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1  
2 **Title: A possible Middle Pleistocene Denisovan from the Annamite Chain of northern Laos**

3  
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67

68 **Abstract:** The Pleistocene presence of the genus *Homo* in continental Southeast Asia is  
69 primarily evidenced by a sparse stone tool record and rare human remains. Here we report the  
70 first Middle Pleistocene hominin specimen from Laos, with the discovery of a molar from the  
71 Tam Ngu Hao 2 (Cobra Cave) limestone cave in the Annamite Mountains. The age of the fossil-  
72 bearing breccia ranges between 164-131 kyr, based on the Bayesian modelling of luminescence  
73 dating of the sedimentary matrix from which it was recovered, U-series dating of an overlying  
74 flowstone, and U-series–ESR dating of associated faunal teeth. Analyses of the internal structure  
75 of the molar in tandem with palaeoproteomic analyses of the enamel indicate that the tooth  
76 derives from a young, likely female, *Homo* individual. The close morphological affinities with  
77 the Xiahe specimen from China indicate that they belong to the same taxon and that Tam Ngu  
78 Hao 2 represents, most likely, a Denisovan.

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80

81 **MAIN TEXT**

82 **Introduction:** From the Early to Late Pleistocene, the presence of *Homo erectus* is well  
83 documented in Asia, notably in China and Indonesia<sup>1-3</sup>. However, the taxonomic attribution of  
84 most Asian late Middle Pleistocene *Homo* specimens remains a matter of contention<sup>4-7</sup>. The  
85 recent description and analysis of the Harbin cranium from China has reignited this debate by  
86 suggesting its attribution to a new species named *Homo longi*<sup>8</sup>, but this taxonomic attribution of  
87 this specimens remains highly debated. In fact, the Harbin cranium shows close morphological  
88 similarities with other late Middle to early Late Pleistocene Asian *Homo* specimens from Dali,  
89 Xujiayao, Xuchang and Hualongdong, whose taxonomy remains unclear<sup>4,9,10</sup>. These fossils are  
90 considered to belong to a different taxon than *H. erectus* and are often grouped under the generic  
91 label ‘archaic humans’<sup>9,10</sup>. Due to the combination of features they exhibit, including  
92 Neanderthal-like traits, it has been suggested that they belong to an Asian sister taxon of  
93 Neanderthals, the Denisovans, even if this attribution to the latter group remains under debate  
94<sup>5,11,12</sup>. The small number of fossils currently securely attributed to this group (Denisova 2, a  
95 lower left molar; Denisova 3, a distal manual phalanx; Denisova 4, an upper left M3; Denisova  
96 8, an upper molar; and the Xiahe mandible)<sup>13-16</sup> prohibits a clear morphological picture of the  
97 overall Denisovan morphology. Their geographic distribution also remains debated. Modern  
98 Papuans, Aboriginal Australians, Oceanic/Melanesian, Philippine Ayta groups and, to a much  
99 lesser extent, mainland Southeast Asian populations, retain a Denisovan genetic legacy<sup>14,17,18,19</sup>.  
100 Combined paleoproteomic and morphometric analyses recently suggested that the Middle  
101 Pleistocene Xiahe mandible from Baishiya Karst Cave belonged to a Denisovan, extending the  
102 known range of this group onto the Tibetan Plateau<sup>15</sup>. However, there is still no fossil evidence  
103 explaining the Denisovans genetic imprint on modern southeast Asian populations and—due to  
104 the paucity of the Middle Pleistocene fossil record—it is still unknown whether one or more  
105 human lineages (co)existed in continental southern Asia. We present here the first unambiguous  
106 Middle Pleistocene *Homo* specimen from mainland southeast Asia and discuss its taxonomic  
107 attribution and implications for human evolution in the region.

108 In December 2018, a hominin permanent lower molar was recovered from a breccia block at  
109 Tam Ngu Hao 2 (Cobra Cave), Huà Pan province, Laos (20°12’41.5”N, 103°24’32.2”E, altitude  
110 1,116 m; Fig. 1, Fig. S1). The tower karst in which the cave was formed is positioned on the  
111 south-eastern side of P’ou Loi Mountain with an entrance located 34 m above the alluvial plain  
112 (Fig. 1A, Fig. S1). The site was discovered during a survey of the area around Tam Pà Ling,

113 where early *Homo sapiens* fossils have previously been recovered<sup>20-22</sup>. The tooth (TNH2-1) is a  
114 mandibular left permanent molar crown germ (Fig. 2A-F; Fig. S2), and the absence of occlusal  
115 and interproximal wear combined with the incipient root formation suggests that the tooth was  
116 unerupted at the time of the individual's death. The morphology of the tooth is compatible with  
117 an attribution to either a first or a second lower molar (Supplementary Material). In either case,  
118 considering the early maturational stage of the root, this tooth belonged to a juvenile individual  
119 corresponding to an age ranging from 3.5 to 8.5 years following modern developmental  
120 standards<sup>23</sup>.

121 To best document THN2-1, morphological description and comparative analyses were  
122 performed. We also developed a specific sampling protocol that allowed us to sample for  
123 palaeoproteomic and future isotopic analyses while preserving the whole occlusal surface  
124 morphology of the crown. Sampling for these destructive analyses took place after microCT  
125 analyses of the entire tooth, ensuring full morphological data were saved. No additional sampling  
126 for ancient DNA analyses was performed at this stage given the old age of the specimen and the  
127 tropical conditions under which the sediment and fossils were deposited. The invasive sampling  
128 strategy to collect dental tissues for molecular analyses only focused on the distal part of the  
129 inferior aspect of the crown, keeping the mesial portion of the crown intact.

130

131

## 132 **Results**

### 133 **Context and Dating**

134 The geological setting, stratigraphy and micromorphology of the sediment sequence were  
135 analysed to obtain a comprehensive, multi-scalar assessment of the depositional context and  
136 taphonomic history of the fossils recovered from the cave (Supplementary Material). The  
137 partially eroded sediments that infill the studied entrance passage comprise a lower and an upper  
138 facies representing two phases of sediment accumulation separated by an erosional surface and  
139 an unknown period of time (Fig. 1B). The lower facies (Lithological Unit 1, LU1) is weakly  
140 cemented and forms an arenitic silty clay deposit that is devoid of fossils (Fig. 1E). The upper,  
141 fossiliferous facies (Lithological Unit 2, LU2) is well cemented and coarse grained, containing  
142 intrakarstic angular limestone clasts and extrakarstic rounded pebbles, forming a very hard  
143 breccia/conglomerate layer from which skeletal elements—and in particular, teeth—were  
144 recovered in high frequencies (Fig. 1D). The change in lithology between the two facies most

145 likely reflects a reconfiguration of the karstic hydrological system as would be associated with a  
146 major flood, eroding space in LU1 onto which the sediments of LU2 were unconformably  
147 overlain. The sediments of LU2 are laterally contiguous and densely packed throughout the  
148 exposure excavated for this study, precluding major reworking of material and confirming the  
149 stratigraphic context of the fossils contained within, including the hominin tooth (see detailed  
150 observations described in Supplementary Material). The upper facies (LU2) is draped with two  
151 carbonate flowstones, indicating a final change in hydrology and the passage of surface water out  
152 of the cave and the precipitation of laminar speleothem (Fig. 1C).

153  
154 Three bovid teeth (TNH2-10/CC10, TNH2-11/CC11, TNH2-12/CC12) recovered from the upper  
155 fossil-bearing breccia (LU2) were directly dated using coupled uranium series and electron spin  
156 resonance (US-ESR), providing a weighted mean age estimate of  $151 \pm 37$  thousand years ago  
157 (kyr) (2-sigma) (Fig. 1B; Tables S1, S2) and an age range of 188-117 kyr. Two large blocks of  
158 breccia (LCC1 and LCC2) from LU2 (upper) and one block of the silty clay unit (LCC3) from  
159 LU1 (lower) were removed for luminescence dating (Fig. 1B). These samples produced coeval  
160 age estimates of  $143 \pm 24$  kyr (LCC1) and  $133 \pm 19$  kyr (LCC2) for the deposition of the LU2  
161 breccia and  $248 \pm 31$  kyr (LCC3) for the underlying LU1 silty clay deposit (Table S3). These  
162 ages are in stratigraphic agreement with the age of the overlying flowstone (CCF1), which was  
163 precipitated earlier than  $104 \pm 27$  kyr based on the weighted mean of U-series age estimates on  
164 four separate sub-samples of flowstone carbonate (Table S4). Bayesian modelling was performed  
165 on all independent age estimates to determine an overall geochronological framework for the site  
166 and tooth (Supplementary Material and Fig. S3). The fossiliferous breccia including the tooth  
167 was deposited between 164-131 kyr (at 68% confidence limit).

168  
169 **Fauna**

170 The Tam Ngu Hao 2 faunal assemblage comprises 186 identified dentognathic specimens (NISP)  
171 dominated by isolated teeth of large mammals, including several megaherbivores (Table S5).  
172 Their analyses reveal typical taphonomic pathways of assemblages from karstic systems in terms  
173 of representation of specimens and types of damage. Due to the energy associated with the  
174 deposition of (LU2), only teeth of large mammals are present in the assemblage, and we note the  
175 absence of small and light teeth of any microvertebrates. Moreover, most teeth are gnawed by  
176 porcupines, known to be a major accumulator agent in the region <sup>24</sup>. Therefore, the poor

177 preservation of specimens as shown in Fig. S15, precludes identification to the species level for  
178 most of the recorded taxa. The fauna bears close affinities to those known from the late Middle  
179 Pleistocene of southern China and northern Indochina and, to a lesser extent, Java, which is  
180 consistent with the sedimentary chronology of the site. It can be assigned to the “*Stegodon-*  
181 *Ailuropoda* faunal complex”<sup>25–28</sup>. We note the absence of Neogene taxa that persist in the Early  
182 Pleistocene and that of two key-species, *Pachycrocuta brevirostris* and *Gigantopithecus blacki*,  
183 which are good indicators of pre-300 kyr faunas in the region<sup>29–31</sup>. The archaic *Stegodon*  
184 persisted in Asia most likely until the end of the Late Pleistocene<sup>32</sup>. We recovered herbivores  
185 including *Tapirus*, *Stegodon*, and Rhinocerotidae, animals that were adapted to canopied  
186 woodlands in the area. We also found animals such as the *Bos* species, small-sized Caprinae and  
187 large-sized Cervidae (possibly *Rusa unicolor*), which are all known to exhibit a great variability  
188 in their preferred habitats, from closed and intermediate forests to open grassland<sup>33</sup>, and feeding  
189 behaviour.

190

#### 191 **Ancient proteins analyses**

192 The enamel from the TNH2-1 tooth specimen was analysed using nanoLC-MS/MS and the  
193 recently developed approach for ancient enamel proteomes<sup>29</sup>. The TNH2-1 proteome is  
194 composed of a common set of enamel-specific proteins, all of which have previously been  
195 observed in Pleistocene enamel proteomes<sup>34–36</sup> (Table S6). The enamel proteome has elevated  
196 levels of diagenetic protein modifications (Fig. S4A-D, Table S7) and preserves serine (S)  
197 phosphorylation within the S-x-E motif previously observed in ancient dental enamel<sup>34,35</sup> (Fig.  
198 S4E). Based on proteome composition and modification, as well as the absence of peptides  
199 matching to any of these proteins in our extraction and mass spectrometry blanks, we consider  
200 our proteomic data as indicative of endogenous proteins deriving from the sampled enamel.

201

202 Unfortunately, no high-confidence peptides overlapped diagnostic amino acid positions with  
203 sequence differences between *H. sapiens*, Denisovans, or Neanderthals, making further  
204 taxonomic assignment based on palaeoproteomics impossible. This is in line with previous  
205 research, which indicated that closely related hominin populations can be distinguished based on  
206 dentine and bone proteomes, while enamel proteomes are less informative in the context of close  
207 phylogenetic proximity<sup>35</sup>. Nevertheless, by comparing the sequences recovered from the TNH2-

208 1 enamel proteome with that of extant hominids for which protein sequences are available, we  
209 find that the specimen belongs to a member of the genus *Homo* (Table S8).

210 The absence of peptides specific to male-diagnostic amelogenin Y (AMELY) suggests that either  
211 the sampled molar was from a female individual or that AMELY-specific peptides were not  
212 observed due to degradation beyond the limit of detection of the instrument.

213

#### 214 **External and internal structural analyses of the tooth**

215 Externally, the TNH2-1 crown displays a coarse wrinkling pattern that is found in Pleistocene  
216 *Homo* (*H. erectus* s.l., European and Asian Middle Pleistocene *Homo* and Neanderthals), but is  
217 rare in modern *H. sapiens*. The mid-trigonid crest is well developed as commonly recorded in  
218 European Middle Pleistocene *Homo* and Neanderthals, while it is generally absent or less  
219 frequent in *H. erectus* s.l. and fossil and extant *H. sapiens*<sup>37</sup>. Below the external surface, the  
220 enamel-dentine junction (EDJ) of the tooth shows the dentine horns of the five main cusps and of  
221 a tuberculum intermedium and a low but uninterrupted mid-trigonid crest (Fig. 2A-H, Fig. S2;  
222 Supplementary Material). The latter feature is generally found in Neanderthals (80-100%  
223 depending on the molar position)<sup>38-40</sup> but is less frequent in *H. erectus* s.l. and *H. sapiens*<sup>41-47</sup>  
224 (Fig. S5). In addition, the EDJ of TNH2-1 shows an internally-positioned metaconid reminiscent  
225 of Neanderthal molars<sup>40</sup> and a low crown topography similar to that of *H. erectus*<sup>41-47</sup>. These  
226 features, as well as a slight buccal shelf present on the EDJ of TNH2-1, are all expressed on the  
227 EDJ of the Denisovan molars from Baishiya Karst Cave (Xiahe, Gansu, China) (Fig. S5)<sup>15</sup>.  
228 TNH2-1 dentine differs from the much higher and proportionally more mesiodistally compressed  
229 EDJ of Neanderthals and *H. sapiens*<sup>39,40</sup>, as well as from the shorter dentine horns and more  
230 densely wrinkled occlusal basin of *H. erectus* s.l.<sup>41-47</sup> (Fig. S5).

231

232 In terms of absolute dimensions, only Asian Middle Pleistocene *Homo* have larger tooth crowns  
233 than TNH2-1 (Tables S9, S10). TNH2-1 crown metrics are within the ranges of variation for *H.*  
234 *erectus* s.l., *H. antecessor*, Asian Middle Pleistocene *Homo* and Neanderthals, but they  
235 statistically differ from the smaller crowns of European Middle Pleistocene *Homo* and from  
236 Pleistocene and Holocene *H. sapiens* (Fig. 2I-J; Tables S10, S11). With respect to tooth crown  
237 tissue proportions, TNH2-1 has a high percentage of crown dentine (Vcdp/Vc: 55.37%) with  
238 moderately thick enamel as shown by absolute and relative enamel thickness values (3D AET:  
239 1.18 mm; 3D RET: 17.00; Table S12). These crown tissue proportions match to those of the



240 nearly unworn M2 of the Xiahe mandible <sup>15</sup> (Vcdp/Vc: 54.62%; 3D AET: 1.47 mm; 3D RET:  
241 18.97) and the upper molar of Denisova 4 (3D RET: 15.27; B. Viola, pers. comm.), but within  
242 the ranges of variation of all comparative fossil and extant human groups (Fig. S6A-C; Tables  
243 S12, S13). Three-dimensional maps of topographic enamel thickness distribution show that  
244 TNH2-1 has the thickest enamel at the top of the hypoconid and hypoconulid cusps and in the  
245 distobuccal quarter of the crown (Fig. S6D). In comparison, all other samples tend to have the  
246 thickest enamel distributed on all buccal cusps and more spread on the buccal aspect of the  
247 crown, even if variable between groups and between molar positions. The M2 of the Xiahe  
248 specimen shows thicker enamel spread along the buccal crown aspect but its distribution pattern  
249 is partly obliterated by occlusal wear.

250  
251 The EDJ shape of TNH2-1 was quantitatively compared with those of Pleistocene and Holocene  
252 human groups using geometric morphometrics (Supplementary materials). Landmark-based and  
253 surface deformation-based approaches were used, with both methods similarly distinguishing  
254 between *H. erectus* s.l., European Middle Pleistocene *Homo* and Neanderthals and *H. sapiens*  
255 using canonical variate and a between-group principal component analyses (Fig. 3, Fig. S7).  
256 Along CV2 and bgPC1, the higher EDJ and more externally set dentine horns of Neanderthals  
257 and *H. sapiens* are discriminated from the lower and more centrally positioned dentine horns of  
258 *H. erectus* molars. The CV1 and bgPC2 axes separate Neanderthals from modern humans, with  
259 the former having more internally placed mesial dentine horns and a more developed  
260 hypoconulid than the latter. TNH2-1 falls outside the ranges of all other groups. It has an  
261 intermediate EDJ shape between the low crown of *H. erectus* (but exceeding the variation of the  
262 latter group along CV1 and bgPC2) and the cusp position of Neanderthal molars (even if outside  
263 their range of variation along CV2 and bgPC1). TNH2-1's closest morphological affinity lies  
264 with the Denisovan specimen Xiahe, which also displays Neanderthal-like features (Fig. 3, Fig.  
265 S7).

## 266 267 **Discussion**

268 Reconstructing dispersals and ultimately evolutionary trajectories of *Homo* in Asia depends on a  
269 currently poor fossil record. The Asian late Middle Pleistocene fossil record is mostly limited to  
270 the eastern part of the continent <sup>4,8-10,15,48</sup>. Any additional human remains from this time period  
271 documenting the evolution of *Homo* in southern Asia might thus help confirm previous

272 hypotheses or reveal new lineages. Proteomic analysis of the TNH2-1 molar indicates that it  
273 belongs to a female individual of the genus *Homo*. Morphometric analyses of the external and  
274 internal crown structural organisation allow us to reject a number of hypotheses regarding  
275 species assignment. TNH2-1 has large crown dimensions and a complex occlusal surface that  
276 differentiates it from the smaller and morphologically simpler teeth of *H. floresiensis*<sup>49</sup>, *H.*  
277 *luzonensis*<sup>50</sup> and *H. sapiens*. The EDJ shape shows a mixture of Neanderthal-like and *H.*  
278 *erectus*-like features, closely resembling the M1 morphology of the Denisovan specimen from  
279 Xiahe (Fig. 2, Fig. S5). The similarities between TNH2-1 and *H. erectus* are mostly related to the  
280 proportionally lower crown, although *H. erectus* molars display even lower molar crowns and a  
281 narrower occlusal basin (Fig. 2, Fig. S5). The Lao fossil shows clear Neanderthal-like features  
282 such as a well-developed mid-trigonid crest and internally-positioned mesial dentine horns, but  
283 differs with its much lower EDJ topography and occlusal basin shape.

284 The differences from Neanderthals that we observe do not preclude TNH2-1 from belonging to  
285 this taxon and would make it the south-eastern-most Neanderthal fossil ever discovered.  
286 However, considering the morphological particularities of TNH2-1 in unison, as well as the  
287 high-degree of morphodimensional similarities with the molars of the Denisovan specimen from  
288 Xiahe, the most parsimonious hypothesis is that TNH2-1 belongs to this sister group of  
289 Neanderthals. If TNH2-1 indeed belongs to a Denisovan, this occurrence, along with the recent  
290 discovery of a Denisovan mandible from the Tibetan Plateau, a high-altitude, hypoxic  
291 environment<sup>15</sup>, would suggest that this Pleistocene Asian population possessed a high degree of  
292 plasticity to adapt to very diverse environments<sup>51</sup>. Available Denisovan dental remains indicate  
293 a mixture of traits consistent with the current paleogenetic evidence that Denisovans and  
294 Neanderthals are sister taxa<sup>13,14,51–53</sup> and are therefore expected to share some craniodental  
295 features<sup>15,54</sup>. This is further supported by recent analyses that identified possible Denisovan  
296 skeletal characteristics based on unidirectional methylation changes including traits that have  
297 been linked to Chinese fossils such as Xujiayao and Xuchang<sup>9,54</sup>. Denisovans are notable for  
298 their large dentition, with some Neanderthal-like crown features<sup>15,48,54</sup>, as well as distinctive  
299 cusp and root morphology<sup>14–16</sup>. In the absence of molecular analyses, looking for these  
300 combined features in the Asian human fossil record, including in fossils like the Penghu 1  
301 mandible from the Taiwan Strait<sup>55</sup>, may help identify more Denisovan specimens (Fig. S8).

302 The alternative hypothesis that TNH2-1 belongs to a group of Neanderthals that made an  
303 incursion into southeast Asia (see for example discussions on fossils that may demonstrate this  
304 dispersal from Maba and Dali)<sup>56,57</sup> cannot be outright rejected.

305  
306 The tooth from Tam Ngu Hao 2 Cave in Laos thus provides direct evidence of a Denisovan or  
307 Neanderthal female individual with associated fauna in mainland Southeast Asia by 164-131 kyr.  
308 This discovery further attests that this region was a hotspot of diversity for the genus *Homo* (Fig.  
309 S8), with the presence of at least five late Middle to Late Pleistocene species: *H. erectus*<sup>58</sup>,  
310 Denisovans/Neanderthals, *H. floresiensis*<sup>49</sup>, *H. luzonensis*<sup>50</sup> and *H. sapiens*<sup>20-22</sup>.

311  
312

313 **Data and material availability:** All mass spectrometry proteomics data have been deposited in  
314 the ProteomeXchange Consortium (<http://proteomecentral.proteomexchange.org>) via the PRIDE  
315 partner repository with the dataset identifier PXD018721.

316

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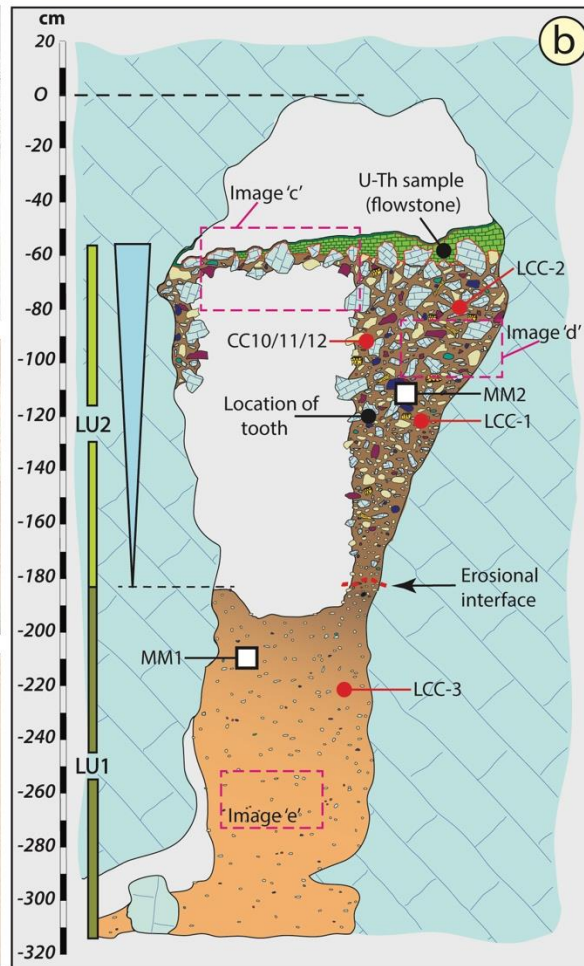
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490 **Author Contributions:**

491 Conceptualization: FD, LS, AMB, TS, VS, TL

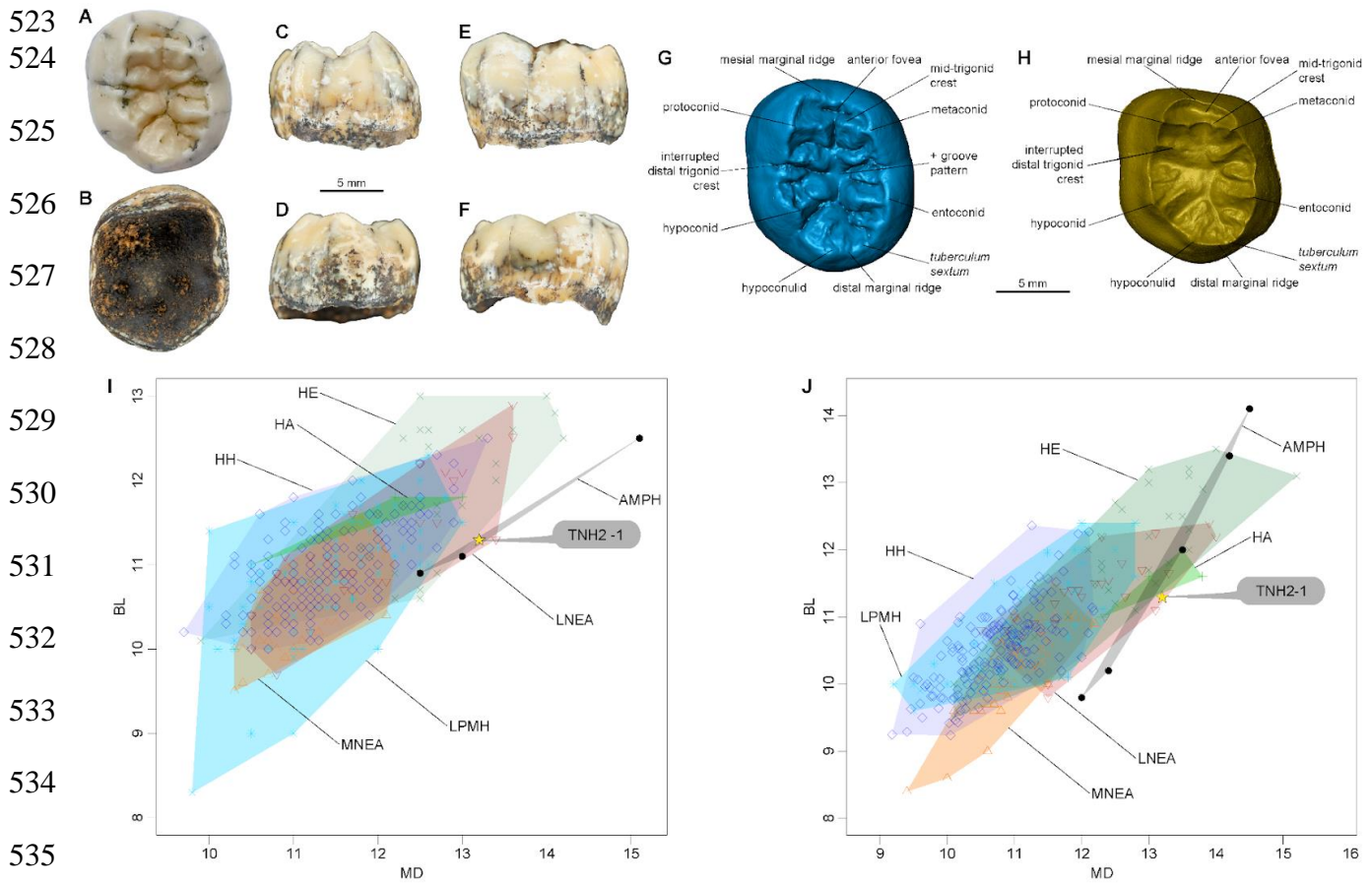
492 Methodology: CZ, MMS, KEW, RJ, RMB

493 Formal analysis: CZ, MMS, AMB, POA, MWM, LP, XS, J-JH, PD, J-LP, QB, FW, PLR, JVO,  
494 EC, HMC, KEW, RJ, J-XZ, RMB  
495 Investigation: FD, CZ, LS, PD, J-LP, ES, SF, TD, QB, AMB, FW, PLR, JVO, EC, HMC, KEW,  
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507 **Supplementary Materials**  
508 Materials and Methods  
509 Supplementary Text  
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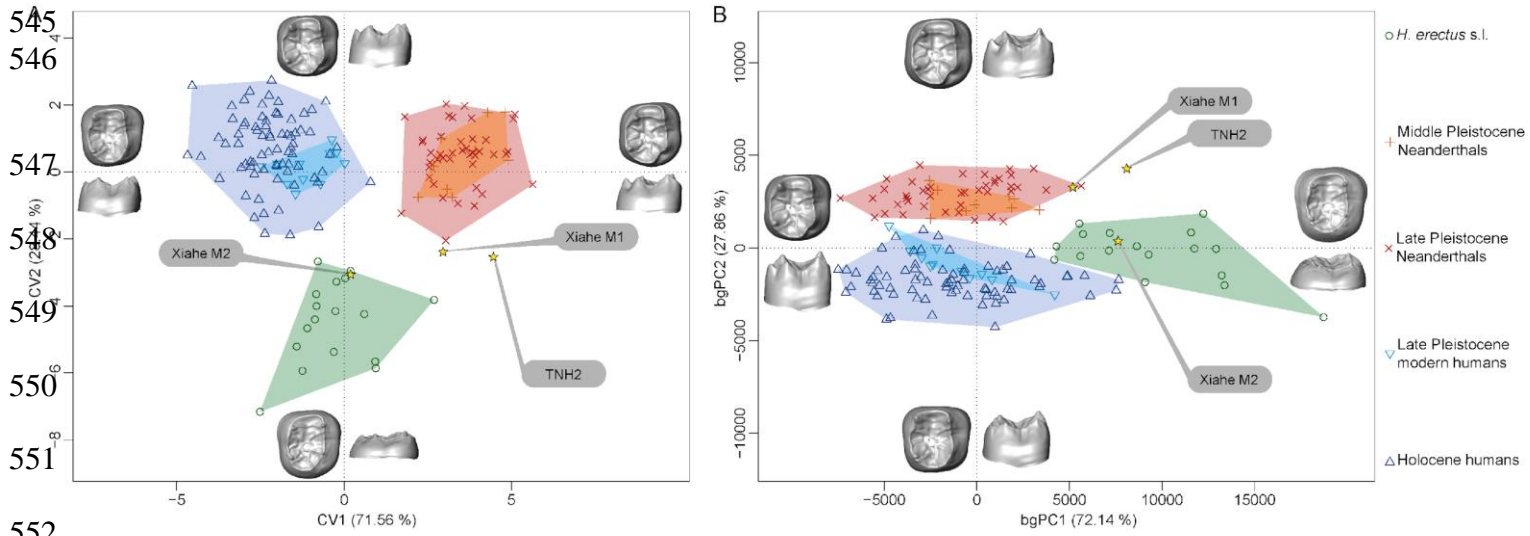


Limestone: substratum of the cave and reworked clasts	Arenitic silty clays	Arenitic breccias and arenitic conglomerates
Thin flowstone draping upper surface	Vein quartz pebbles	Altered granite pebbles
Sandstone clasts (upper: breccias, lower: pebbles)	Quartzite pebbles	Pisolites
LU Lithological Unit	LCC Luminescence sample	Flowstone capping fossiliferous breccia
CC US-ESR sample	MM Micromorphology sample	Teeth - bones

515 **Fig. 1. Geomorphological context and stratigraphy of TNH2.** A) aerial view of the site. The  
516 red circle indicates the entrance of Tam Ngu Hao 2 cave; B) stratigraphy and sampling locations  
517 of the infilling of the cave, showing Lithological Unit 1 and 2 (LU1 and LU2) with the erosional  
518 interface between these layers indicated by a dashed red line; Micromorphological  
519 (microstratigraphic) samples (MM1 and MM2) are also shown. Encircled numbers denote  
520 approximate positions of photographs in C, D & E; C) view of the flowstone capping the upper  
521 remaining part of LU2. D) detail of the arenitic breccia/conglomerate of LU2 ; E) detail of the  
522 arenitic silty clay of LU1.



537 **Fig. 2. Morphological and metrical features of the TNH2-1 specimen.** A-F, pictures of TNH2  
 538 in occlusal (A), inferior (B), mesial (C), distal (D), buccal (E) and lingual (F) views. G-H, virtual  
 539 renderings of the outer enamel surface (G) and enamel-dentine junction (H) in occlusal view  
 540 showing the main morphological features. I-J, bivariate scatter plots of the mesiodistal and  
 541 buccolingual crown dimensions of TNH2 compared with the M1s (I) and M2s (J) of *H. erectus*  
 542 (HE), *H. antecessor* (HA), Middle Pleistocene Neanderthals (MNEA), Late Pleistocene  
 543 Neanderthals (LNEA), Asian Middle Pleistocene *Homo* (AMPH), Late Pleistocene modern  
 544 humans (LPMH) and Holocene humans (HH).



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554

555 **Fig. 3. Canonical variate analysis (A) and between-group principal component analysis (B)**

556 **of the EDJ deformation-based shape comparison of TNH2-1, *H. erectus* s.l., the Denisovan**

557 **specimen from Xiahe, Neanderthals and *H. sapiens*.**