

The rise and fall of proboscidean ecological diversity

Authors: Juan L. Cantalapiedra¹, Óscar Sanisidro¹, Hanwen Zhang^{2,3}, María T. Alberdi⁴, José L. Prado⁵, Fernando Blanco⁶, Juha Saarinen⁷.

¹ Universidad de Alcalá, GloCEE – Global Change Ecology and Evolution Research Group, Departamento de Ciencias de la Vida, 28805, Madrid, Spain.

² School of Earth Sciences, University of Bristol, Wills Memorial Building, Queens Road, Bristol, BS8 1RJ, United Kingdom.

³ Department of Earth Sciences, Natural History Museum, Cromwell Road, London, SW7 5BD, United Kingdom.

⁴ Departamento de Paleobiología, Museo Nacional de Ciencias Naturales (CSIC), José Gutiérrez Abascal 2, 28006 Madrid, Spain.

⁵ INCUAPA, CONICET-UNICEN. Universidad Nacional del Centro de la Provincia de Buenos Aires. Del Valle 5737. B7400JWI- Olavarría, Argentina.

⁶ Museum für Naturkunde, Leibniz-Institut für Evolutions- und Biodiversitätsforschung, Invalidenstr. 43, 10115 Berlin, Germany.

⁷ Department of Geosciences and Geography, University of Helsinki, FI-00014, Helsinki, Finland.

*Correspondence to: jlopezcant@gmail.com

Abstract

Proboscideans were keystone Cenozoic megaherbivores, and present a highly relevant case study to frame the tempo and magnitude of recent megafauna extinctions against long-term macroevolutionary patterns. Surveying the entire proboscidean fossil history using model-based approaches, we show that the dramatic Miocene explosion of proboscidean functional diversity was triggered by their biogeographic expansion beyond Africa. Ecomorphological innovations drove niche differentiation, and communities that accommodated several disparate proboscidean species in sympatry became commonplace. The first burst of extinctions took off in the Late Miocene (~ 7 Ma). Importantly, this and subsequent extinction trends show high

32 ecomorphological selectivity, and went hand-in-hand with palaeoclimate dynamics. The global
33 extirpation of proboscideans began escalating from 3 Ma with further extinctions in Eurasia, and
34 then a dramatic increase in Africa extinctions at 2.4 Ma. Overhunting by humans may have
35 served as a final double jeopardy in the Late Pleistocene, following climate-triggered extinction
36 trends that began long before hominins evolved suitable hunting capabilities.

37 **Main text**

38 The worldwide extirpation of megafaunas was a radical upheaval in the recent evolutionary
39 history of terrestrial ecosystems, with Late Pleistocene human activities often considered
40 culpable^{1,2}. Yet, understanding these extinctions in view of long-term macroevolutionary
41 dynamics of the megafauna lineages has been critically lacking³. Proboscideans, being keystone
42 megaherbivores in Cenozoic terrestrial ecosystems, were among the most affected groups. For
43 centuries^{4,5}, their fossil record has elucidated an evolutionary history of success and decline in
44 equally dramatic measures, with three endangered living elephant species⁶ representing a mere
45 vestige of a once formidable clade high in both taxonomic diversity and ecomorphological
46 disparity, spread across Africa, Eurasia and the Americas. Therefore, proboscidean evolution
47 poses an invaluable case study for palaeobiologists to explore causes of uneven distribution in
48 biodiversity across phylogeny and evolutionary history. Hereby, we comprehensively re-examine
49 the rich fossil record of proboscideans in its entirety, and investigate the interactions between
50 proboscidean diversification and the tempo and mode of their ecomorphological evolution. Our
51 approach assesses clade and community-level dynamics, placing emphasis on the timing and
52 processes behind the sudden decimation of this group in the Quaternary. Our research presents a
53 blueprint for evaluating any plausible impact of Pleistocene humans on megafauna extinctions¹
54 against long-term macroevolutionary factors of decline.

55 We compiled a dataset for extinct and living proboscidean taxa with unprecedented detail,
56 consisting of 185 species, 2,130 fossil occurrences and 17 ecomorphological traits (including
57 body size, craniodental morphology, mode of mastication, tusk morphology, and foot posture;
58 see Extended Data Fig. 1), which are pertained to multiple fundamental aspects of proboscidean
59 functional ecology (e.g. breadth of dietary preference and feeding envelope, food processing,
60 energy requirements, home range, social grouping, sexual selection, and mode of locomotion).
61 Based on these traits, and through nonmetric multidimensional scaling (NMDS) analysis, we

62 constructed a two-dimensional functional morphospace of proboscideans that condenses 93% of
63 their ecomorphological variation (Fig. 1b). A first empirical dimension of proboscidean
64 functional morphospace (NMDS-1) encapsulates greater dental masticatory durability (DMD),
65 through increasing mesiodistal length, number of transverse loph(id)s, and relative crown height
66 (degree of hypsodonty) of the molars. These trends eventually facilitated dentitions with a high
67 proal shearing effectiveness during mastication, key to enhanced dietary plasticity and
68 processing of low-quality food⁷⁻⁹. A second dimension reflects larger-scale craniodental
69 modifications, including but not limited to: development of sharp or obtuse dental loph(id)s,
70 curvature of upper tusks, presence of shovel-like lower tusks and elongated mandible (Extended
71 Data Fig. 2). This functional space yields eight proboscidean functional types (PFTs), each
72 representing a cluster of ecomorphologically similar species, and thus experiencing similar
73 evolutionary pressures¹⁰ (Fig. 1a-c, and Extended Data Fig. 2 and 3). We reconstruct the global
74 and continental diversification trajectories of proboscideans based on occurrence data¹¹, and use
75 multidimensional phylogenetic models¹² to assess the mode of evolution of lineages across the
76 above-mentioned functional space (see below). We inspect mechanisms behind community
77 assembly and disassembly processes using general linear mixed-effects models (see methods).

78 **Early diversification and the dispersal outside Afro-Arabia**

79 During the first half of their history, until some 30 Ma, proboscideans were restricted to Afro-
80 Arabia. Species diversity across this interval rose steadily, but with no substantial ecological
81 diversification, and only two out of the eight PFTs evolved. Despite a good coverage in the Early
82 Eocene fossil record, initial diversification rate estimates should be read with caution given a
83 sizable gap in the record between around 51 and 40 Ma (see the broad 95% credibility intervals
84 in Fig. 1d). But, from then on, our estimates show that net diversification remained high
85 compared to other mammalian clades^{13,14} (average diversification was 0.08 species/Myr; see Fig.
86 1d). In fact, this rate was sustained until the collapse of the group in the Quaternary (Fig. 1d).
87 Importantly, our global diversification estimates aggregate clade-wise rate variations, and do not
88 imply rate homogeneity across lineages and geographic regions (Fig. 1d). With diversity
89 doubling every 9 Myr on average, we found little evidence of an asymptotic global carrying
90 capacity constraining proboscidean diversity (Extended Data Fig. 4), a scenario where speciation
91 and extinction rates are expected to balance each other rendering zero net diversification¹⁵. The

92 absence of a global diversity limit in proboscideans is likely the result of processes operating at
93 different scales: geographic expansion, local niche partitioning, and ecology-dependent species
94 survival.

95 The latest Oligocene and Early Miocene (25 to 20 Ma) witnessed the expansion of several
96 proboscidean lineages beyond of Afro-Arabia (termed ‘Proboscidean Datum Event’ by students
97 of Eurasian biochronology¹⁶ or PDE; Fig. 1c). Current understanding of proboscidean evolution
98 emphasises the importance of the PDE, since following this bio-event proboscideans became
99 ubiquitous in megaherbivore guilds across Eurasia as well as Africa^{16–18}. Around 16 Ma,
100 proboscideans incurred into North America, setting a new biogeographic milestone. Alongside
101 the progressive increase in seasonality and landscape heterogeneity during the Neogene¹⁹, major
102 biogeographic events like the PDE are expected to increase clades’ global carrying capacity²⁰,
103 boosting speciation opportunities²¹ and phenotypic evolution²², stemming from the exposure to a
104 wider array of environmental settings and biotic interactions. However, the impact of the PDE on
105 proboscidean macroevolutionary patterns has never been quantified. To further explore the
106 relevance of the PDE, we use multidimensional phylogenetic procedures¹² to test scenarios of
107 unconstrained, decelerating, and constrained evolution, as well as a suite of models where
108 African lineages evolved in a different fashion to those evolving in other continents (see
109 Methods). The best fit model shows that rates of ecomorphological evolution in African lineages
110 were slow and ameliorated over time (Extended Data Fig. 5), underlining the modest ecological
111 diversity of African lineages before the Neogene. Lineages outside Africa, on the other hand,
112 evolved 25 times faster (Extended Data Fig. 5). The inception of Eurasian lineages triggered
113 unprecedented radiation in both diversity and disparity of proboscideans, culminating in the
114 establishment and expansion of several new PFTs (Fig. 1c).

115 **Local and clade-scale patterns**

116 The Neogene expansion outside Africa drastically changed the course of proboscidean evolution,
117 both in their global and local diversity and disparity. The sustained emergence of functional
118 novelties translated into niche differentiation in proboscideans, facilitating faunas which
119 accommodated several sympatric proboscidean species^{23,24}. Linear models offer an insight into
120 trends in local proboscidean diversity, and how this relates to the masticatory functional
121 morphology of the constituent proboscidean taxa (see methods). Before the PDE, alpha diversity

122 remained relatively low. But following the PDE — around 22 Ma — and until 8 Ma, African and
123 Eurasian faunas were overall richer in proboscidean species, and this richness generally
124 increased over time (Extended Data Fig. 6). Over this period, community-level diversity is
125 significantly correlated with higher local mean NMDS-1 value ($P = 0.004$; Extended Data Fig.
126 7). Proboscideans responsible for this signal were the PFT 5 (e.g., amebelodonts or “shovel-
127 tuskers”) and PFT 6 (e.g., stegolophodonts) taxa, which underwent evolutionary modifications
128 towards enhanced DMD, as well as divergent feeding biomechanics (i.e., enlarged functional
129 lower incisors in PFT 5²⁵, and foreshortened jaws for greater mechanical efficiency of
130 mastication in PFT 6⁹). Higher prevalence of these functionally specialised taxa reduced the
131 intensity of interspecific competition, allowing herbivore guilds to accommodate more
132 proboscidean species.

133 In sum, increasing global ecomorphological diversity, combined with niche partitioning in
134 sympatric proboscidean species, produced high initial speciation rates in Eurasia and the
135 Americas, and also fuelled sustained net diversification in Africa. Together with the poor
136 performance of phylogenetic models where ecomorphological evolution decelerates as a function
137 of time (Extended Data Fig. 5), the multilayer approach presented here reinforces an
138 expansionist scenario of global proboscidean diversification, mediated by geographic dispersals,
139 amelioration of local competition, and relentless exploration of different functional
140 ecomorphological adaptations.

141 Local processes also impacted proboscidean macroevolutionary patterns via ecology-mediated
142 diversification. The dynamic and heterogeneous late Neogene and Quaternary biospheres
143 imposed more austere resource limitations, accelerating the evolution of multi-loph(id) dentition
144 optimised for effective proal shearing (highest DMD) in stegodonts and elephantids (PFTs 7 and
145 8)^{9,26}. As testified by successive peripatric speciation events during the evolution of
146 mammoths^{27,28}, novel adaptive responses of peripheral populations to marginal conditions²¹
147 played a key role in the diversification of PFT 7 and PFT 8 taxa. At the broadest scale, our
148 ecology-dependent diversification models¹⁴ reveal that these taxa with further increasing DMD⁹
149 show higher macroevolutionary persistence, signified by higher speciation rate and moderate
150 extinction risk (Fig. 2, Extended Data Fig. 8 and 9), a finding consistent with the rapid expansion
151 of PFTs 6, 7 and 8 in the late Neogene and Quaternary (Fig. 1c). Furthermore, ecology-selective

152 survivorship alone (species sorting) is a macroevolutionary force capable of influencing
153 evolutionary trends²⁹, and has probably been a major factor propelling lineages into unexplored
154 regions of the proboscidean functional space.

155 **Late Neogene turnover**

156 The onset of C₄ grass-dominated habitats around 8 Ma, which brought about less productive and
157 markedly seasonal conditions³⁰, brought dramatic changes to the evolutionary context of
158 megafauna communities^{26,31}. After the C₄ shift, proboscidean communities began to deplete. Our
159 community-level analyses depict a sustained decrease of local proboscidean richness over time
160 (Extended Data Fig. 6 and 7), as decreasing productivity in seasonal savannah-grassland
161 biomes³² could not sustain multiple competitively robust proboscidean species with further
162 enhanced DMD and greater foraging flexibility^{8,33,34}. For the first time, herbivore guilds
163 dominated by these new PFTs (6, 7 and 8) contain fewer proboscidean species than those with
164 more ecomorphologically disparate proboscidean assemblages ($P < 0.001$). Through most of the
165 Neogene, extinction was highly biased towards the more archaic proboscidean taxa with
166 primitive dentition (Extended Data Fig. 10). In the Late Miocene, extinctions eroded a broader
167 spectrum of the functional space (Fig. 3b). Deinotheres (PFT 2) and trilophodont gomphotheres
168 (dominant components of PFTs 3 and 5) were major casualties. Interestingly, whereas
169 gomphotheres are characterised by high volatility (high speciation and extinction rates),
170 deinotheres showed remarkable ‘living-fossil’-like macroevolutionary tendencies^{35,36}, being
171 persistently low in diversity and deficient in apomorphy acquisition (Fig. 1c, 2b).

172 Meanwhile, the expansion of PFTs 6, 7 and 8 sustained further net positive diversification.
173 Increased climatic perturbations and ecosystem heterogeneity would have promoted allopatric
174 speciation and local selection^{21,27,28}. Given the decline of local diversity, the maintenance of
175 global diversity would have stemmed from enhanced taxonomic provinciality between regions
176 (beta diversity). At the macroevolutionary scale, this shift is more clearly reflected in Eurasian
177 diversification trends (Fig. 1d). Around 7 Ma, sustained extinctions produced a brief decline in
178 Eurasian diversity (Extended Data Fig. 4). Eurasian proboscidean diversity managed to recover
179 at around 5 Ma, as speciation rates underwent a four-fold increase and extinction stabilised
180 between 6 and 3 Ma (Fig. 1d).

181 **The Quaternary collapse**

182 With newly-established PFTs showing ample diversification potential, no evidence of a diversity
183 equilibrium being approached (a maximal diversity of 33 species is estimated around 3.2 Ma),
184 and no signal of evolutionary stagnation, nothing would have predicted the abrupt collapse of the
185 clade in Quaternary times⁷. Although speciation rate in Eurasia remained high, extinction
186 aggravated at around 3 Ma, progressively hampering net diversification, as exemplified by the
187 rapid coming and going of numerous dwarfed insular species before the Late Pleistocene. In
188 Africa, extinction rates suddenly quintupled the background Neogene rates around 2.4 Ma (Fig.
189 2a), a pulse that saw decline in African elephantid (PFT 8 taxa) diversity³⁷. These events mark
190 the first phase of broad-scale global proboscidean demise, which first wiped out the few relict
191 PFT 5 species in North America , and then the last deinotheres (PFT 2) in Africa. The axe also
192 reached the brevirostrine gomphotheres (PFT 6) and mammutids (PFT 4), removing broader
193 portions of the functional morphospace than ever before (Fig. 3b). The decimation of
194 proboscidean communities continued (Extended Data Fig. 7). There is a substantial shift towards
195 a higher local mean NMDS-1 from 2.4 Ma onwards (Extended Data Fig. 7), reflecting a wipeout
196 of “mastodont-grade” taxa in Eurasia and Africa and their replacement by PFTs 7 and 8^{8,33,38,39}.
197 Yet, the estimated global proboscidean diversity by the end of this first phase was still around 23
198 species, although a third of which were endemic island dwarfs⁴⁰. A second phase of
199 proboscidean demise is recovered in Eurasia and the Americas, starting ca. 160 and 75 kya,
200 respectively (Fig. 1E). These extinction shifts represent a 24- and 16-fold increase in severity
201 with respect to Neogene basal levels in Eurasia and the Americas, respectively (Fig. 3a). The
202 already severely depleted mammutids and gomphotheres (PFTs 4 and 6) completely disappeared.
203 Proboscideans with the most proal shearing-adapted dentitions (PFTs 7 and 8)⁹, which had
204 weathered the first extinction wave, were hit hard too (Fig. 1c and 3b).

205 The cumulative evidence points to a major role of environmental forcing in the collapse of
206 proboscideans. Extinction severity in Africa and Eurasia during the last 10 Myrs track global
207 temperature trends, according to environment-dependent extinction models¹⁴ (Fig. 3c). Decrease
208 in average global temperature⁴¹ (measured through $\delta^{18}\text{O}$) and increase in the amplitude of
209 temperature fluctuations yielded a significant relationship with extinction rates. Our models
210 uncover continent-wide associations of extinction and environmental deterioration, but they do

211 not imply homogeneity in causality across lineages and smaller geographic scales. Plummeting
212 productivity and ecological reshuffling⁴² (including enhanced competition with other ungulate
213 clades that evolved specialist grazing ecomorphs) due to rapid Plio-Pleistocene climatic
214 fluctuations set a tight grip on proboscidean diversity limits, stemming from a sustained decrease
215 in the capacities of ecosystems to support diverse megaherbivore communities throughout the
216 Plio-Pleistocene³¹. Remarkably, the onset of the second, most acute extinction regime is not
217 recovered in Africa, and pre-dates the settlement of *Homo sapiens* in Eurasia and Americas (Fig.
218 3a). The final dismantling of proboscidean communities and the downfall of the clade was likely
219 initiated by the severe Holarctic climatic changes from the MIS 6 glaciation (ca. 190 kyr) to the
220 particularly harsh and cool climatic conditions of the MIS 4 stadial (ca. 74 kyr). Nevertheless,
221 our results do not contradict a scenario whereby anthropogenic pressure contributed to the
222 further catastrophic decline of proboscideans towards the end of the Pleistocene, as indicated by
223 the timing of extinction of many of the remaining taxa^{43,44}. Significantly, some of the shifts in
224 ecosystem functioning that triggered the decline of proboscideans were already in place before
225 they had an obvious impact on diversity trajectories, which highlights the importance of
226 integrating ecomorphological and local-scale approaches when establishing the causes of
227 megafauna declines.

228 In summary, the fossil evidence shows that proboscidean evolution was a complex and
229 continuous innovative process that allowed this group to overcome 60 million years of severe
230 environmental shifts. The high diversification rate estimated for the clade is consistent with the
231 notion that *K* strategies should promote speciation by ensuring the persistence of marginal
232 populations in new environments²¹. Increasingly resource-efficient ecomorphological novelties
233 would have complemented such strategies, enhancing species persistence and speciation, and in
234 turn propelling ecological innovation. Proboscideans were far from a case of lineage senescence
235 when their demise started, showing that large branches of the tree of life may vanish long before
236 exhausting their evolutionary potential^{7,45}.

237

238

239

240 **References**

- 241
- 242 1. Surovell, T., Waguespack, N. & Brantingham, P. J. Global archaeological evidence for
243 proboscidean overkill. *Proc National Acad Sci USA* **102**, 6231–6236 (2005).
- 244 2. Smith, F. A., Smith, R. E. E., Lyons, S. K. & Payne, J. L. Body size downgrading of mammals
245 over the late Quaternary. *Science* **360**, 310–313 (2018).
- 246 3. Faith, J. T., Rowan, J., Du, A. & Barr, W. A. The uncertain case for human-driven extinctions
247 prior to *Homo sapiens*. *Quaternary Res* **96**, 88–104 (2020).
- 248 4. Cuvier, G. Mémoires sur les espèces d'éléphants vivants et fossiles. *Mémoires de l'Institut des*
249 *Sciences et Arts* **2**, 1–22 (1800).
- 250 5. Osborn, H. F. The Ancestral Tree of the Proboscidea. Discovery, Evolution, Migration and
251 Extinction Over a 50,000,000 Year Period. *Proc Nat Acad Sci USA* **21**, 404–412 (1935).
- 252 6. IUCN. The IUCN Red List of Threatened Species. Version 2021-1.
253 <https://www.iucnredlist.org> (2021).
- 254 7. Maglio, V. J. Origin and Evolution of the Elephantidae. *T Am Philos Soc* **63**, 1 (1973).
- 255 8. Zhang, H., Wang, Y., Janis, C. M., Goodall, R. H. & Purnell, M. A. An examination of
256 feeding ecology in Pleistocene proboscideans from southern China (*Sinomastodon*, *Stegodon*,
257 *Elephas*), by means of dental microwear texture analysis. *Quatern Int* **445**, 60–70 (2017).
- 258 9. Saegusa, H. Stegodontidae and *Anancus*: Keys to understanding dental evolution in
259 Elephantidae. *Quaternary Sci Rev* **231**, 106176 (2020).
- 260 10. Hempson, G. P., Archibald, S. & Bond, W. J. A continent-wide assessment of the form and
261 intensity of large mammal herbivory in Africa. *Science* **350**, 1056–1061 (2015).
- 262 11. Silvestro, D., Salamin, N., Antonelli, A. & Meyer, X. Improved estimation of
263 macroevolutionary rates from fossil data using a Bayesian framework. *Paleobiol* **45**, 546–570
264 (2019).
- 265 12. Clavel, J., Escarguel, G. & Merceron, G. mvMORPH: an R package for fitting multivariate
266 evolutionary models to morphometric data. *Methods Ecol Evol* **6**, 1311–1319 (2015).
- 267 13. Cantalapiedra, J. L., Fernández, M. H., Azanza, B. & Morales, J. Congruent phylogenetic
268 and fossil signatures of mammalian diversification dynamics driven by Tertiary abiotic change.
269 *Evolution* **69**, 2941–2953 (2015).
- 270 14. Silvestro, D., Antonelli, A., Salamin, N. & Quental, T. B. The role of clade competition in
271 the diversification of North American canids. *Proc Nat Acad Sci USA* **112**, 8684–8689 (2015).

- 272 15. Sepkoski, J. J. A kinetic model of Phanerozoic taxonomic diversity I. Analysis of marine
273 orders. *Paleobiology* **4**, 223–251 (1978).
- 274 16. Tassy, P. The Proboscidean Datum Event: how many proboscideans and how many events?
275 in *European Neogene Mammal Chronology 237–252* (Plenus Press, 1989). doi:10.1007/978-1-
276 4899-2513-8_16.
- 277 17. Made, J. van der. The evolution of the elephants and their relatives in the context of changing
278 climate and geography. in *Elefantentreich - Eine Fossilwelt in Europa* (ed. Meller, H.)
279 (Landesamt für Denkmalpflege und Archäologie Sachsen-Anhalt - Landesmuseum für
280 Vorgeschichte, 2010).
- 281 18. Saarinen, J. J. *et al.* Patterns of maximum body size evolution in Cenozoic land mammals:
282 eco-evolutionary processes and abiotic forcing. *Proc Royal Soc B* **281**, 20132049 (2014).
- 283 19. Fortelius, M. *et al.* Evolution of Neogene Mammals in Eurasia: Environmental Forcing and
284 Biotic Interactions. *Annu Rev Earth Pl Sc* **42**, 579–604 (2014).
- 285 20. Marshall, C. R. & Quental, T. B. The uncertain role of diversity dependence in species
286 diversification and the need to incorporate time-varying carrying capacities. *Phil Trans Roy Soc*
287 *B* **371**, 20150217–13 (2016).
- 288 21. Vrba, E. S. Evolution, species and fossils: how does life evolve? *S. African J. Sci.* **76**, 61–84
289 (1980).
- 290 22. Cantalapiedra, J. L., Prado, J. L., Fernández, M. H. & Alberdi, M. T. Decoupled
291 ecomorphological evolution and diversification in Neogene-Quaternary horses. *Science* **355**,
292 627–630 (2017).
- 293 23. Calandra, I., Göhlich, U. B. & Merceron, G. How could sympatric megaherbivores coexist?
294 Example of niche partitioning within a proboscidean community from the Miocene of Europe.
295 *Naturwissenschaften* **95**, 831–838 (2008).
- 296 24. Sanders, W. J. Proboscidea from Kanapoi, Kenya. *J Hum Evol* **140**, 102547 (2020).
- 297 25. Wang, S. *et al.* Evolution of Protanancus (Proboscidea, Mammalia) in East Asia. *J Vertebr*
298 *Paleontol* **35**, e881830 (2015).
- 299 26. Lister, A. M. The role of behaviour in adaptive morphological evolution of African
300 proboscideans. *Nature* **500**, 331–334 (2013).
- 301 27. Lister, A. M., Sher, A. V., Essen, H. van & Wei, G. The pattern and process of mammoth
302 evolution in Eurasia. *Quatern Int* **126**, 49–64 (2005).

- 303 28. Wei, G. *et al.* New materials of the steppe mammoth, *Mammuthus trogontherii*, with
304 discussion on the origin and evolutionary patterns of mammoths. *Sci China Earth Sci* **53**, 956–
305 963 (2010).
- 306 29. Stanley, S. M. *Macroevolution, patterns and processes*. (W.H. Freeman and Company,
307 1979).
- 308 30. Edwards, E. J. *et al.* The Origins of C4 Grasslands: Integrating Evolutionary and Ecosystem
309 Science. *Science* **328**, 587–591 (2010).
- 310 31. Faith, J. T., Rowan, J., Du, A. & Koch, P. L. Plio-Pleistocene decline of African
311 megaherbivores: No evidence for ancient hominin impacts. *Science* **362**, 938–941 (2018).
- 312 32. Kaya, F. *et al.* The rise and fall of the Old World savannah fauna and the origins of the
313 African savannah biome. *Nat Ecol Evol* **2**, 241–246 (2018).
- 314 33. Saarinen, J. & Lister, A. M. Dental mesowear reflects local vegetation and niche separation
315 in Pleistocene proboscideans from Britain. *J Quaternary Sci* **31**, 799–808 (2016).
- 316 34. Rivals, F., Semperebon, G. M. & Lister, A. M. Feeding traits and dietary variation in
317 Pleistocene proboscideans: A tooth microwear review. *Quaternary Sci Rev* **219**, 145–153 (2019).
- 318 35. Vrba, E. S. Evolutionary Pattern and Process in the Sister-Group Alcelaphini-Aepycerotini
319 (Mammalia: Bovidae). in *Living Fossils* (eds. Eldredge, N. & Stanley, S. M.) 62–79 (Springer,
320 1984).
- 321 36. Herrera-Flores, J. A., Stubbs, T. L. & Benton, M. J. Macroevolutionary patterns in
322 Rhynchocephalia: is the tuatara (*Sphenodon punctatus*) a living fossil? *Palaeontology* **60**, 319–
323 328 (2017).
- 324 37. Todd, N. E. Trends in Proboscidean Diversity in the African Cenozoic. *J Mamm Evol* **13**, 1–
325 10 (2006).
- 326 38. Rivals, F., Mol, D., Lacombat, F., Lister, A. M. & Semperebon, G. M. Resource partitioning
327 and niche separation between mammoths (*Mammuthus rumanus* and *Mammuthus meridionalis*)
328 and gomphotheres (*Anancus arvernensis*) in the Early Pleistocene of Europe. *Quatern Int* **379**,
329 164–170 (2015).
- 330 39. Sanders, W. J. & Haile-Selassie, Y. A New Assemblage of Mid-Pliocene Proboscideans from
331 the Woranso-Mille Area, Afar Region, Ethiopia: Taxonomic, Evolutionary, and Paleoecological
332 Considerations. *J Mamm Evol* **19**, 105–128 (2012).
- 333 40. Geer, A. A. E. *et al.* The effect of area and isolation on insular dwarf proboscideans. *J*
334 *Biogeogr* **43**, 1656–1666 (2016).

- 335 41. Westerhold, T. *et al.* An astronomically dated record of Earth's climate and its predictability
336 over the last 66 million years. *Science* **369**, 1383–1387 (2020).
- 337 42. Vrba, E. S. Habitat Theory in Relation to the Evolution in African Neogene Biota and
338 Hominids. in *African Biogeography, Climate Change, and Hominid Evolution* (eds. Bromage, T.
339 G. & Shrenk, F.) 19–39 (1999).
- 340 43. Stuart, A. J. Late Quaternary megafaunal extinctions on the continents: a short review. *Geol J*
341 **50**, 338–363 (2015).
- 342 44. Jukar, A. M., Lyons, S. K., Wagner, P. J. & Uhen, M. D. Late Quaternary extinctions in the
343 Indian Subcontinent. *Palaeogeogr Palaeoclim Palaeoecol* **562**, 110137 (2021).
- 344 45. Raup, D. M. *Extinction: Bad Genes or Bad Luck?* (Norton, 1991).
- 345 46. Cantalapiedra, J. L. *et al.* Conserving evolutionary history does not result in greater diversity
346 over geological time scales. *Proc Royal Soc B* **286**, 20182896 (2019).
- 347 47. Ronquist, F. *et al.* MrBayes 3.2: Efficient Bayesian Phylogenetic Inference and Model
348 Choice Across a Large Model Space. *Systematic Biol* **61**, 539–542 (2012).
- 349 48. Hublin, J.-J. *et al.* New fossils from Jebel Irhoud, Morocco and the pan-African origin of
350 *Homo sapiens*. *Nature* **546**, 289–292 (2017).
- 351 49. O'Connell, J. F. *et al.* When did *Homo sapiens* first reach Southeast Asia and Sahul? *Proc*
352 *National Acad Sci* **115**, 8482–8490 (2018).
- 353 50. Ardelean, C. F. *et al.* Evidence of human occupation in Mexico around the Last Glacial
354 Maximum. *Nature* **584**, 87–92 (2020).
- 355 51. Paradis, E. *Analysis of Phylogenetics and Evolution with R (Second Edition)*. (Springer,
356 2012).
- 357 52. Revell, L. J. phytools: an R package for phylogenetic comparative biology (and other things).
358 *Methods Ecol Evol* **3**, 217–223 (2012).
- 359 53. Bates, D., Mächler, M., Bolker, B. & Walker, S. Fitting Linear Mixed-Effects Models Using
360 lme4. *J Stat Softw* **67**, (2015).
- 361 54. MacLatchy, L. M., Desilva, J., Sanders, W. J. & Wood, B. Hominini. in *Cenozoic Mammals*
362 *of Africa* (eds. Werdelin, L. & Sanders, W. J.) 471–545 (University of California Press, 2010).
- 363
- 364
- 365

366 **Methods**

367 **Occurrence dataset**

368 We gathered a species-level occurrence dataset for proboscideans drawn from the literature and
369 online databases. We reviewed the dataset for taxonomic rigour and used the literature to
370 maximise the precision of spatiotemporal occurrence data. Since occurrences with broad
371 temporal ranges flatten diversification rates, we excluded occurrences with broad temporal
372 ranges, unless this significantly decimated the occurrence available for a given taxon (see details
373 in Supplementary Methods).

374 We retain 1,427 records and 129 species from the NOW database
375 (<https://nowdatabase.org/now/database>) and 626 records and 33 species from the PBDB
376 (<https://paleobiodb.org>). A further 73 occurrences for 37 species were added based on an
377 assortment of other published studies. 22 of these species were not originally included in NOW
378 and PBDB, resulting in a 12% increment of the total diversity studied. The final dataset includes
379 2,130 occurrences for 185 proboscidean species, with an average temporal precision of 1.06 Myr
380 (median 0.8 Myr; 95% CI: 0.002675 — 3.7 Myr).

381 **Phylogenetic dataset**

382 We draw extensively from the pre-existing systematic palaeontological literature, to build an
383 informal proboscidean supertree that incorporates current understanding of their phylogenetic
384 relationships, but at the same time acknowledges the uncertainty —both topological and
385 temporal— derived from all the available sources.

386 The baseline topology included polytomies where branching relationships between species
387 remain uncertain. We combined this topology with the oldest credible occurrence of each
388 species, and used a tip-dating procedure⁴⁶ implemented in MrBayes⁴⁷ to generate a posterior
389 distribution of trees that are consistent with all the information available to date, while
390 accounting for temporal and topological uncertainties. From the posterior distribution, 100 trees
391 were retained for further analyses. Further details are provided in the Supplementary Methods.

392 **Functional ecology and proboscidean functional types (PFTs)**

393 Our aim is to capture the evolutionary patterns behind the rise and demise of functional diversity
394 (FD) of proboscideans over time. To delimit different proboscidean functional ecomorphs, we
395 consider functional traits that synthesise morphologies with a clear role in the interactions of
396 each species with its environment, condensing aspects like breadth of dietary preference and
397 feeding envelope, food processing, energy requirements, home range size, social grouping,
398 sexual selection and locomotion. A total of 17 discretised traits were defined and analysed: body
399 size, molar structure, number of transversal loph(id)s, hypsodonty, horisodonty, presence of
400 acute lophs, presence of obtuse lophs, occlusal topography, degree of enamel folding, skull
401 shape, morphology of upper tusks, morphology of lower tusks, mandible shape, symphyseal

402 morphology of mandible, pattern of mastication, dental replacement, and foot posture. An
403 extensive description of the traits, their sources, and their ecological relevance is presented in the
404 Supplementary Methods. We collected information to score these traits based on direct
405 observations by several of us (H.Z., M.T.A., J.L.P., and J.S.), and from the literature. The
406 categories of the 17 functional traits are summarised in the Extended Data Fig. 1.

407 We were able to gather comprehensive ecomorphological information for 180 species and
408 translated 93% of the observed disparity in a two-dimensional NMDS space (by means of a
409 Gower distance matrix; see Supplementary Methods). This allows us to reduce the potential
410 covariation in some of the traits, while making it easier to model ecomorphological evolution in
411 a phylogenetic framework and as part of ecology-dependent models. We also used the Gower
412 distance matrix to compute ecological disparity (measured as the sum of variances), and
413 extinction selectivity over time. Based on the NMDS space, we used *k-means* to identify 8
414 proboscidean functional types (PFTs). See detailed procedures in the Supplementary Methods. A
415 reconstruction of representative species of the eight PFTs is shown in Figure 1a.

416 **Speciation and extinction rates over time**

417 Detailed fossil occurrence datasets provide the opportunity to estimate speciation and extinction
418 rates through time while controlling for sampling biases. We applied Bayesian procedures
419 (implemented in the software *PyRate*) that analyse occurrence data to simultaneously estimate
420 preservation rates, origination and extinction ages for each species, as well as speciation and
421 extinction rates through time. Complexity in speciation and extinction rates trajectories is
422 modelled by identifying the number of significant rate shifts and their temporal assignment using
423 a reversible jump Markov Chain Monte Carlo (RJMCMC). Temporal ranges associated with
424 each occurrence data point are treated as dating uncertainties, with higher data uncertainty
425 yielding larger uncertainty around parameter estimates. Occurrence ages were randomly
426 resampled 20 times from the uniform distribution defined by their temporal ranges, generating 20
427 datasets. We run *PyRate* independently on these 20 datasets. The number of generations of the
428 MCMC chains was tailored for each dataset based on exploratory analyses. The MCMC run for
429 the global dataset (including NMDS correlations and a run focussed on the last 10 Myr) was run
430 for 10 million generations. The African, Eurasian and American datasets were run for 5, 7 and 5
431 million generations, respectively. We sampled every 1000 iterations, and discarded the 10% as
432 burning. The resulting samples obtained from each resampled dataset were enough to ensure ESS
433 above 200 in all key parameters. To report and visualise parameter estimates, 500 random
434 samples were kept from each run, resulting in a total of 10,000 samples from each analysis. Rate
435 shifts were allowed to take place once every million years (the default setting in *PyRate*, -
436 `min_dt 1`). We allowed preservation rates to shift between the Eocene and Oligocene (33.9 Ma),
437 between Aquitanian and the Burdigalian (20.44 Ma), and then for boundaries between these
438 periods: Burdigalian (15.97 Ma) / Serravallian (11.63 Ma) Tortonian / Messinian (5.333 Ma) /
439 Pliocene (2.58 Ma) / Pleistocene (0.0117 Ma) / Holocene. This definition of boundaries was

440 aimed at obtaining good posterior estimates of sampling rates and was based on exploratory
441 analyses of occurrence frequency and comparison between different shift configurations.

442 Detailed diversification profiles for the last 10 Myr were obtained by limiting the *PyRate*
443 analyses (`-edgeShift 10 0`) and allowing rate shifts to take place every 10 kyr (`-min_dt`
444 `0.01`), taking advantage of the finer record in Upper Miocene, Pliocene and Quaternary times.
445 The fine timing of extinction shifts for the last 10 Myr was compared to the earliest records of
446 *Homo sapiens* in each continent⁴⁸⁻⁵⁰ (see Fig. 3a). Importantly, *PyRate* estimates extinction times
447 for all the species, and thus our extinction-rate profiles are not based on a literal reading of last
448 appearance datum (LADs). Our approach is thus conservative regarding the comparison of the
449 timing of extinction shifts with the first records of *Homo sapiens*, since *PyRate* places the timing
450 of the last extinction regime closer to the appearance of modern humans than raw FADs would
451 do.

452 Finally, we computed diversity curves through time using *PyRate*-implemented functions that
453 combine estimates of sampling rates over time and occurrence distribution, yielding diversity
454 estimates that account for the shifting quality of the paleontological record through time.

455 **Effect of ecology on speciation and extinction**

456 Ecology may exert a very strong impact on diversification rates (speciation and extinction),
457 producing species sorting. To test for a role of ecological function on diversification in
458 Proboscidea, we drew on the possibility to run the Covar model in *PyRate*. Under this model,
459 diversification trajectories are estimated as described above, but each species is able to depart
460 from the overall baseline trend based on a correlation with a continuous variable. In our case, the
461 two axes of our NMDS space were tested for such a correlation. The resulting coefficients
462 (Extended Data Fig. 8) were used to estimate the departure from baseline speciation and
463 extinction for each species (Fig. 2 and Extended Data Fig. 9).

464 **Effect of palaeoclimate on extinction**

465 A major question regarding megaherbivore declines from the Late Miocene onwards is to what
466 extent were species extinctions the results of climatic perturbations. To quantitatively test for an
467 association between the decline of proboscideans (i.e. extinction rates) and palaeoclimate, we use
468 climate-dependent diversification models¹⁴ at the continental level. In order to avoid correlations
469 stemming from broad Cenozoic trends, we restricted our analyses interval to the last 10 Myr. As
470 a proxy for palaeoclimate, we use the recently published record of $\delta^{18}\text{O}$ variations in deep-sea
471 benthic foraminifera⁴¹. We computed two variables from the $\delta^{18}\text{O}$ data. We estimated the
472 average $\delta^{18}\text{O}$ score in 0.1 Myr bins, as a proxy for temperature (the original data comes with
473 points every 20 kyrs). We also test for a role of the severe short-term environmental fluctuations
474 that characterised the Plio-Pleistocene. To do so, we estimated the absolute of the first
475 differences of the 20 kyrs curve, and then averaged these values into 0.1 bins. Both climate-
476 dependent diversification models were run using the *PyRateContinuous.py* script in the *PyRate*

477 software¹¹ for 1,000,000 generations, sampled the Bayesian search every 1000 iterations, and
478 discarded the first 10% as burnin. Distribution of the correlation parameters for each continent
479 are shown in Figure 3c.

480 **Phylogenetic modelling**

481 We assessed modes of proboscidean ecomorphological evolution using phylogenetic modelling.
482 To do so, we applied the information yielded from the NMDS axes, which are continuous
483 representations of the many functional ecomorphological aspects of proboscideans. Given the
484 multidimensional nature of our data, we used multivariate phylogenetic models¹², which allow us
485 to jointly model the two dimensions of the NMDS space while accounting for the covariance of
486 the axes, as well as their different associated evolutionary rates and/or optimal values.

487 We fit two sets of models. In the first set, the entire clade is modelled together, and thus
488 represents rates and trends which are shared across all lineages. This set of models include
489 brownian motion (BM), early burst (EB), and Ornstein–Uhlenbeck (OU) stationary model,
490 representing unconstrained diffusion, decaying-rate scenario (e.g. adaptive radiations,
491 ecomorphological stagnation), and constrained or limited evolution, respectively.

492 In a second set of models, we explore the role of biogeography on the tempo and mode of
493 proboscidean functional evolution. In particular, we investigate the role of the Proboscidean
494 Datum Event (PDE), this is, the dispersal of proboscidean lineages beyond Afro-Arabia at
495 around 22 Ma. In this set of models, we partition the tree in lineages evolving in and outside
496 Africa using functions in the *ape*⁵¹ and *phytools*⁵² R packages (see Supplementary Methods), and
497 fit different combinations of the models described above to these two partitions. Further
498 procedural details of models and specific R functions are provided in the Supplementary
499 Methods.

500 All models were run on 100 trees. The best performing model is the EBBM (EB model in Africa;
501 BM model outside Africa). Parameter estimates and fit of the models are shown in Extended
502 Data Fig. 5.

503 **Trends in community-level proboscidean diversity**

504 We assessed the temporal trends of local species richness and the relationship of averaged dental
505 masticatory durability (DMD; mean NMDS-1 scores) in local richness using mixed-effects
506 models. We tested for the different trends in community richness in four different phases. The
507 African phase spans proboscidean communities before 22 Ma, an age that marks the
508 establishment of several lineages outside Africa (the Proboscidean Datum Event). The second
509 phase, called the after-PDE phase, spans from 22 to 8 Ma. The C₄ expansion phase, includes
510 communities between 8 and 2.4 Ma, reflecting the new ecological settings brought about
511 globally by this new type of grassland biome, which have been previously hypothesised to have a

512 major impact in megafauna communities³¹. The decline phase spans from 2.4 Ma to the present,
513 reflecting the interval where extinction intensified and drove the clade to its diversity decline.

514 Models where local species richness was a function of age, and where richness was a function of
515 mean NMDS-1 scores (dental masticatory durability, DMD, which was scaled from 0 to 1, to
516 ease interpretations; see Extended Data Fig. 7b) were evaluated while taking into account age
517 uncertainty and geography-dependent varying effects such as sampling heterogeneity and
518 taxonomic practices (see Supplementary Methods). Mixed effects models were run using the
519 function *lmer* in the R package *lme4*⁵³. Corrected Akaike Information Criterion (AICc) scores
520 show that the different phases have an effect both on the slope and the intercepts of the
521 relationships, both for age and DMD models (AICc Weights are 1 in both cases suggesting a
522 strong preference for the most complex models). The parameters of the best models for each
523 variable are provided in Extended Data Fig. 6. A representation of the best models is shown in
524 Extended Data Fig. 7.

525

526

527 **Methods references**

528 46. Cantalapiedra, J. L. *et al.* Conserving evolutionary history does not result in greater diversity
529 over geological time scales. *Proc Royal Soc B* **286**, 20182896 (2019).

530 47. Ronquist, F. *et al.* MrBayes 3.2: Efficient Bayesian Phylogenetic Inference and Model
531 Choice Across a Large Model Space. *Systematic Biol* **61**, 539–542 (2012).

532 48. Hublin, J.-J. *et al.* New fossils from Jebel Irhoud, Morocco and the pan-African origin of
533 *Homo sapiens*. *Nature* **546**, 289–292 (2017).

534 49. O’Connell, J. F. *et al.* When did *Homo sapiens* first reach Southeast Asia and Sahul? *Proc*
535 *National Acad Sci* **115**, 8482–8490 (2018).

536 50. Ardelean, C. F. *et al.* Evidence of human occupation in Mexico around the Last Glacial
537 Maximum. *Nature* **584**, 87–92 (2020).

538 51. Paradis, E. *Analysis of Phylogenetics and Evolution with R (Second Edition)*. (Springer,
539 2012).

540 52. Revell, L. J. phytools: an R package for phylogenetic comparative biology (and other things).
541 *Methods Ecol Evol* **3**, 217–223 (2012).

542 53. Bates, D., Mächler, M., Bolker, B. & Walker, S. Fitting Linear Mixed-Effects Models Using
543 *lme4*. *J Stat Softw* **67**, (2015).

544 54. MacLatchy, L. M., Desilva, J., Sanders, W. J. & Wood, B. Hominini. in *Cenozoic Mammals*
545 *of Africa* (eds. Werdelin, L. & Sanders, W. J.) 471–545 (University of California Press, 2010).

546

547 **Acknowledgements:** We thank D. Silvestro, J. Calatayud and J. Clavel for technical assistance,
548 and F. Bibi, M.J. Benton and W.J. Sanders for helpful comments. S. Wang and G. van den Bergh
549 provided important constructive comments on the informal supertree. Gratitudes are extended to
550 E.M. Dunne, D. Mothé, F. Rivals and two other anonymous reviewers, whose comments and
551 recommendations improved the presentation of this paper. This is PBDB publication #398.

552 **Funding:** This work was supported by the Talent Attraction Program of the Madrid Government
553 (2017-T1/AMB5298), the German Research Foundation (DFG, AOBJ: 637491), and the
554 Academy of Finland (post-doctoral research fund nr. n:o 315691).

555 **Author contributions:** J.L.C., O.S. and J.S. conceptualised the research. J.L.C, O.S., J.S., H.Z.,
556 M.T.A. and J.L.P. gathered the data. J.L.C., O.S. and F.B. designed and performed the analysis.
557 J.S. wrote the description and relevance of functional traits. J.L.C. and O.S. wrote the first
558 version of the paper, and H.Z. and J.S. contributed to the final version (main text and
559 supplement).

560 **Competing interests:** Authors declare no competing interest.

561 **Data and materials availability:** Datasets are available in the Supplementary Data file 1 and
562 *FigShare* (doi: 10.6084/m9.figshare.14035109).

563 **Code availability:** *PyRate* is a Python v.3. based program available at
564 <https://github.com/dsilvestro/PyRate>. Computation in R used functions in packages *mvMORPH*,
565 *phytools*, *ape* and *lme4*. Input files can be obtained from Supplementary Data file 1 and from
566 *FigShare* (doi: 10.6084/m9.figshare.14035109).

567

568

569

570

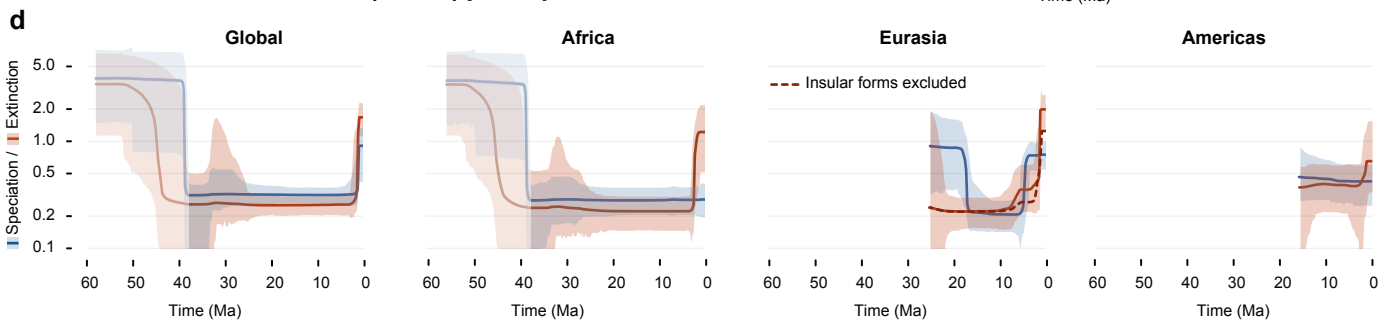
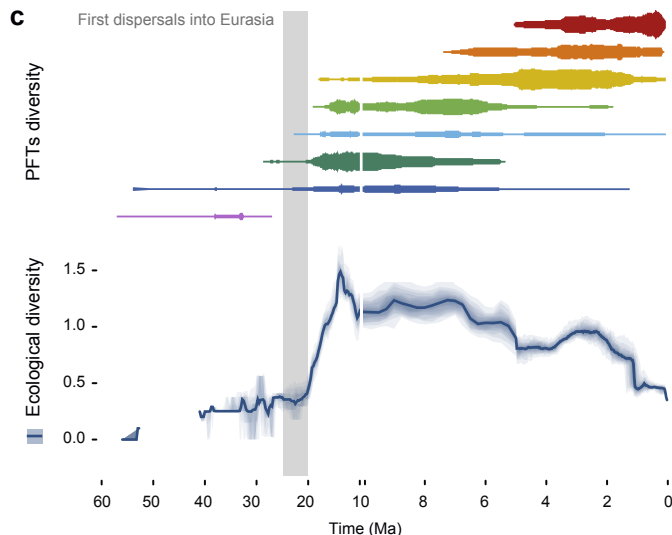
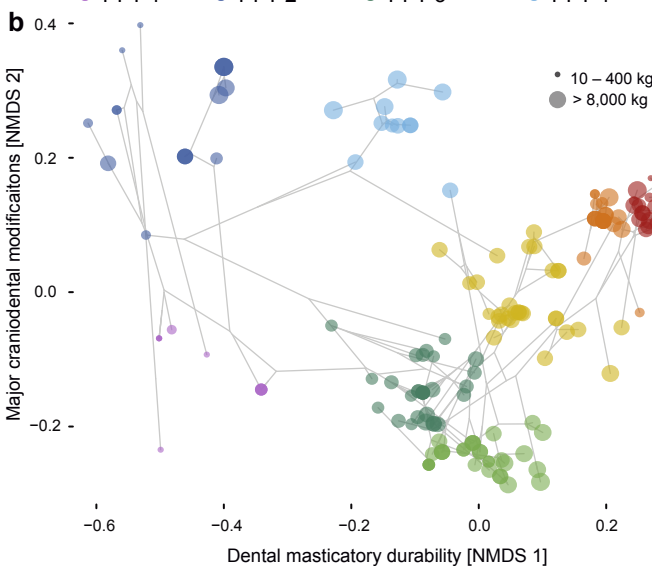
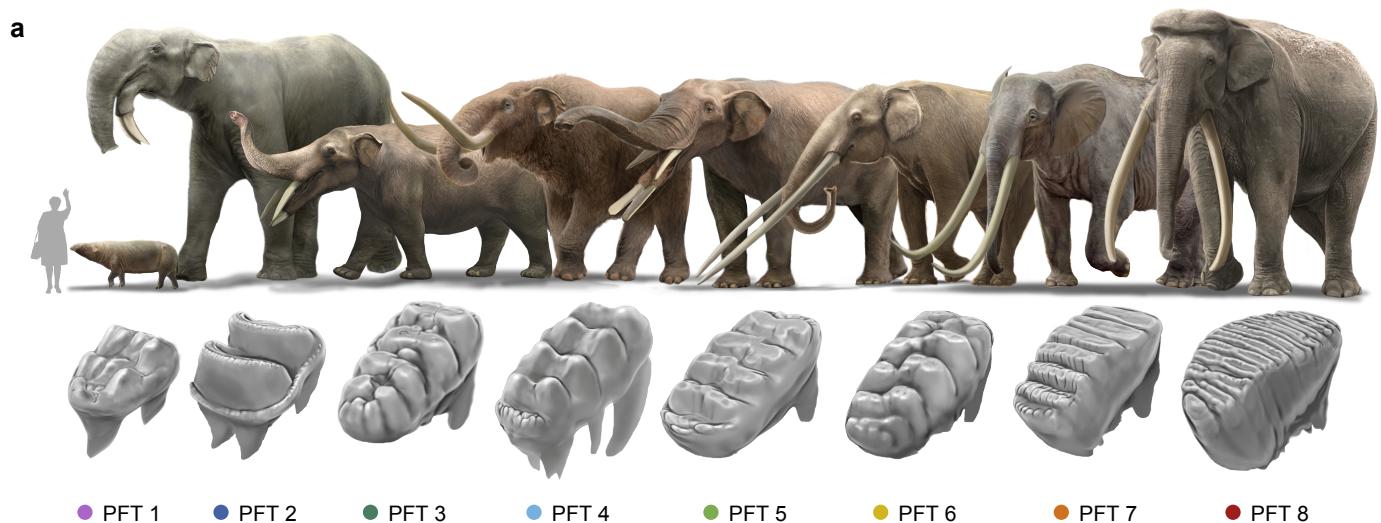
571

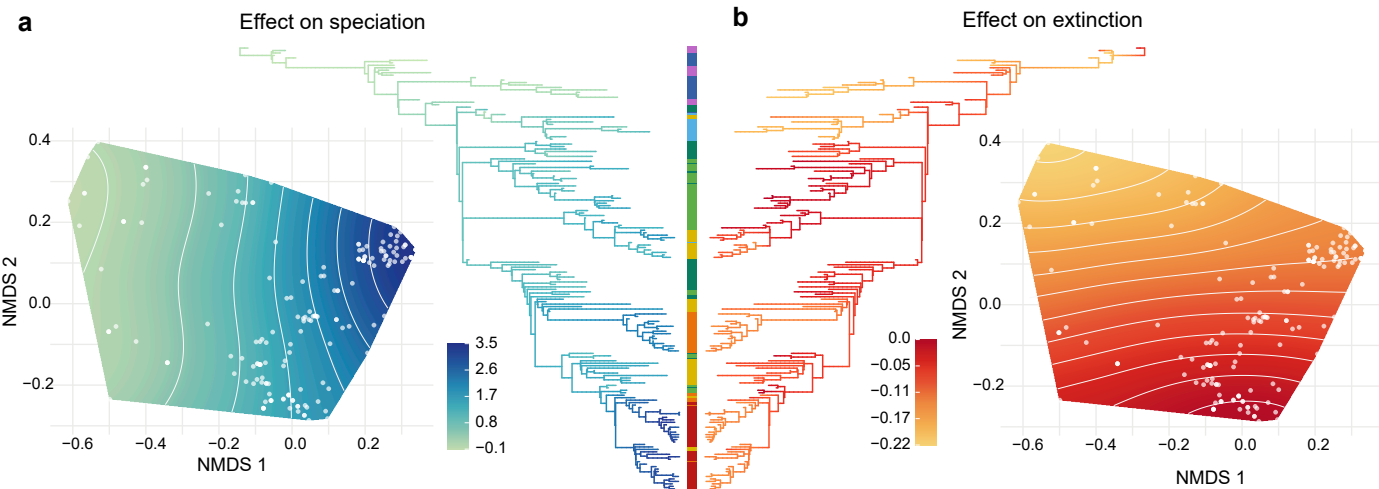
572 **Figure 1. Ecological diversity and diversification in proboscideans. a,**
 573 Reconstruction of representative genera for the eight proboscidean functional types
 574 (PFTs), including their upper third molars (molars not shown to scale). **PFT 1**
 575 *Moeritherium* – sublophodont molar with distinct cusps; **PFT 2** *Deinotherium* -
 576 bilophodont molar with crenulated ridges; **PFT 3** *Gomphotherium* - longirostrine
 577 mandible with pronounced lower incisors, basic bunodont molar with rounded cusps;
 578 **PFT 4** *Mammut* - brevirostrine mandible with lower incisors extremely vestigial or
 579 absent; zygodont molar (with ridged cusps); **PFT 5** *Amebelodon* - longirostrine mandible
 580 with shovel-shaped lower incisors, bunolamellar molar; **PFT 6** *Anancus* - brevirostrine
 581 mandible with lower incisors lost, complex bunodont molar achieving enhanced DMD;
 582 **PFT 7** *Stegodon* - brevirostrine mandible lacking lower incisors, brachydont (low-
 583 crowned) proal shearing molar comprising of numerous lamellae; **PFT 8** *Palaeoloxodon*
 584 - brevirostrine mandible lacking lower incisors, hypsodont (high-crowned) proal
 585 shearing molar comprising of numerous lamellae adjoined together by cementum. **b,**
 586 Two-dimensional functional space with colour-coded assignation to PFTs based on our
 587 17 ecomorphological traits. **c,** species diversity of the PTFs through time (log-scaled),
 588 showing the timing of the PDE (grey bar), and ecological disparity, measured as the
 589 sum of variances. **d,** Global and continental speciation and extinction rates through
 590 time; rates are log-scaled; shaded areas represent 95% credible intervals. Artwork by
 591 O.S..

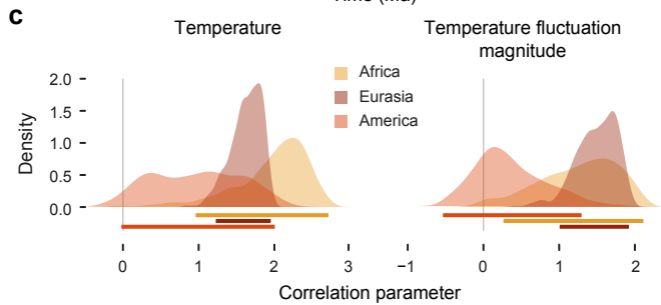
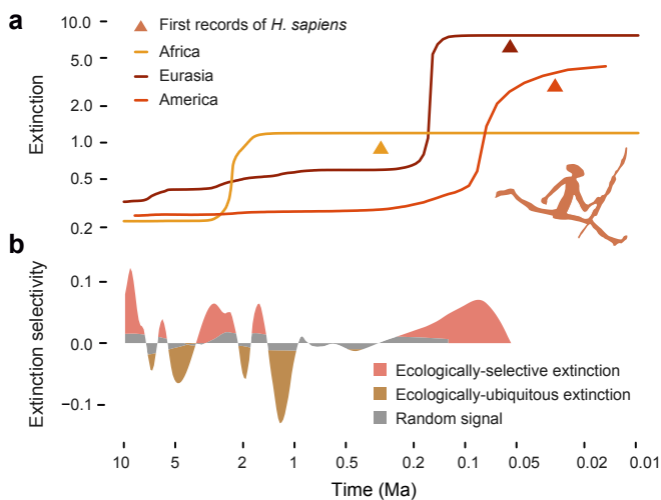
592
 593 **Figure 2. Effect of ecomorphology on diversification.** Ecomorphology-predicted
 594 departure of speciation (**a**), and extinction (**b**). These are derived from the correlations
 595 of rates with NMDS axes. Based on the NMDS scores of each species, we can plot their
 596 expected departure from baseline rates on the NMDS space and the proboscidean
 597 phylogeny (see also Extended Data Fig. 8 and 9). The central vertical bar represents
 598 PFT assignment.

599 **Figure 3. Extinction trends in the last 10 Myr. a,** Continental extinction rates through
 600 time on log-scaled axes, where triangles mark the timings at which earlier presence of
 601 *Homo sapiens* sensu lato⁵⁴ became established on the respective landmasses (Africa 315

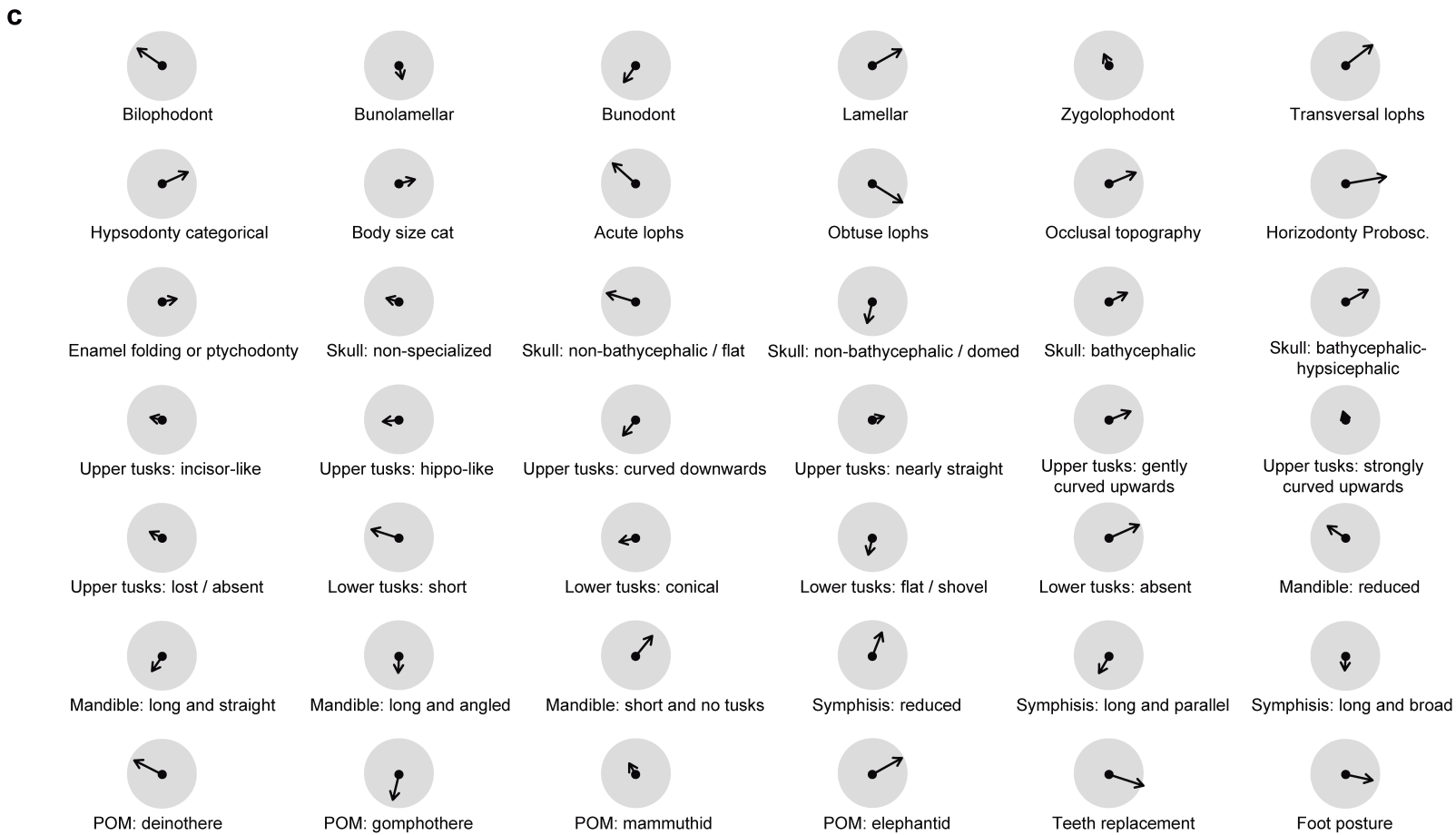
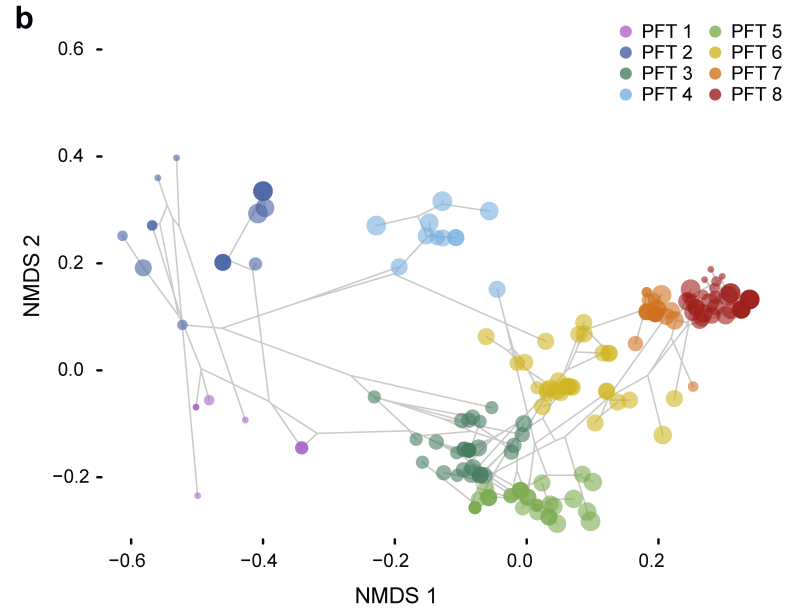
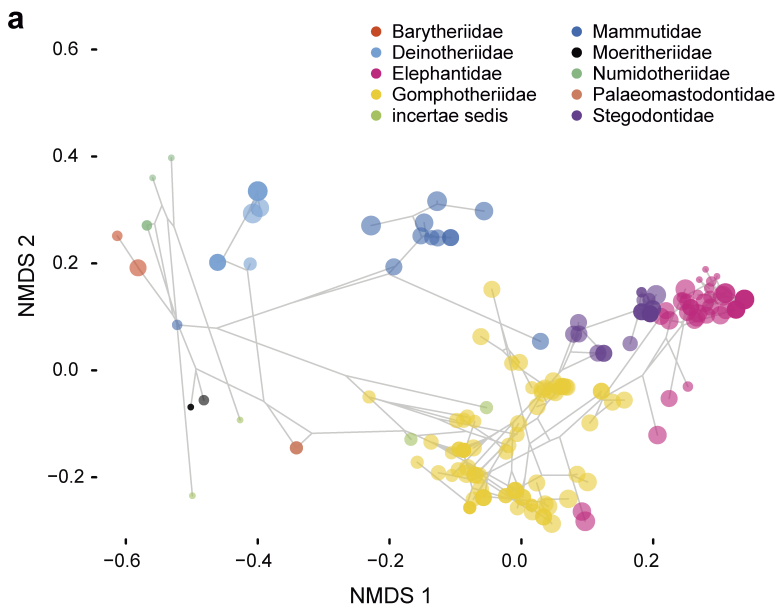
602 kya; Eurasia 55 kya; Americas 30 kya) ⁴⁸⁻⁵⁰. **b**, Extinction selectivity shows whether
603 extinction is ecologically-restricted (positive values) or hits on broad regions of the
604 functional space (negative values); the grey band represents values consistent with a
605 random signal (non-significant *P*-values). **c**, posterior distributions of correlation
606 parameters of extinction with paleotemperature trends for each continent; horizontal
607 bars show the 95 highest probability density.

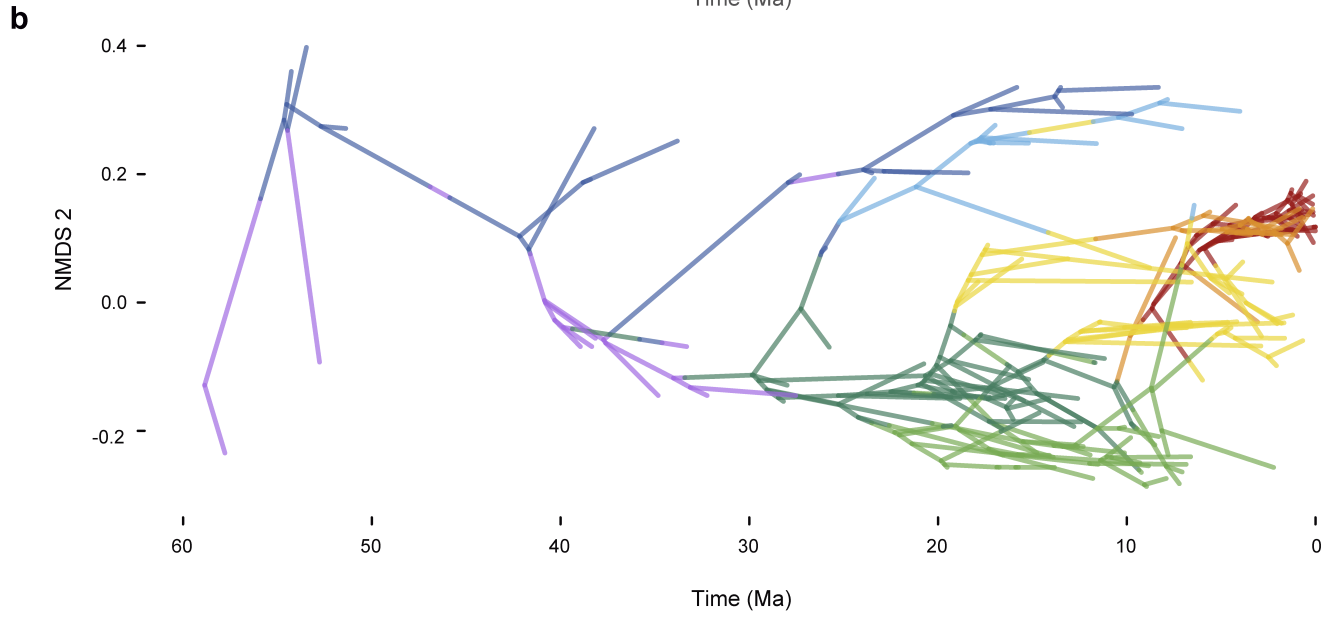
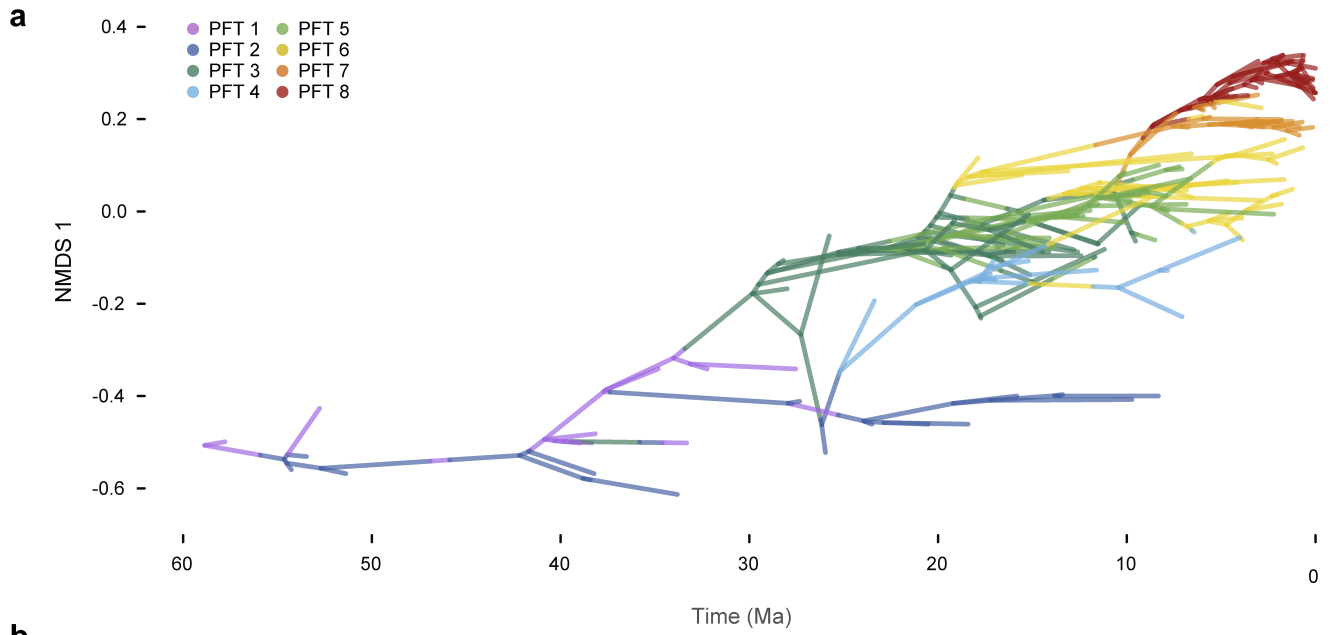


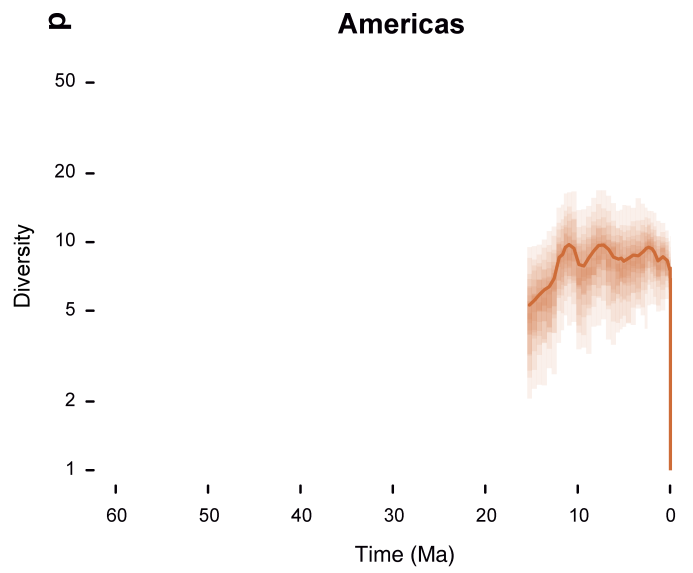
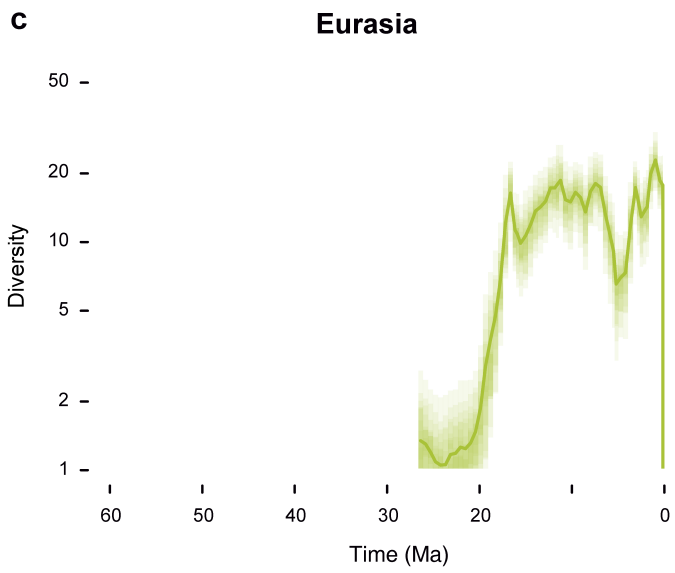
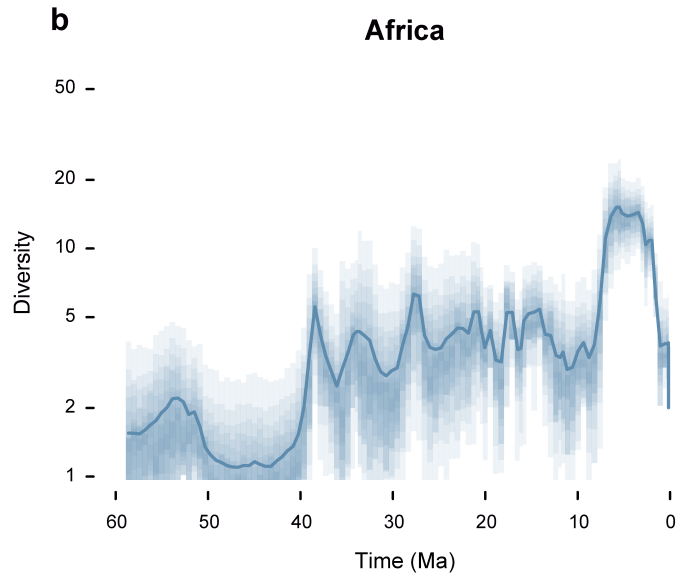
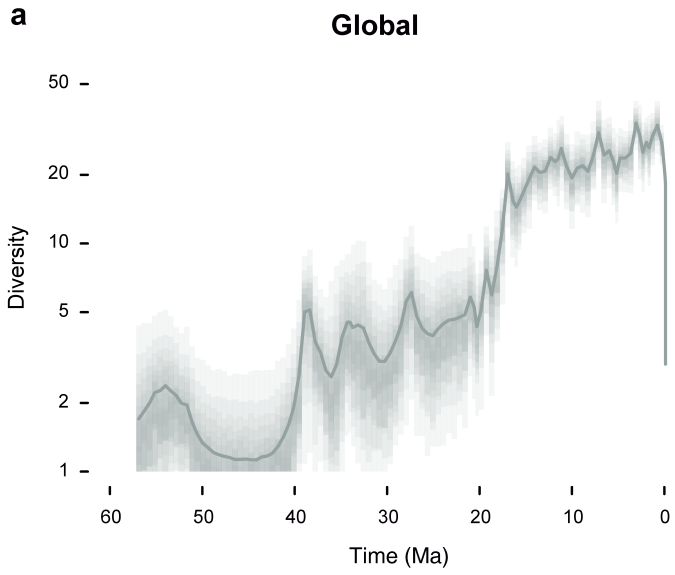




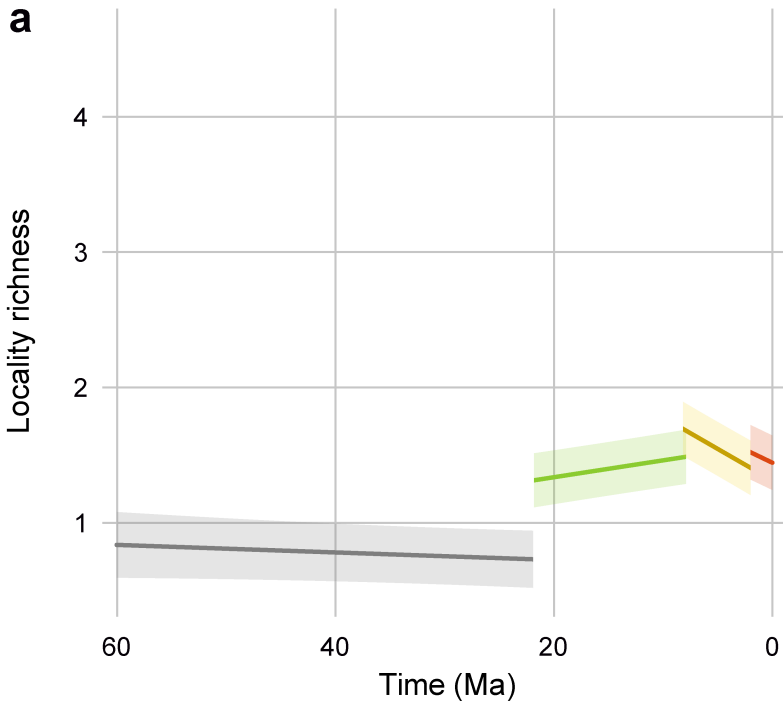
Trait	levels	order
tooth type	1: bunodont 2: bilophodont 3: buno-lamellar 4: lamellar 5: zygodont	unordered
number of transversal lophes	0: 0 lophes 1: 2 lophes 2: 3 lophes 3: 4 - 5 lophes 4: 6 - 7 lophes 5: 8 - 9 lophes 6: 10 - 14 lophes 7: more than 14 lophes	ordered
presence of acute lophes	0: absent 1: present	unordered
presence of obtuse lophes	0: absent 1: present	unordered
occlusal topography	0: raised elements 1: flat	unordered
hypsodonty	0: brachyodont 1: mesodont 2: hypsodont	ordered
horizodonty	1: 2 cusps pairs 2: 3 cusps pairs 3: 4 - 5 cusps pairs 4: 6 - 7 cusps pairs 5: 8 - 9 cusps pairs 6: 10 - 14 cusps pairs 7: more than 14 cusps pairs	ordered
enamel folding	0: absent 1: present	unordered
body size	1: 10-400 kg 2: 400 - 1000 kg 3: 1000 - 2000 kg 4: 2000 - 3000 kg 5: 3000 - 5000 kg 6: 5000 - 8000 kg 7: > 8000 kg	ordered
skull shape	0: non-specialised (e.g. <i>Eritherium</i> , <i>Phosphatherium</i>) 1: non-bathycephalic, flat (e.g. <i>Deinotheriidae</i> , <i>Palaeomastodon</i> , <i>Choerolophodon</i>) 2: non-bathycephalic, domed (e.g. <i>Gomphotherium</i> , <i>Afrochoerodon</i>) 3: bathycephalic, non-hypsicephalic (e.g. <i>Loxodonta</i> , <i>Stegodon</i>) 4: bathycephalic, hypsicephalic (e.g. <i>Elephas</i> , <i>Mammuthus</i>)	unordered
upper tusks	0: I2s with generalised incisor-like morphology (e.g. <i>Phosphatherium</i>) 1: I2s hippo-like (e.g. <i>Moeritherium</i>) 2: down-curved (e.g. <i>Gomphotherium</i> , <i>Platybelodon</i> , <i>Eozygodon</i>) 3: nearly straight, very little down or up-curve (e.g. <i>Cuvieronius</i> , <i>Anancus</i> , <i>Stegotetrabelodon</i>) 4: gentle up-curve (e.g. <i>Mammuthus americanum</i> , <i>Loxodonta</i> , <i>Elephas</i>) 5: very strong up-curve (e.g. <i>Mammuthus</i> , <i>Choerolophodon corrugatus</i>) 6: lost/absent, no evidence of functional role (<i>Prodeinotherium hobleiyi</i>)	unordered
lower tusks	0: short, i2 always present (e.g. <i>Phosphatherium</i> , <i>Moeritherium</i>) 1: developed, conical or oval in section, straight or curved (e.g. <i>Omanitherium</i> , <i>Gomphotherium</i> , <i>Deinotherium</i>) 2: developed and flattened, shovel-like (e.g. <i>Amebelodon</i> , <i>Platybelodon</i>) 3: absent or very reduced (e.g. <i>Mammuthus</i> , <i>Primelephas</i> , <i>Loxodonta</i>).	unordered
mandible	0: short with great angle of ventral deflection (<i>Prodenotherium</i> , <i>Deinotherium</i>) 1: long and straight (relative to the occlusal border; e.g. <i>Gomphotherium</i> , <i>Konobelodon</i>) 2: long with great angle of ventral deflection (e.g. <i>Stegotetrabelodon</i> , <i>Tetralophodon</i>) 3: short and with missing or reduced lower tusks (e.g. <i>Mammuthus</i> , <i>Elephas</i>)	unordered
mandibular symphysis	0: reduced, with or without incisors (earliest taxa, <i>Elephas</i>) 1: long, with parallel borders (e.g. <i>Konobelodon</i> , <i>Tetralophodon</i> , <i>Gomphotherium stenheimense</i> , <i>Deinotherium</i>) 2: long and rostrally broadened (e.g. <i>Amebelodon</i> , <i>Protanancus</i> , <i>Platybelodon</i> , <i>Gomphotherium angustidens</i>)	unordered
pattern of mastication (POM)	0: Deinotherium-pattern of mastication (e.g. <i>Deinotherium</i> , <i>Numidotherium</i> , <i>Phosphatherium</i>) 1: Gomphotherium-pattern of mastication (e.g. <i>Gomphotherium</i>) 2: Mammuthian-pattern of mastication (e.g. <i>Mammuthus</i> , <i>Zygodont</i>) 3: Elephantian-pattern of mastication (e.g. <i>Elephas</i> , <i>Stegodon</i> , <i>Mammuthus</i>)	unordered
teeth replacement	0: P4/p4 present when M3/m3 in function 1: P4/p4 absent when M3/m3 in function.	unordered
foot posture	0: plantigrade 1: subunguligrade	unordered



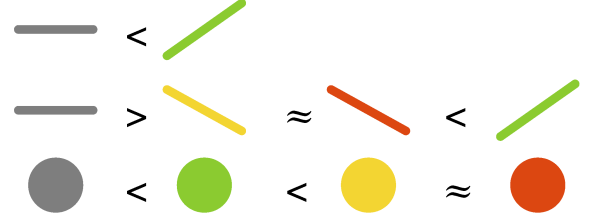




	phase	mean intercept	lower CI	upper CI	mean slope	lower CI	upper CI
richnes ~age*phase	African phase 60 - 22 Ma	0.687	0.459	0.916	-0.003	-0.007	0.001
	After PDE 22 - 8 Ma	1.505	1.304	1.706	0.013	0.008	0.017
	C ₄ expansion 8 - 2.4 Ma	1.615	1.417	1.814	-0.047	-0.055	-0.039
	Decline 2.4 - 0 Ma	1.694	1.490	1.899	-0.039	-0.046	-0.032
richnes ~mean_DMD*phase	African phase 60 - 22 Ma	0.723	0.486	0.959	-0.055	-0.232	0.122
	After PDE 22 - 8 Ma	1.399	1.183	1.615	0.209	0.175	0.243
	C ₄ expansion 8 - 2.4 Ma	1.538	1.323	1.754	-0.156	-0.193	-0.119
	Decline 2.4 - 0 Ma	1.617	1.400	1.834	-0.196	-0.225	-0.166

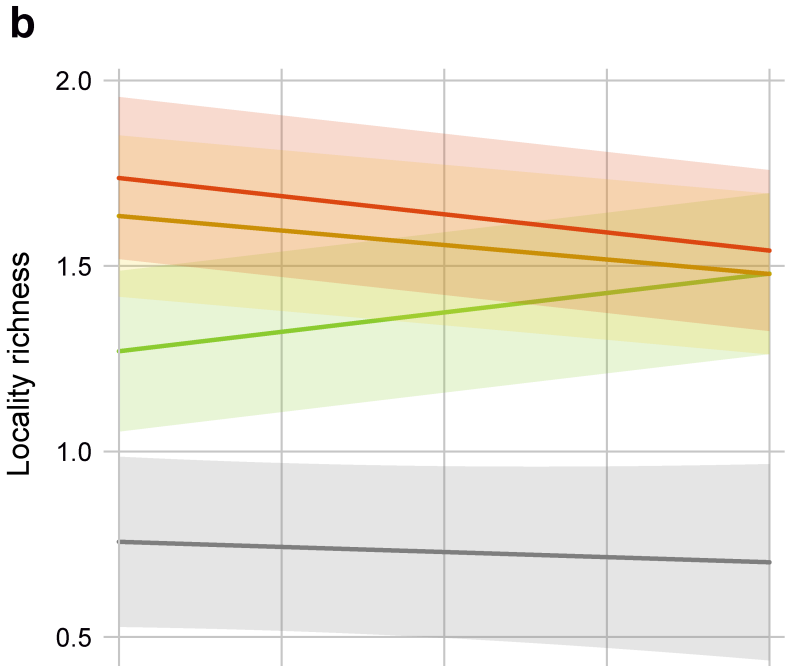


Richness ~ age * phase



Phase

- A: African phase (60 – 22 Ma)
- B: After Proboscidean Datum Event (22 – 8 Ma)
- C: C₄ Expansion phase (8 – 2.4 Ma)
- D: Decline phase (2.4 – 0 Ma)



Richness ~ Mean DMD * phase

