; B. *L. issiodorensis*,

458 H MV2015; C. *L. canadensis*, USNM212620; D. *L. rufus*, AMNH 24666. Not to scale.

459 Fig. 5. Comparison of occlusal view of upper cheek tooth rows of *Lynx* spp.. A. *L.*

460 *issiodorensis*, A1. H MV2012, Longdan; A2. AMNH 96-B1042 (reversed/mirrored),

461 Xiazhuang, and A3. AMNH 62-B756, both Shouyang; B. *L. hei* sp. nov., H MV2010,

462 Longdan; C. *L. rufus*, C1. AMNH M14822; C2. AMNH M243108; D. *L. pardinus*, D1.

463 AMNH M169492; D2. USNM 152619; E. *L. canadensis*, E1. USNM 077095; E2. USNM

464 212617; F. *L. lynx*, F1. AMNH 41337; F2. IOZ T0139.

465 Fig. 6. Comparison of buccal view of *Lynx* spp. A. *Lynx canadensis*, USNM 025726; B. *L.*

466 *rufus*, USNM 243108; C. *L. hei* sp. nov., H MV 2010; D. *L. issiodorensis*, ; E. *L.*

467 *issiodorensis*, Longdan, H MV2016; F. *L. issiodorensis*, Xiazhuang, Shouyang, AMNH 96-

468 B,1024. Not to scale.

469 Fig. 7. Crania of *Lynx hei*, and selected living felid species, lateral, ventral and dorsal

470 views. A. *L. hei*, H MV2010; B. *L. rufus*, AMNH M24665; C. *L. candaensis*, USNM33126;

471 D. *L. pardinus*, AMNH M169492; E. *L. lynx*, IOZ09419, F. *Caracal caracal*, AMNH

472 M116512.

473 Fig. 8. Craniodental measurements and ratio of *Lynx hei* sp. nov. and related taxa. A. P3  
474 length vs width/length ratio; B. P4 length vs width/length ratio; C. P4 length vs blade  
475 width/length ratio; D. P4 length vs parastyle length/length ratio; E. P4 length vs P3/P4 length  
476 ratio; F. condylobasal length vs tooththrow length; G. tooththrow length vs diastema/tooththrow  
477 length ratio; H. condylobasal length vs rostrum width cross P4 and m1/condylobasal length  
478 ratio; I. condylobasal length vs rostrum width cross C/condylobasal length ratio.

479 Fig. 9. Bayes Inference tip-dating analysis, with constraint on *Otocolobus* with  
480 *Prionailurus*. See the results of other phylogenies in appendix.

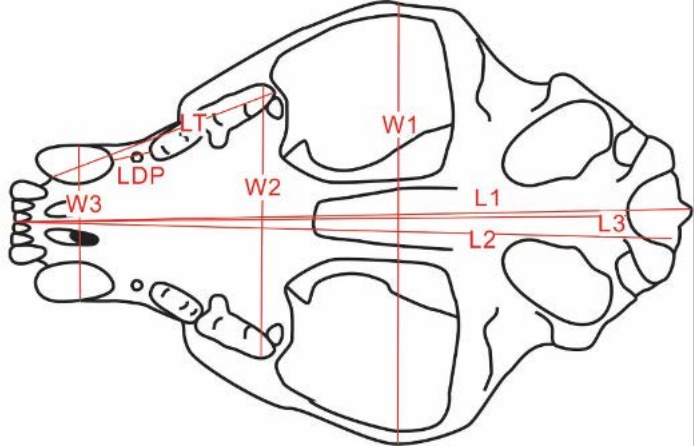
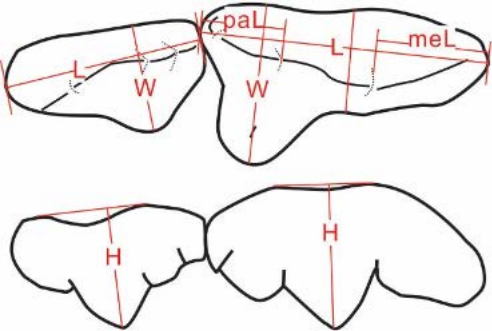
481 Fig. 10. Model of diversification pattern of *Lynx* proposed in this paper. The specimen of  
482 *L. issiodorensis* HVM2015 is from Longdan, China.

483

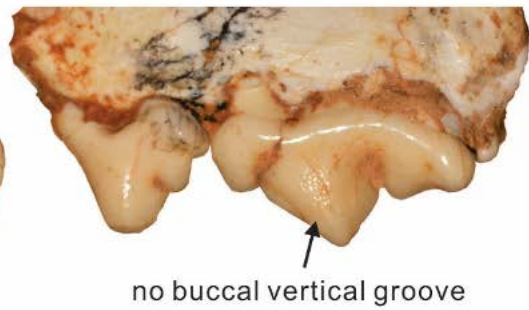
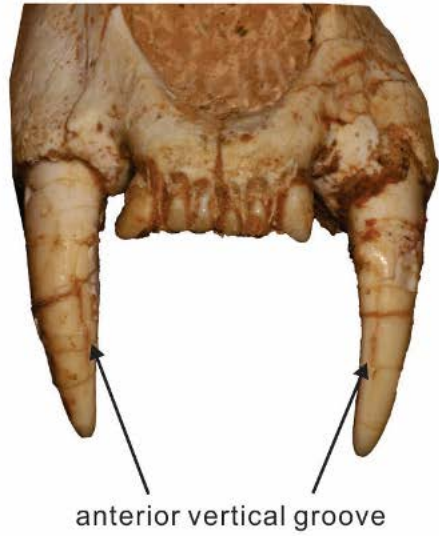
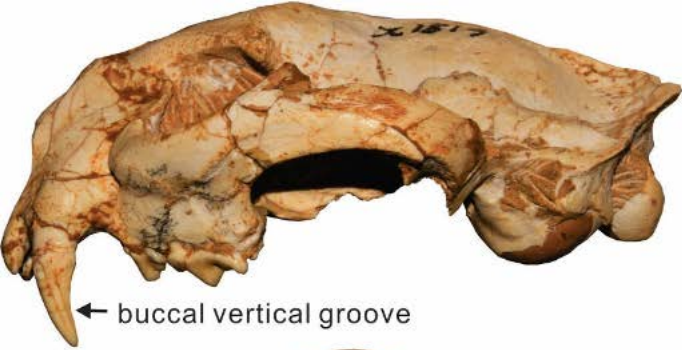
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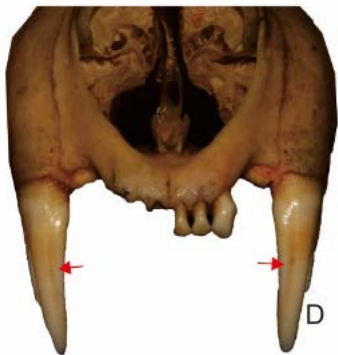
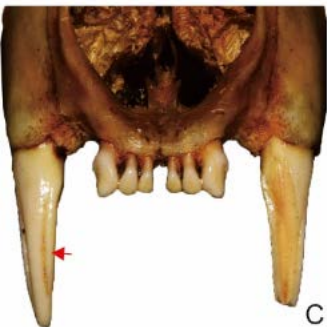
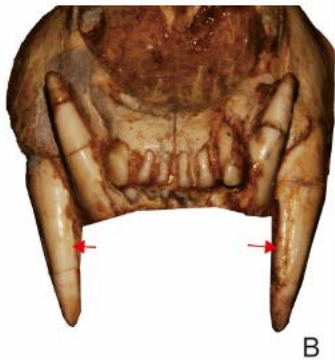
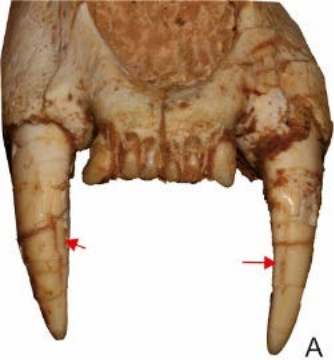


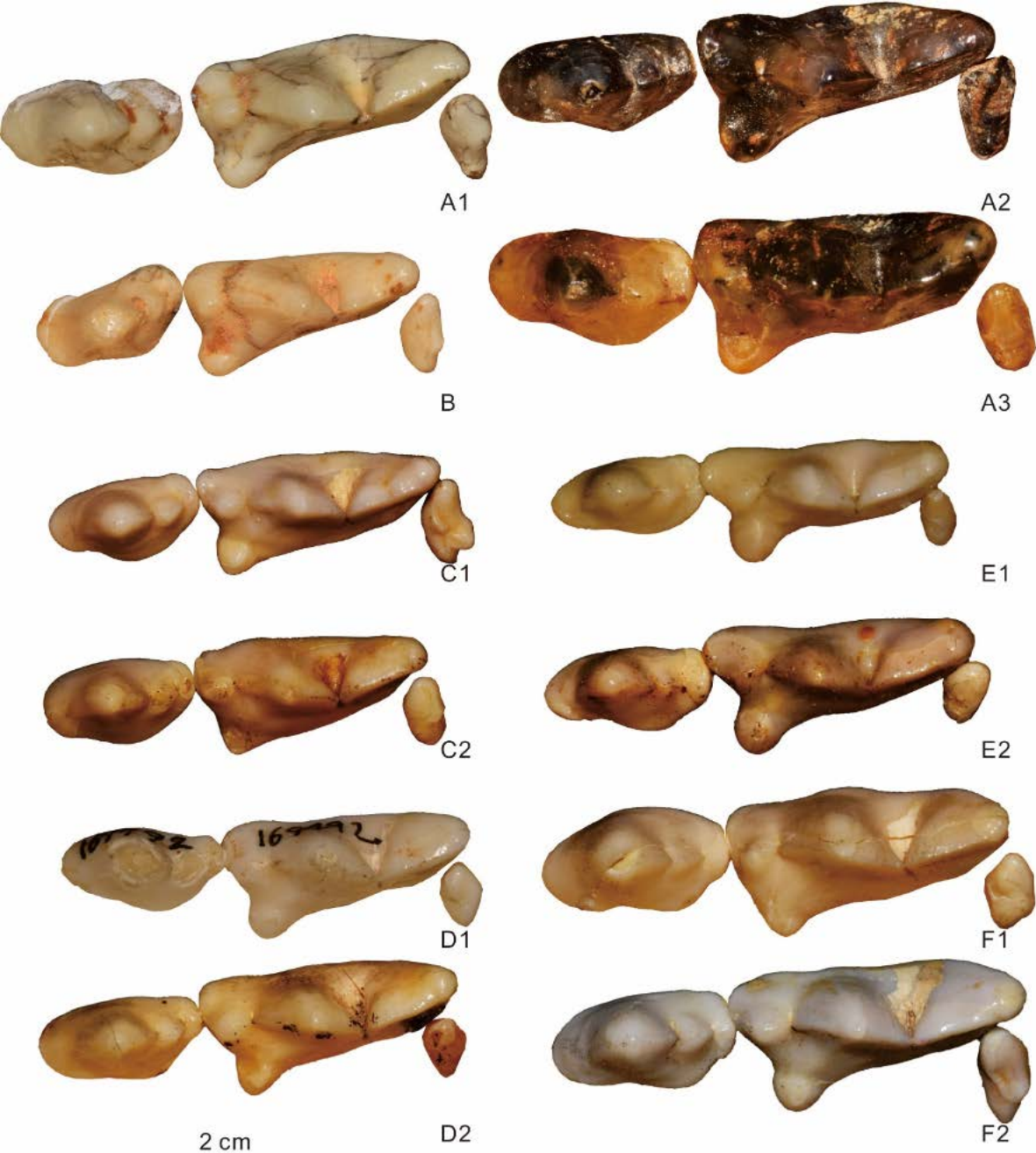
● Longdan

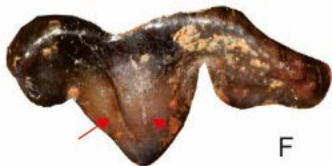
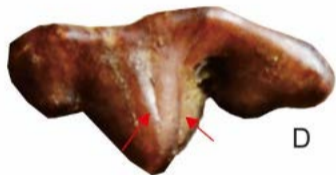




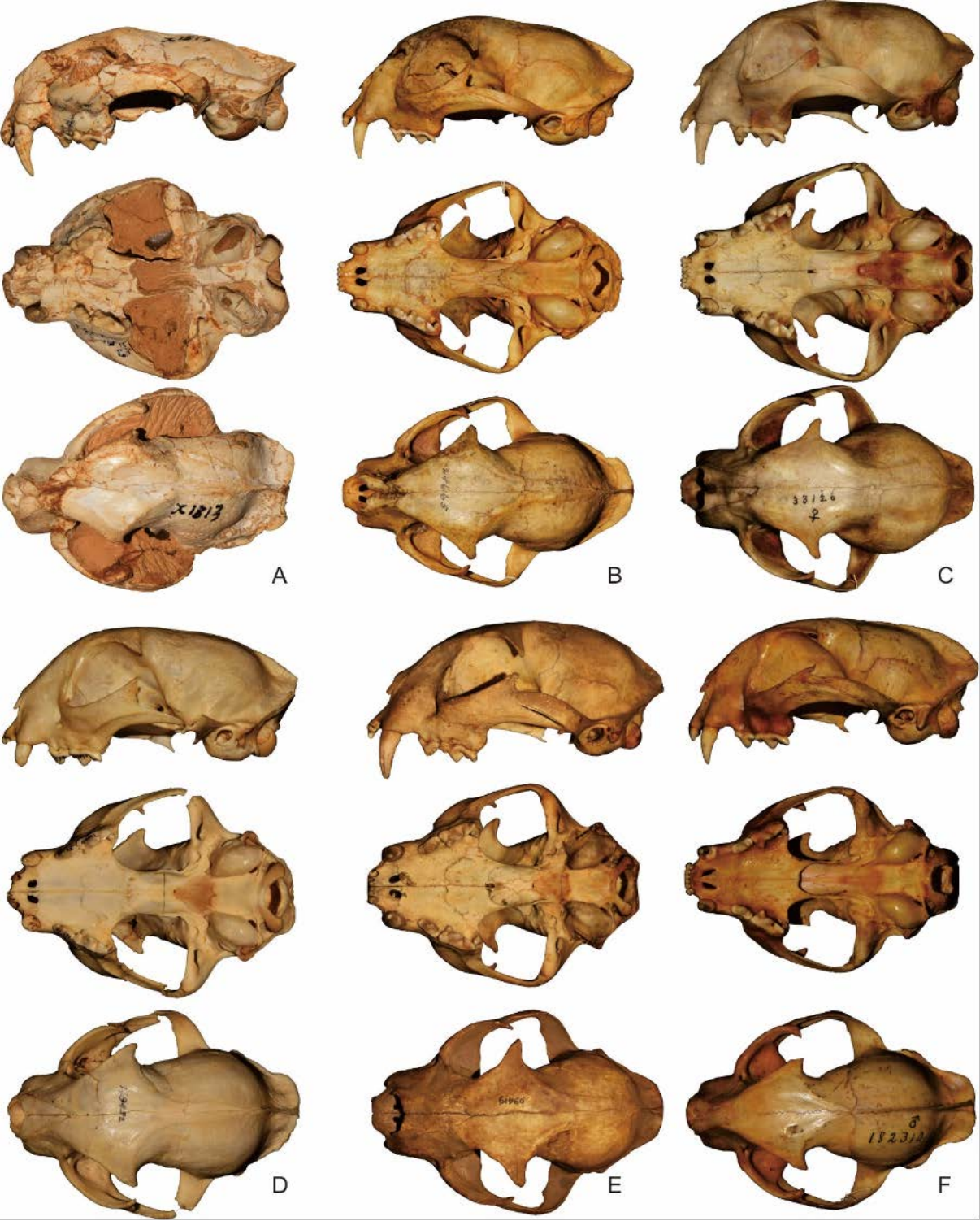


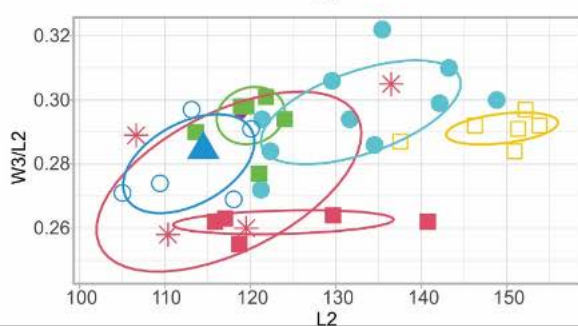
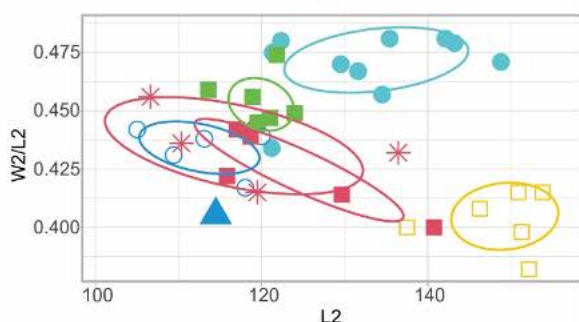
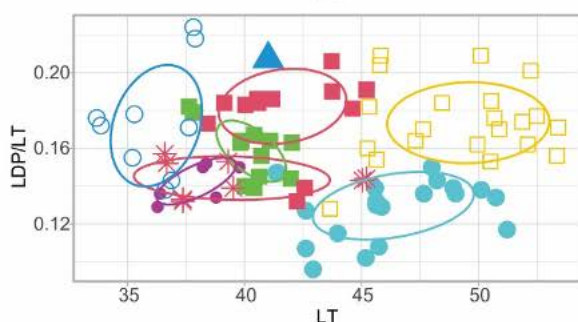
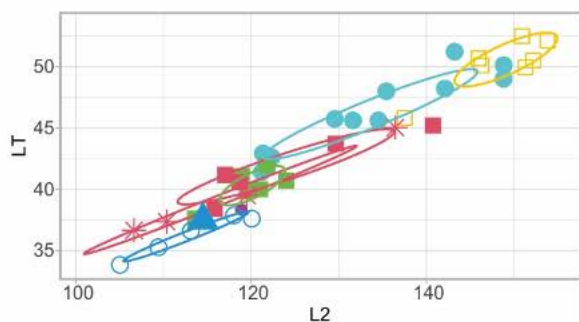
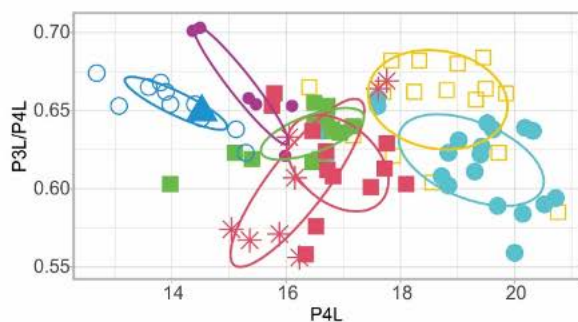
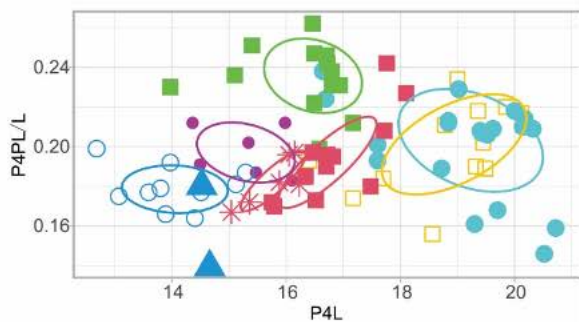
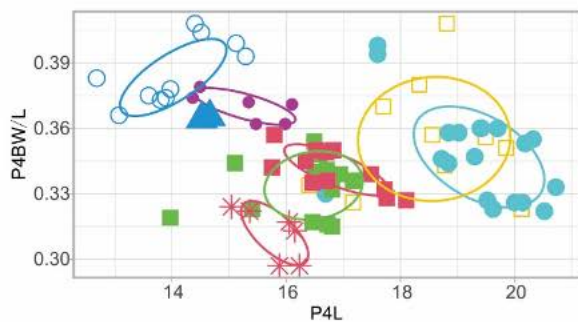
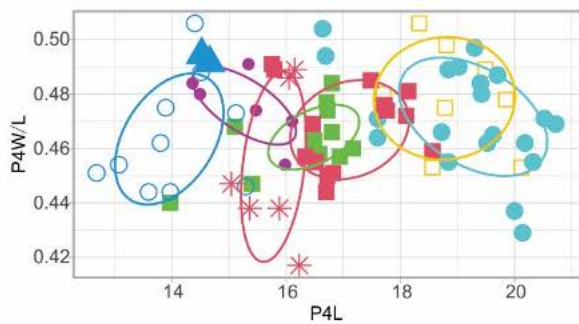
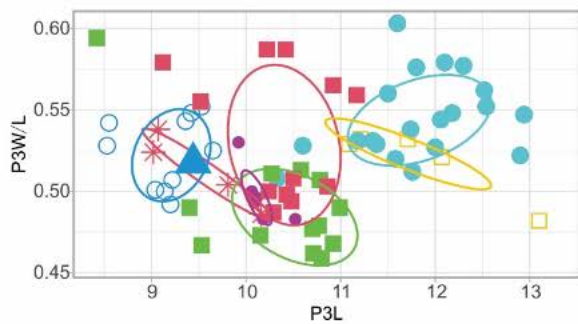






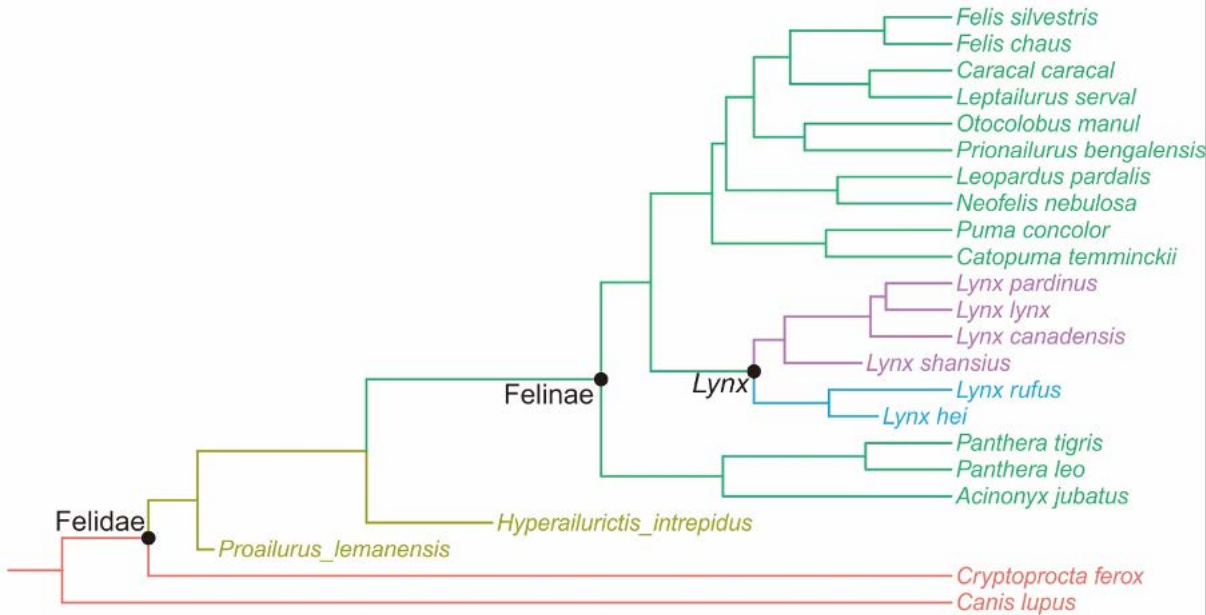






Taxon

- *Lynx rufus*
- *Lynx shansius*
- ▲ *Lynx hei*
- *Lynx canadensis*
- *Lynx lynx*
- ◆ *Lynx pardinus*
- ✱ *Caracal caracal*
- *Catopuma temminckii*





1993	old No.	No.	note	Character	0	1	2	3
		1		canine Upper canine morphology height	reduced (<0.13)	crown not elongated (~13-20% of c	moderately elongate (~20-25%)	25-35
		2		canine Upper canine buccal groove	0	1	2	3
		3		canine Upper canine lingual groove	0	1	2	
		4		canine Lower canine buccal groove	0	1	2	
	4	5		canine lower canine morphology	caniniform and large	small	very small and incisiform	
		6		M1 M1 transverse shortening	not shortened (>2)	shortened (<2)		
		7		M1 M1 cusp	presence of large, multicusp M1	presence of reduced, multi-cuspid	M no distinct cusp	
		8		P4 P4 protocone morphology	presence of large, usually cusped p	protocone small but still distinct	protocone very small or absent	
		9		P4 P4 protocone angulation	P4 protocone angulation	strongly anteriorly directed	lingually directed	
		10		P4 P4 protocone position	anterior to the paracone-parastyle ju	at the paracone-parastyle junction	posterior to the paracone-parastyle junction	
		11		P4 P4 parastyle anteroposterior length	absent or cingulum-like	small (<0.2)	moderate (0.2-0.25)	large (>0.25)
		12		P4 P4 buccal concavity	absent	one	two	
		13		P4 P4 preparastyle size	absent	small	large	
		14		P4 P4 preparastyle direction	nearly perpendicular to parastyle	with distinct angle with parastyle	nearly in line with parastyle	
		15		P4 P4 buccal vertical groove	absent	present		
		16		P4 Relative size of P4	P4 short relative to PL (<25% of PL	P4 long relative to PL (>25% of PL)		
		17		P3 Relative length P3/P4	P3 long relative to P4 (>60% of LP4	P3 moderate (ca. 50-60% of LP4)	P3 short (ca. 35-50% of LP4)	P3 very short (<35% of LP4)
		18		P3 P3 posterior cingulum	absent or weak	distinct	cuspid-like	
		19		P3 P3 anterior accessory cusp	absent	present	large	
		20		P3 P3 anterior cingulid cusp	absent	often present		
		21		P3 anterior end of P3 anterior ridge	close to the anterior border	distinctly lingually		
9		22		P3 P3 lingual ridge	absent	present		
		23		P3 Greatest width of P3	at main cusp	posterior part	distal part	
		24		P2 P2 morphology	large with multiple cusps	present but very small, usually only	absent	
		25		P2 P2 root	double	single		
		26		P1 P1 morphology	present	present but very small	absent	
		27		m3 m3 presence	Present	absent		
		28		m2 m2 presence	Present	absent		
		29		m1 m1 metaconid-talonid morphology	large talonid basin with distinct hypc	large without basin	small	absent
		30		m1 m1 metaconid and talonid differential	differential	undistinguishable		
		31		m1 m1 widest part	posterior to the paraconid-protoconid	at the paraconid-protoconid notch	slightly anterior to the paraconid-pro	near the middle part of the paraconid
		32		m1 m1 robustness (W/L)	slender (mostly <45%)	robust (~50%)		
		33		m1 anterior border	vertical or slightly posteriorly incli	distinctly posteriorly inclined		
		34		m1 Relative size of m1	short relative to mandible length (<1	long relative to mandible length (>15%)		
		35		p4 p4 relative length	long (>80% of m1 length)	short (<80% of m1 length)		
		36		p4 p4 paraconid direction	posterior inclined	vertical		
		37		p4 p4 paraconid size	distinctly smaller than the posterior	similar to the posterior accessory cu	larger than the posterior accessory cuspid	
		38		p4 p4 posterior cingulid	absent or weak	clearly distinguished	cuspid-like	
		39		p4 middle constriction	generally absent	generally present		
		40		p3 p3 size (versus m1)	small (<0.3%) or absent	relatively small (0.3-0.5)	large	
		41		p3 p3 height	same or slightly lower than the p4	distinctly lower than the p4 (<0.7)		
		42		p3 p3 anterior accessory cusp	absent	indistinct	distinct	
		43		p3 p3 posterior cingulid	absent or weak	clearly distinguished	cuspid-like	
		44		p2 p2 presence	Present	absent		
		45		p1 p1 presence	Present	absent		
		46		cranium dorsal concavity between C-P4	absent	present		
		47		cranium zygomatic arch depth	narrow	deep		
		48		cranium Anterior border of orbit	at anterior border of P3	P3	P3/P4 to P4 paracone	P4 metacone
		49		cranium Infraorbital fenestra position	at anterior border of P3	P3	at around P3/P4	posterior to P3/P4
		50		cranium Infraorbital fenestra size	very small (Acinonyx)	small (Felis)	large (Panthera)	very large (Smilodon)
		51		cranium Snout area elevation compared with braincase	snout area low	snout area distinctly elevated		
		52		cranium cranial dorsal contour	nearly straight	with distinct angle	rounded	
		53		cranium forehead width	distinctly wider than the rostrum	similar to the rostrum		
		54		cranium extension of posterior border of the nasals relative to maxi	not extended	similar	more extended	
		55		cranium posterior arch nasals	weakly arched	arched		
		56		cranium posterior border shape of the nasals	smooth	pointed		
		57		cranium anterior process of frontal	weak	strong		
		58		cranium nasal width	narrow	wide		
		59		cranium nasal middle constriction (Rostral constriction)	absent	present		
		60		cranium Nasal curvature	absent	present		
		61		cranium depression at the posterior end of nasal	absent	present		
		62		cranium Frontal bone, outer surface depression	absent	present		
		63		cranium anterior border of jugal	strongly anteriorly inclined	moderately anteriorly inclined		
		64		cranium Lower rim of the orbit	vertical	flat		
		65		cranium Size of the jugal postorbital process	small, rounded, often almost absent	rounded triangle	long, pointed	
		66		cranium Angle of the occipital condyles	low (~45-60)	high (~70-100)		
		67		cranium rostrum width vs palatine width	narrow, <0.7	wide 0.7-0.8	very wide >0.8	
		68		cranium palatine width versus CBL	widest part of the palate comparativ	widest part of the palate comparativ	widest part of the palate comparatively narrow	wide (>45%)
		69		cranium Alisphenoid canal	Present	absent		
		70		cranium occipital condyle position	low, much lower than the middle po	high, close to the middle point of inion and mastoid process		
		71		cranium Orientation of P3 and P4	tiny angle or nearly in line	small angle	large angle	
		72		cranium extension of bony choana	at or slightly posterior to the M1	distinctly posterior to the M1		
		73		cranium palate concavity medial to the M1	absent or weak	distinct		
		74		cranium position of rounded foramen	very close to the orbital fissure	at the middle of orbital fissure and oval foramen		
		75		cranium pterygoid fossa	absent	small	distinct	
		76		cranium middle ear opening	moderate	enlarged		
		77		cranium sagittal crest	always present	often or always absent		
		78		cranium postorbital bar	absent	often present		
		79		cranium size of ectotympanic	distinctly smaller than the entotymp	enlarged to similar size of ectotympanic		
		80		cranium anterior projection of jugal	absent	present		
		81		cranium postorbital constriction	absent or indistinct	distinct		
		82		cranium posterior expansion of frontal	absent or very weak	distinct		
		83		mandible number of mental foramina	mostly double	mostly single		
		84		mandible mandible height at canine	very low < 0.9	moderate (0.9-1.1)	high (1.1-1.3)	very high (>1.3)
		85		mandible lower border profile	nearly straight	distinctly convex		
		86		mandible Mandibular fossa termination	posterior to the camassial	well anterior of the posterior edge of the camassial, frequently terminating around camassial saddle or even at the m1/p4 junction		
		87		mandible Posterior part of horizontal dentary ramus height relative t	slender (~15-19%)	robust (~20-23%)	very robust >0.23	





Appendix for

A small *Lynx* from Longdan, Gansu Province and the  
diversification of *Lynx* lineage during the Plio-Pleistocene

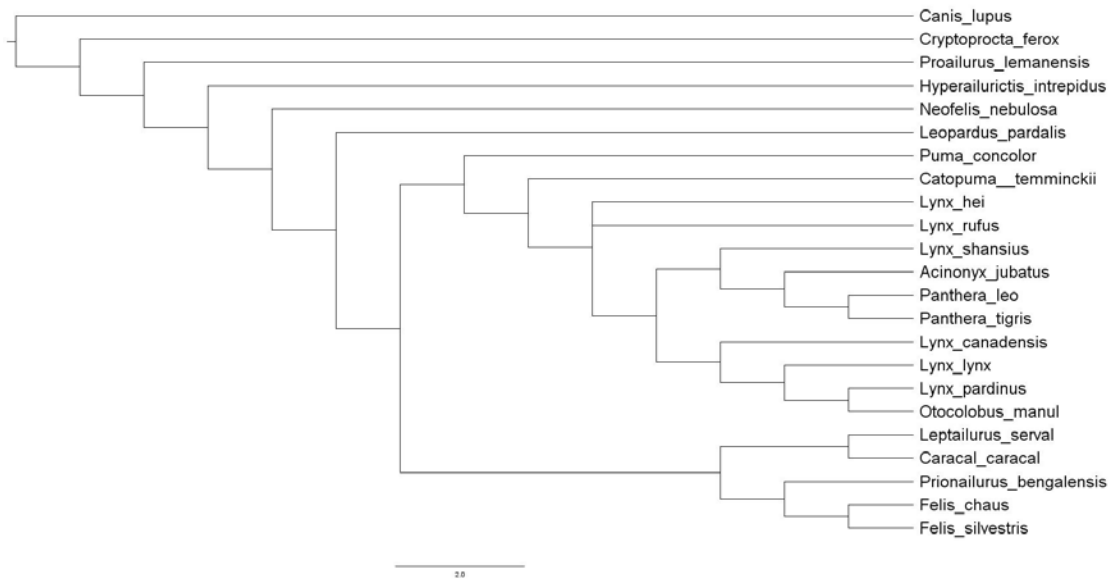


Fig. S1. Maximum parsimony analysis, without constraint.

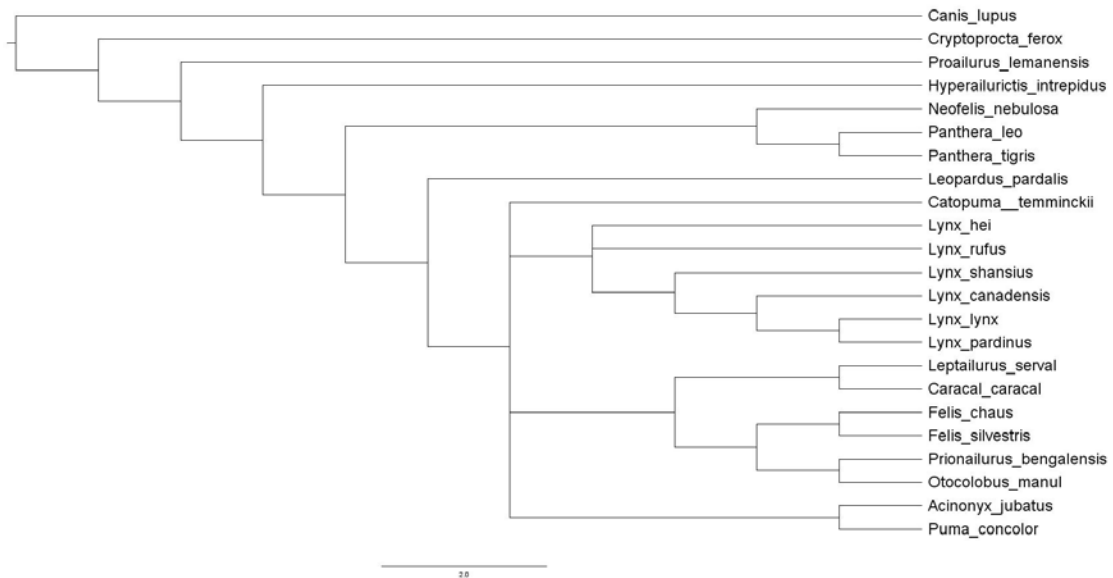


Fig. S2. Maximum parsimony analysis, with constraint on *Panthera* with *Neofelis*, *Acinonyx* with *Puma*, and *Otolobus* with *Prionailurus*.

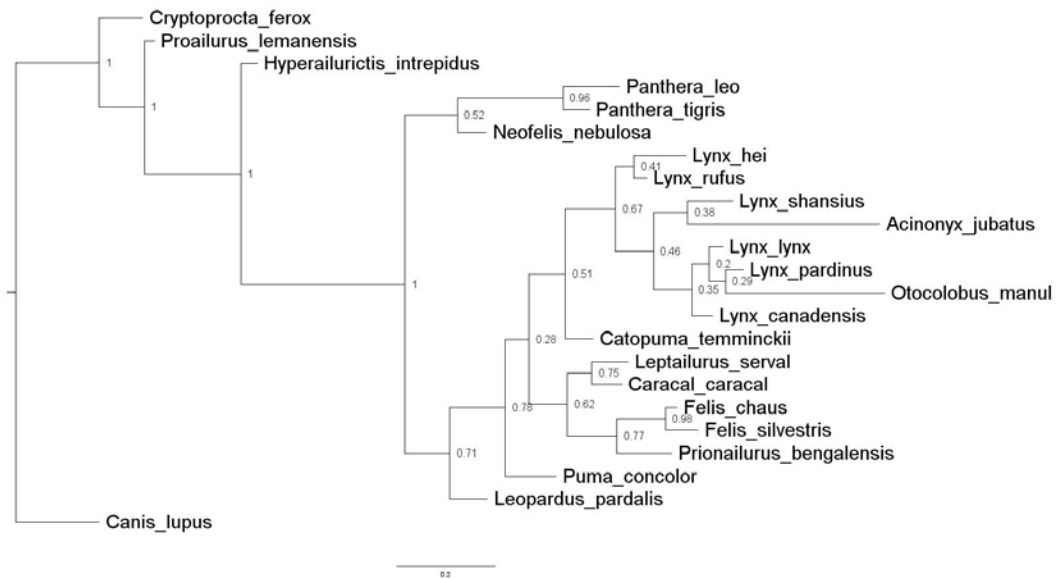


Fig. S3. Bayes Inference analysis, without constraint.

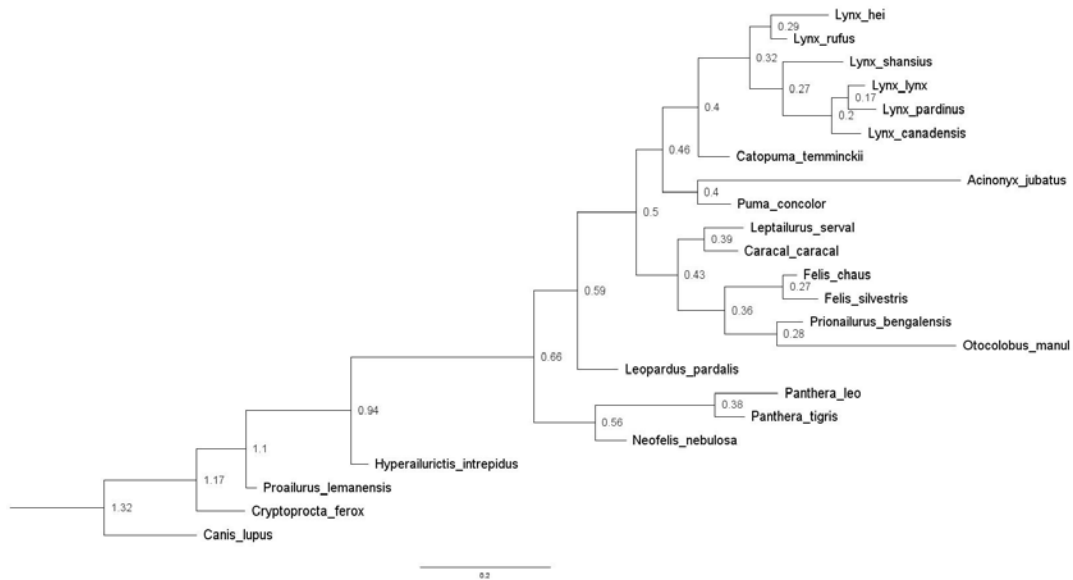


Fig. S4. Bayes Inference analysis, with constraint on *Acinonyx* with *Puma*, and *Otolobus* with *Prionailurus*.

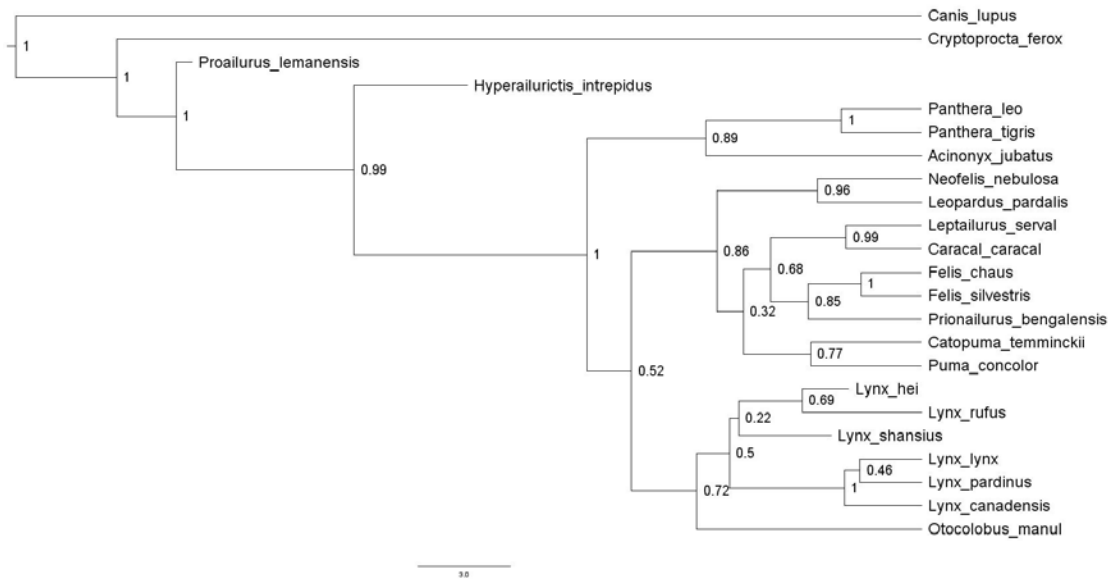


Fig. S5. Bayes Inference tip-dating analysis, without constraint.

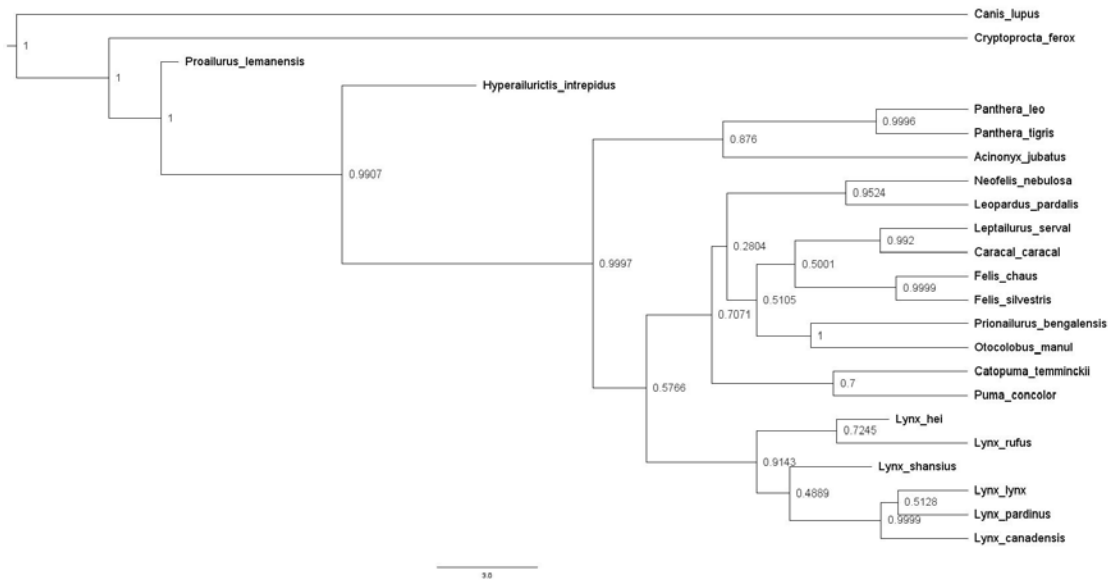


Fig. S6. Bayes Inference tip-dating analysis, with constraint on *Otocolobus* with *Prionailurus*.