

1 **New data on morphological evolution and dietary adaptations of *Elephas recki* from**
2 **the Plio-Pleistocene Shungura Formation (Lower Omo Valley, Ethiopia)**

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5 **Nouvelles données concernant l'évolution morphologique et les adaptations**
6 **alimentaires d'*Elephas recki* de la formation plio-pléistocène de Shungura (basse vallée**
7 **de l'Omo, Éthiopie)**

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10 **Evolution of *Elephas recki* from the Shungura Formation**

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37 **ABSTRACT**

38 The ~~abundant and diverse~~ proboscideans, abundant and diverse throughout the Cenozoic, are
39 essential terrestrial megaherbivores for studying morphological adaptations and reconstructing
40 paleoenvironments in Africa. This new study of the lineage of *Elephas recki* (or "*Elephas recki*
41 complex") from the Plio-Pleistocene deposits of the Shungura Formation aimed at investigating
42 further the interplay between morphological and behavioral adaptations within proboscideans. For
43 this, we used the most complete and well-preserved third molars of the series for characterizing
44 morphological changes, while describing dietary behavior by applying mesowear angle analysis
45 to all identifiable cheek teeth. Biometric character changes indicated that evolutionary change was
46 not a gradual, progressive trend through the Shungura sequence deposition. Rather, we observed
47 a complex trend involving sudden changes. The mesowear angle analysis suggested that *Elephas*
48 *recki* from the Shungura Formation had consistently herbaceous monocot (such as grass and
49 sedges) dominated grazing dietary habits throughout its evolution, except for some slight
50 tendencies of mixed browse/graze diet in certain time intervals. We therefore did not observe any
51 correlation between morphological change and feeding preferences over two million years. This
52 delay between the acquisition of the observed diet and purportedly consequent morphological
53 changes questions the classical interpretations of factors at play in this evolutionary sequence.

54
55 **Keywords:** *Elephas recki*, Enamel thickness, Hypsodonty index, Mesowear, Shungura Formation

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56 **RÉSUMÉ**

57 Les proboscidiens, un groupe ~~abondant et diversifié~~ de mégaherbivores terrestres abondant et
58 diversifié durant le Cénozoïque, revêtent une grande importance dans l'étude des adaptations
59 morphologiques et dans la reconstruction des paléoenvironnements en Afrique. Cette nouvelle
60 étude de la lignée *Elephas recki* (ou "complexe *Elephas recki*") des dépôts plio-pléistocènes de la
61 Formation de Shungura a pour objectif d'investiguer plus avant les interrelations entre adaptations
62 morphologiques et adaptations comportementales au sein des proboscidiens. Pour cela, nous
63 utilisons les troisièmes molaires les plus complètes et les mieux préservées de cette série pour
64 caractériser les changements morphologiques, tout en décrivant le comportement alimentaire en
65 appliquant l'analyse angulaire des méso-usures à toutes les dents identifiables. Les variations des
66 caractères biométriques indiquent une évolution différant d'une tendance graduelle et progressive
67 pendant le dépôt de la séquence de Shungura. Au lieu de cela, nous observons une tendance
68 complexe impliquant des changements abrupts. L'analyse angulaire des méso-usures suggère
69 qu'*Elephas recki* de la Formation de Shungura avait un comportement alimentaire invariable de
70 paisseur (consommateur de monocotylédones herbacées telles que les graminées et les
71 cypéracées), à l'exception de légères tendances à une alimentation mixte paisseur/brouleur à
72 certains moments. Nous n'avons donc observé aucune corrélation entre changement
73 morphologique et préférences alimentaires durant plus de deux millions d'années. Ce décalage
74 entre l'acquisition du régime alimentaire observé et les changements morphologiques
75 supposément résultants remet en question les interprétations classiques concernant les facteurs à
76 l'œuvre dans cette séquence évolutionnaire.

77

78 Mots clefs : *Elephas recki*, Épaisseur de l'émail, Index d'hypsodontie, Méso-usures, Formation
79 de Shungura

80 INTRODUCTION

81

82 Proboscidea Illiger, 1811 is an order of mammals that includes elephants, the largest extant land
 83 animals on extant Earth. Their evolutionary history is well-documented and fossils indicate an
 84 incredible diversity (Shoshani & Tassy 2005; Gheerbrant & Tassy 2009; Cantalapieadra *et al.*
 85 2021). Most of their evolution occurred in Afro-Eurasia, with later migrations-dispersals to the
 86 Americas (Coppens 1978; Tassy 1988). The "true" elephants, i.e. the family Elephantidae,
 87 originated in Africa during the Late Miocene (Maglio 1973; Kalb *et al.* 1993), and include three
 88 emblematic lineages of extant and recently extinct crown elephantids (*Loxodonta*, *Elephas*, and
 89 *Mammuthus*). Following a steep decline during the last two million years (Cantalapieadra *et al.*
 90 2021), and despite some successful diversifications (*Mammuthus*, *Palaeoloxodon*) in northern
 91 Eurasia, there are currently only three species of elephants in the world: the Asian elephant,
 92 *Elephas maximus*, and two species found in Africa, the savanna elephant *Loxodonta africana*
 93 living in the bush or savanna, and the forest elephant *Loxodonta cyclotis* in the forest (Owen-Smith
 94 1988; Roca 2001; Sukumar 2003; Sanders 2010; Sanders 2024). Despite this relatively low
 95 diversity, these megaherbivores play particularly important roles in ecosystems (e.g., Owen-Smith
 96 1988; Fritz *et al.* 2002).

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97 In the past, the study of morphological changes in African elephant fossils (prominently in their
 98 dentition) has been provided important proxies for-in biochronology and trends of-used-to
 99 approximate environmental changes (Beden 1987, 1983, 1980; Saegusa & Gilbert 2008; Sanders
 100 & Haile-Selassie 2012). This contributed to the narrative of a late Neogene spread of drier
 101 environments and related biomes (prominently savanna) having a driving impact on faunal
 102 evolution, and notably on hominids (Bobe 2006; Cerling *et al.* 2013, 2011, 1997; Strömberg 2011).
 103 Further attention was latter brought to the ecology of African elephantids, notably to their isotopic
 104 ecology (e.g., Cerling *et al.* 2015, 1999; Uno *et al.* 2011) alongside with other taxa. Lister (2013)
 105 demonstrated a decoupling between these morphological trends and paleoecological signals at a
 106 regional scale (e.g., eastern Africa), suggesting a significant delay between environmental
 107 changes, dietary behaviors, and morphological responses. However, Saarinen and Lister (2023)
 108 recently observed a stepwise increase in crown height (hypsodonty) in true elephants happened
 109 rapidly but in distinct stage after 5 Ma, specifically corresponding to peaks in arid climatic
 110 phases. A decoupling between these morphological trends and paleoecological signals was

111 demonstrated by Lister (2013) and Saarinen & Lister (2023) at regional scale (i.e., eastern Africa),
112 suggesting substantial delay between environmental changes, dietary behavior, and morphological
113 responses.

114 Our purpose is to further explore the modalities and factors of this decoupling, by investigating
115 the evolutionary history of a well-documented, single lineage at a local scale. The rich fossil record
116 of the chronostratigraphically well-controlled Shungura Formation (Lower Omo Valley,
117 southwestern Ethiopia, Turkana Depression; Arambourg 1948; Howell & Coppens 1974;
118 Boisserie *et al.* 2008) is particularly ~~adapted-suitable for~~ pursuing this objective. Our research
119 focused on *Elephas recki*, the most abundant proboscidean in the Shungura Formation, commonly
120 found in the Plio-Pleistocene paleontological sites of eastern Africa (Maglio 1972, 1973; Gilbert
121 & Asfaw 2008). The remains ~~of~~ attributed to *Elephas recki* (classically considered as a coherent
122 evolutionary lineage, but for some authors may include different lineages; see below) from
123 Shungura are considered as essential in establishing a reliable biochronological framework for the
124 region as well as for paleoenvironmental reconstructions, given that they display considerable
125 modifications of the craniodental apparatus seemingly in response to the spread of more open
126 environments and greater competition for grazing resources (Maglio 1973; Cerling *et al.* 1999;
127 Sanders & Haile-Selassie 2012; Zhang 2020; Sanders 2024).

130 MATERIAL AND METHODS

132 FOSSIL SAMPLE

133 The fossils examined in this study were collected from the Shungura Formation by the
134 International Omo Research Expedition (IORE, 1967-1976) and the Omo Group Research
135 Expedition (OGRE, since 2006). The Plio-Pleistocene deposits of the Shungura Formation are
136 well-known for their almost continuous stratigraphic sequence, i.e. ca. 800 m of sediments of
137 fluvio-lacustrine and volcanic origins divided into 12 geological members (from the lowermost to
138 the uppermost levels: Basal, A to H, and J to L) subdivided into variable numbers of units (e.g.,
139 12 for Member B, 5 for Member F) corresponding generally to sedimentological cycles (Heinzelin,
140 1983). Each member was defined by a widespread volcanic ash at its base (except for the Basal
141 Member), and many other volcanic layers are found within members. The chronology of this

142 formation is one of the best known thanks to radiochronology (K/Ar and $^{40}\text{Ar}/^{39}\text{Ar}$ methods) and
 143 by magnetostratigraphy, indicating a time span from 3.75 Ma to 1.09 Ma (Fig. 1A; see recent age
 144 model in Gardin *et al.* 2024 and references herein). Thanks to these features, the Shungura record
 145 is particularly suitable for studying-tracking the evolution, paleoecology and environmental
 146 context of various vertebrate groups, ~~as well as their paleoecology and the contemporary~~
 147 environments at community level (Bobe & Eck 2001; Alemseged 2003; Bobe 2011).

148 In the Shungura stratigraphic sequence, ~~fossil~~-elephantid fossils are abundant and constitute a
 149 significant proportion of the collected mammals (Coppens 1978; Beden 1980; Alemseged *et al.*
 150 2003, 2007). There are at least five species known from Shungura, including *Elephas ekorensis*,
 151 *Loxodonta exoptata*, *Loxodonta adaurora*, and *Loxodonta atlantica*. The fifth one, *Elephas recki*,
 152 is the most common, comprising 98 % of elephantid specimens (Coppens 1978). Based notably
 153 on the Shungura record, Beden (1980) divided *E. recki* into five subgroups recorded throughout
 154 the stratigraphic sequence of the Shungura Formation (Fig. 1B,D) and were exclusively identified
 155 through changes in cheek tooth morphology (Beden 1980). The same author later refined their
 156 stratigraphic occurrences (Beden 1987): *E. recki brumpti* (members A and B), *E. recki*
 157 *shungurensis* (members C to F), *E. recki atavus* (members F and G), *E. recki ileretensis* (from
 158 Member K to lower Member L, til Unit L-4 included, and possibly also from Member J), and *E.*
 159 *recki recki* (upper Member L, from Unit L-5 to top).

160 There is an on-going debate about these systematic ~~biology of~~ biology of attributed within the "*Elephas*
 161 *recki* complex" (Todd 2005; Zhang 2020; Sanders 2024). The disagreement is specifically about
 162 whether the earliest and youngest subspecies should be placed within the genus *Elephas* or,
 163 instead, within *Phanagoroloxodon* and *Palaeoloxodon*, respectively. Based on cranial
 164 morphopology, Saegusa & Gilbert (2008) and Zhang (2020) advocated for relating *E. recki*
 165 *ileretensis* and *E. recki recki* within *Palaeoloxodon* ~~(a subgenus of *Elephas* for the former authors,~~
 166 ~~a genus on its own for the latter)~~. However, the implications of this view result ~~this view~~ remains
 167 discussed in the literature (Sanders 2024). First, the inclusion of *Elephas recki brumpti* within
 168 *Phanagoroloxodon* is not universally approved (Sanders 2024). Second, *Palaeoloxodon* is
 169 considered as a subgenus of *Elephas* by various authors (e.g., Saegusa & Gilbert 2008) instead of
 170 a genus on its own as proposed by Zhang (2020). Third, recent paleogenetic work (Meyer *et al.*
 171 2017; Palkopoulou *et al.* 2018) suggested that recent Eurasian specimens of *Palaeoloxodon* are
 172 nested within *Loxodonta*, which conflicts with phylogenetic results based on morphology (Zhang

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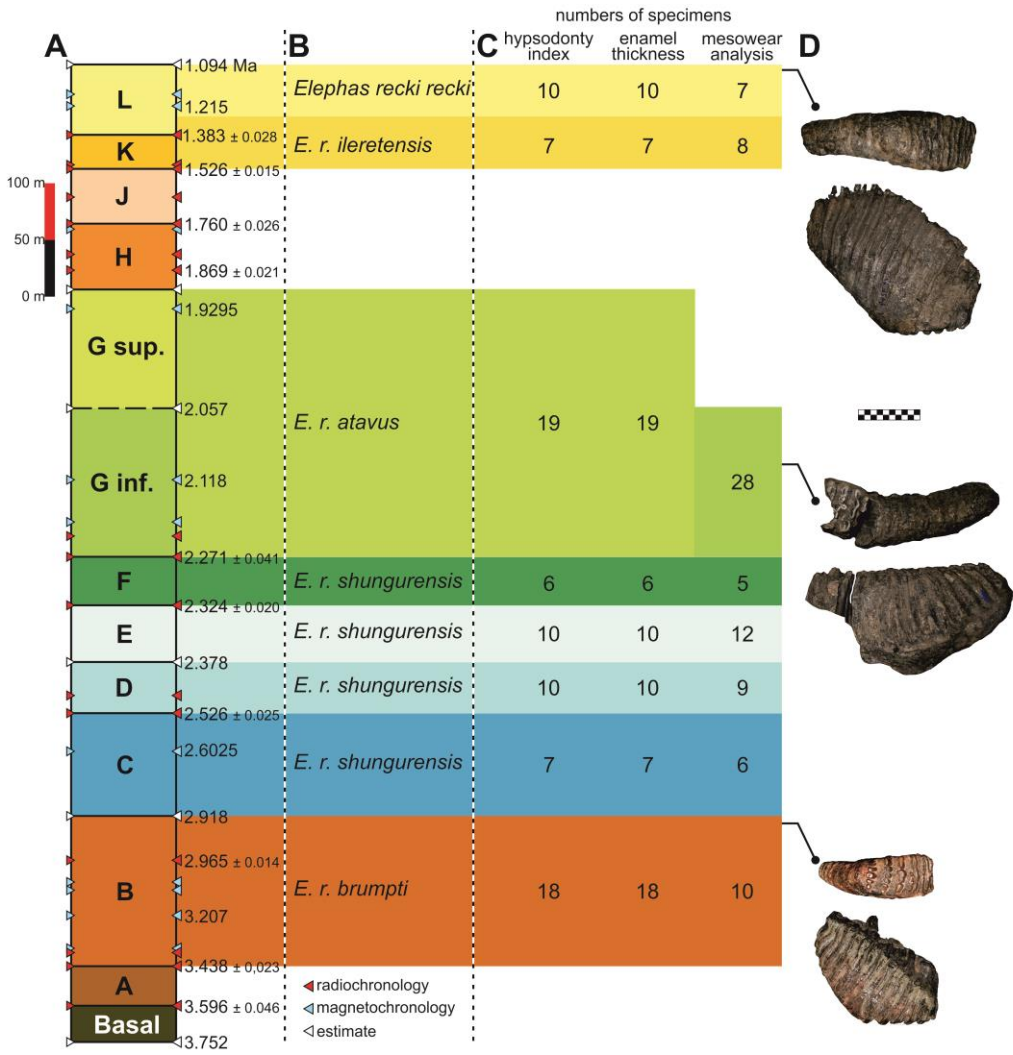
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173 [2020](#)). Consequently, ~~for~~ this paper, ~~which purpose is not intending~~ to test conflicting taxonomic
174 hypotheses, we choose to retain the classical use of subspecific division within *Elephas recki* [for](#)
175 [two reasons: the vast majority of this material can still be interpreted as belonging to a single](#)
176 [anagenetic lineage, and its subspecific subdivision retains a practical dimension](#). We note however
177 that subspecies is a term more relevant to geographic variations within a species than to
178 evolutionary stages, and that the subspecies of *E. recki* are more adequately viewed [either as](#)
179 [arbitrary, practical temporal stages displaying marked overlaps, or as substitutes for chronospecies](#)
180 [\(see Sanders 2024 for a more thorough discussion\)](#). [Although we believe that further examinations](#)
181 [of the Shungura record may eventually lead to the definition of chronospecies, this paper that does](#)
182 [not consider all relevant data to perform a proper taxonomic revision is not the right place to take](#)
183 [such decisions](#).

184 In our study, we also focused on analyzing only well-preserved molars of *Elephas recki*. Indeed,
185 mammalian teeth are highly resistant to weathering, which makes them common skeletal remains
186 found in the fossil record. They provide vital information about morphology, past dietary habits,
187 vegetation, environmental conditions, and climate (Beden 1980; Maglio 1973; Saarinen *et al.*
188 2015; Sanders 2010; Ungar 2010; [Saarinen & Lister 2024](#)). As mammalian teeth are abundant,
189 most research on mammalian dietary studies has relied mainly on the analysis of molar teeth.

190 For biometric and dental wear analyses, we utilized a total of 140 lower and upper molars,
191 including five pairs of antimeres for which average values were used (Table [S4S5](#)). We used a
192 total of 89 lower and upper third molars for biometric analysis, and 86 diagnosable cheek teeth for
193 dental wear analysis (Fig. 1C). Among these, 33 samples were analyzed in combination with both
194 methods. Upper and lower molars display statistically similar values for biometric features and
195 mesowear analysis, as stated in the supplementary information (Tables S1-S3).

196 The specimens for our study were selected based on the quality of preservation. We only chose
197 molars that had good preservation quality for both methods. Molars were chosen from all members
198 of the Shungura Formation, except for the Basal Member and Member A, as no well-preserved
199 fossil evidence was available from these members. We also excluded molars from members H
200 and J due to poor preservation quality of relatively few samples. This created a further gap of
201 approximately 355 thousand years. A similar gap is present in our mesowear sample, with an
202 additional lack of coverage for the approximately 170 thousand years represented by the upper
203 part of Member G.

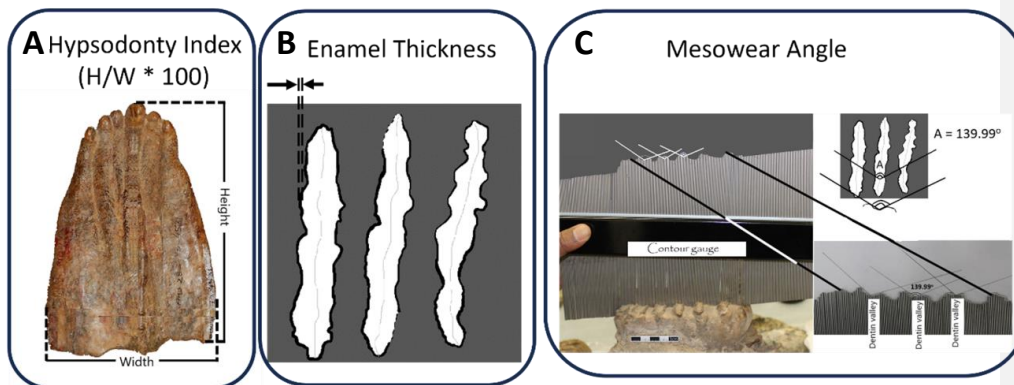


205 FIG. 1. – **A**, chronostratigraphic composite column of the Shungura Formation following the stratigraphic
 206 and radiochronological work described by Heinzelin (1983) and with ages derived from (Feibel *et al.* 1989;
 207 Brown *et al.* 2006; McDougall and Brown, 2008; McDougall *et al.* 2012; Kidane *et al.* 2014); **B**, distribution
 208 of the subspecies of the lineages of *E. recki* through the members of the Shungura Formation and
 209 corresponding numbers of analyzed specimens; **C**, sample sizes of analyzed molars from each member; **D**,
 210 morphological and biometric changes over time exemplified in lateral (bottom) and occlusal (top) views
 211 for, from top to bottom, *Elephas recki brumpti* (OMO 3/0-1974-961, M³ from B-12), *E. recki atavus*
 212 (OMO 75-1969-3196, M³ from G-4 to G-13), and *E. recki recki* (OMO K 7-1969-3204, M³ from Unit L-
 213 9). Scale bar is 10 cm for all.

214 METHODS

215 Given our interest for dental morphological evolution in response to dietary change, we selected
 216 two continuous features commonly used to depict dental evolution through time and which
 217 changes are interpreted as morphological adaptations to dietary changes (Maglio 1973; Beden
 218 1987, 1980): hypsodonty index and enamel thickness (Fig. 2A-B). We compared them to
 219 mesowear angle analysis (Fig. 2C), a recently developed proxy of dietary behavior in
 220 proboscideans (Saarinen *et al.* 2015).

221



222 Fig. 2. – Measurements acquired from teeth of *Elephas recki* from the Shungura Formation. **A**,
 223 measurements involved in the calculation of the hypsodonty index (HI); **B**, measurement of the enamel
 224 thickness (ET) on sketched occlusal surface of molar plates; **C**, measurements of mesowear angles (dentin
 225 valley angle, in white on the sketch) between ridge of enamel bands (in black) plates of a molar.
 226
 227

228 *Hypsodonty Index (HI)*

229 This index (height/width of the crown $\times 100$) is a measure of the relative crown height of elephantid
 230 molars, ~~which the~~ increase of which through time has been related to increasing abrasiveness of
 231 ingested food caused by a combined effect of plant phytoliths and exogenous mineral particles
 232 (Janis & Fortelius 1988; Damuth & Janis 2011). Using this ratio helps to eradicate the issues of
 233 individual size differences that would make it difficult to compare relative crown height in ~~any~~
 234 ~~acceptable~~ meaningful way (Maglio, 1973). According to Maglio (1973), Arambourg (1938) was
 235 the first to develop a method to quantify this biometric characteristic and then later Cooke (1947)
 236 modified this index by multiplying the height-to-width ratio by 100, ~~as described in Maglio's~~

237 ~~monograph (1973)~~. The height of a molar crown should be measured parallel to the vertical axis
238 of the plate or loph(id), from the base of the enamel covering to the apex of the tallest pillar and
239 the width of a molar should be measured across its widest plate or lophid, including cementum
240 (see Fig. 2A).

241 242 *Enamel Thickness (ET)*

243 The thickness of molar enamel is a reliable biometric characteristic for studying the evolution of
244 ~~Elephantidae~~ molars, as noted by Beden (1980) and Maglio (1973). We followed the procedure
245 suggested by Maglio (1973). This involves taking several measurements from different locations
246 on the occlusal surface of a worn enamel band of multiple plates of a molar (series of worn enamel
247 figures of plates along the molar). A minimum of three measurements is recommended on a single
248 enamel band (see Fig. 2B), after which we calculate the average of those measurements.

249 250 *Mesowear analyses*

251 To investigate the dietary preferences within our elephantid sample, we used a relatively a new
252 technique introduced by Saarinen *et al.* (2015). This technique consists in using the macroscopic
253 relief and shape of worn molar surfaces, which, according to the seminal work by Fortelius &
254 Solounias (2000), can serve as an indicator of the relative amount of abrasive plant material,
255 primarily grass-~~material~~, consumed by herbivorous mammals. The ~~Mean-mean Mesowear~~
256 mesowear Angle-angle (MWA) has been calculated for each molar sample by taking the average
257 angle from the dentine valleys of three ~~blades~~lamellae, following the method outlined by Saarinen
258 *et al.* in 2015 (see Fig. 2C) and adapted by Saarinen & Lister (2023). However, we have also made
259 a modification to the sampling method by measuring two lamellae ~~blades~~ from a single molar. A
260 t-test was conducted on some of the representative individual measurements, but no significant
261 difference was found between the series including three samples per individual and that including
262 two samples only (see Supplemental materials Table ~~S3A~~S4A-C and associated information). This
263 modification is useful for including smaller specimen fragments with well-preserved dental
264 surfaces, which can help to increase the sample size.

265 266 *Statistics*

267 We used both parametric and non-parametric statistical tests to analyze our data. To analyze the
268 variations in biometric and MWA values within *E. recki* across the Shungura sequence we
269 conducted several statistical tests including the Mann-Whitney test to assess whether the
270 differences in the median values of the biometric characters (HI and ET) across Shungura members
271 are statistically significant. Secondly, we utilized One-way ANOVAs to examine the variation
272 among the MWA values throughout the Shungura sequence. Additionally, to investigate the
273 differences between lower and upper molars, we carried out a t-test on both biometric and MWA
274 values, showing no significant differences (Tables S1-S3). Finally, we employed Ordinary Least
275 Square (OSL) linear regression and Pearson's correlation analysis tests to evaluate the relationship
276 between biometric evolution and dietary adaptation.

277

278 ABBREVIATIONS

279	ET	Enamel thickness
280	HI	Hypsodonty index
281	MWA	Mesowear Angle analysis
282	LG	lower part of Member G (units G-1 to G-13)
283	LL	lower part of Member L (units L-1 to L-4)
284	UG	upper part of Member G (units G-14 to G-29)
285	UL	upper part of Member L (units L-5 to L-9)

286

287

288 RESULTS

289

290 HYPSONDONTY INDEX (HI)

291 Descriptive statistics for this variable are provided in Table 1 and distributions are illustrated by
292 Fig. 3. The Kruskal-Wallis test showed a significant difference ($p \ll 0.001$) between HI median
293 values of Shungura stratigraphic members, suggesting the assumption that HI values increased
294 progressively through time is not accurate. Our results indicate a complex trend in the crown height
295 of this group of elephantids, which contradicts the expected gradual change over time. Instead, we
296 observe a change in the trend slope, with values for members K and L specimens much higher than
297 expected if the rate of change observed from B to G remained constant through time (Fig. 3). We

298 also conducted a Mann-Whitney multiple comparison test and found highly significant differences
 299 in HI median values notably between members K and L, and between these higher members and
 300 members lower in the sequence (Table 2). Other significant differences appeared between
 301 members B and F, B and LG, D and F, D and LG, F and K&LL.

302

303 TABLE 1. – Statistical summary of biometric characters by members/submembers.

304

Members	B	C	D	E	F	G	K & LL	UL	
N	17	7	9	9	7	19	7	10	
HI	Min	100.8	123.8	102.4	117.5	121.4	102	154.8	183
	Max	151.3	159.6	142.5	143.1	167.4	169.5	188	239.9
	Mean	122.2	134.8	118.9	129.8	145.1	135.5	173.2	202.7
	Media	121.4	129.5	119	131.1	141	137.1	176	202
	n								
SD	14.50	12.18	13.73	8.91	18.72	17.95	11.86	16.23	
ET	Min	3.8	2.6	2.8	3	2.6	2.7	2.4	2.3
	Max	4.5	3.5	3.5	3.6	3.6	4	3.6	3
	Mean	4.12	3.15	3.27	3.19	3.24	3.37	3.01	2.60
	Media	4.22	3.20	3.35	3.18	3.21	3.40	2.99	2.60
	n								
SD	0.246	0.310	0.245	0.218	0.330	0.311	0.474	0.236	

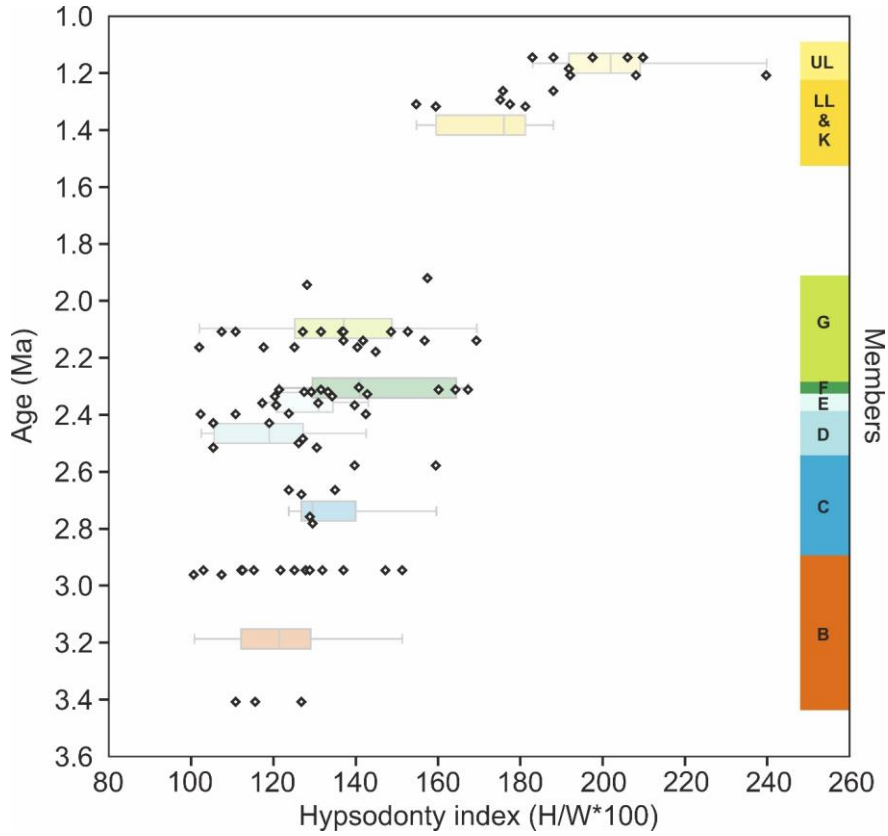
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306 TABLE 2. – Mann-Whitney pairwise tests significant for HI and ET- values.

307

	B	C	D	E	F	G	K & LL	UL
B		1.6E-04	3.7E-05	3.7E-05	1.6E-04	1.2E-06	1.6 E-04	2.0E-05
C	0.057		0.3875	0.957	0.746	0.084	0.701	0.005
D	0.571	0.057		0.417	0.872	0.267	0.392	8.5E-04
E	0.118	0.525	0.064		0.592	0.065	0.483	4.4E-04
F	0.010	0.224	0.015	0.112		0.292	0.442	0.003
G	0.025	0.729	0.027	0.302	0.312		0.102	4.4E-05
K & LL	1.8E-04	0.005	0.001	0.001	0.021	3.4E-04		0.076
UL	2.2E-05	7.6E-04	2.8E-04	2.8E-04	7.6E-04	1.5E-05	0.0011	

308 Lower left values: test results for HI; upper right values: test results for ET; gray shaded, and bold face
 309 values are highlighting significant difference (p<0.05) and highly significant (p<0.01) between members,
 310 respectively.

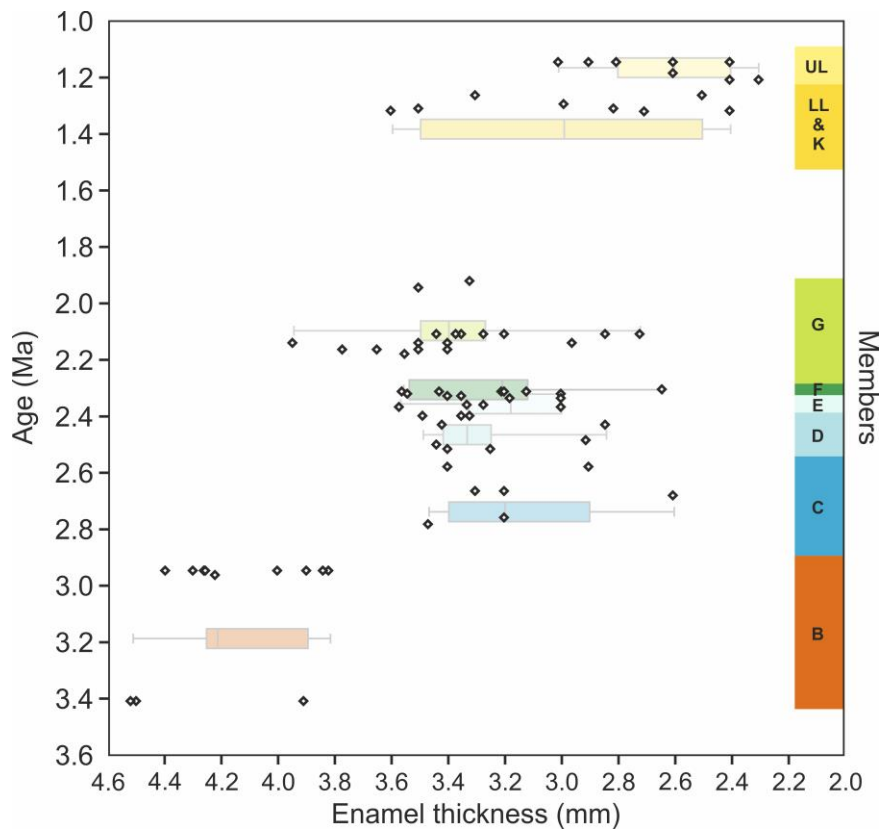


311
 312 FIG. 3. – Box plots and individual values for hypsodonty index (HI) through time in *Elephas recki* from
 313 different members of the Shungura Formation. Boxes represent the interquartile range, bar is median, and
 314 whiskers represent the lower and the upper 25 % of the data.

315
 316 ENAMEL THICKNESS (ET)

317 Descriptive statistics for this variable are provided in Table 1 and distributions are illustrated by
 318 in Fig. 4. The Kruskal-Wallis test indicates that there are statistically significant differences ($p <$
 319 0.05) among ET values. Results of the Mann-Whitney pairwise test is reported in Table 2. Member
 320 B and upper part of Member L samples display mostly highly significant differences with all other
 321 samples. This is suggestive of two episodes of sudden changes in ET values, dropping from a mean
 322 value of 4.2 mm for Member B (*Elephas recki brumpti*, N=18) to a mean value of 3.25 mm for all

323 samples from C to LL (N=61). There is no overlap between Member B sample and the rest of the
 324 samples, except a limited one with the larger LG sample (N=17; Fig. 4). For the upper part of
 325 Member L (N=10), again we observed a drop in ET to a mean value of 2.6 mm, however with a
 326 less significant difference with K and LL sample than with other samples. Despite the temporal
 327 gap between Member G and Member K samples, we did not observe significant differences
 328 between them (Table 2).
 329



330
 331 FIG. 4. – Box plots and individual values for enamel thickness (ET) through time in *Elephas recki* from
 332 different members of the Shungura Formation. Boxes represent the interquartile range, bar is median, and
 333 whiskers represent the lower and the upper 25 % of the data.
 334

335 MESOWEAR ANGLE (MWA)

336 Descriptive statistics for MWA are provided in Tables 3 and 4. Mesowear distributions by
 337 members and subspecies are illustrated by Fig. 5A-B. The results of the ANOVA on MWA ($F =$
 338 1.33 , $df = 7$, and $p = 0.250304$) indicated that is no significant difference between members of the
 339 Shungura Formation for this variable. Following Saarinen *et al.* & Lister (2023) who classified
 340 samples as browsers if their mean MWA is below 113.106° , mixed feeders if their mean MWA is
 341 between 113.106° and 124.117° , and grazers if their mean MWA is above 124.117° , we determined
 342 that *Elephas recki* from Shungura ~~tend to display~~ a predominantly grazing diet, with a ~~tendency~~
 343 ~~towards limited more occurrences of~~ mixed diets for members B, D, F, K, and subspecies *E. r.*
 344 *brumpti* and *P. r. ileretensis* (Fig. 5; Tables 3 and 4).

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346 TABLE 3. – Statistical summary of MWA measurements by members.

347

Members	B	C	D	E	F	LG	K	L
N	8	6	8	11	5	27	8	7
Min	113.5	116.3	109.2	108.7	109.7	112.0	115.0	120.7
Max	130.0	135.0	135.0	139.5	129.3	152.3	131.0	129.0
Mean	122.30	125.66	118.61	128.78	122.34	127.98	122.90	124.30
Median	124.20	126.83	117.70	132.70	128.30	128.30	123.25	124.00
SD	6.957	8.380	8.808	9.563	9.059	10.512	4.809	2.841

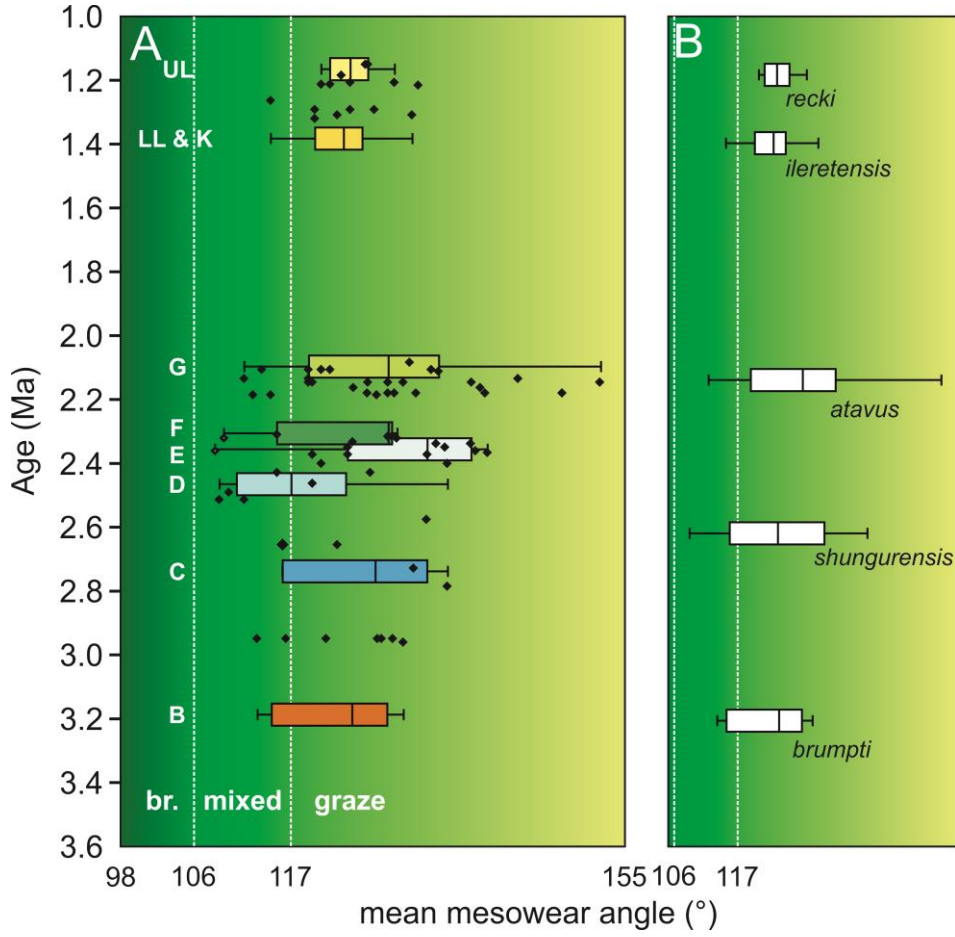
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349 TABLE 4. – Statistical summary of MWA measurements by subspecies.

350

Subspecies	<i>brumpti</i>	<i>shungurensis</i>	<i>atavus</i>	<i>ileretensis</i>	<i>recki</i>
N	8	30	27	8	7
Min	113.5	108.7	112.0	115.0	120.7
Max	130.0	139.5	152.3	131.0	129.0
Mean	122.30	124.37	127.98	122.90	124.30
Median	124.20	124.00	128.30	123.25	124.00
SD	6.957	9.176	10.548	4.809	2.841

351



352
 353
 354 FIG. 5. – Box plots and individual values for mesowear angles (MWA) through time in *Elephas recki* from
 355 the Shungura Formation. **A**, by members; **B**, by subspecies. Boxes represent the interquartile range, bar is
 356 median, whiskers represent the lower and the upper 25 % of the data. Limits for dietary categories browse
 357 (br.)-mixed and mixed-graze at 113-106 and 124-117, respectively, were obtained-revised by Saarinen &
 358 Lister (Saarinen et al. 2023[5]).
 359

360 DISCUSSION

361

362 The evolution of hypsodonty in herbivorous mammals has long been believed as one of the best
363 examples of a sustained macroevolution ~~drived~~ caused by abrasive diets (Damuth & Janis 2011;
364 Kaiser *et al.* 2013). This trend has well recognized in the evolutionary history of different
365 mammalian groups (Simpson, 1945). It is also an indication of convergent evolution in various
366 clades since, considered evidence of an adaptation for an abrasive herbivorous diet generally and
367 mainly consisting of grasses (Williams and Kay 2001; Cerling *et al.* 2011).

368 ~~As proposed by~~ According to Maglio (1973) ~~in his monograph, the~~ in African representatives
369 of the genus *Elephas*; ~~the~~ the rate of change in molar structure, mainly for the HI and ET, was
370 extremely rapid. He also ~~outlined~~ highlighted that the functional ~~meaning~~ implications of the
371 enamel thinning is less ~~obvious~~ salient, however, for Plio-Pleistocene elephantid lineages that had
372 a reduced enamel thickness while being subjected to a highly abrasive diet linked to grass feeding
373 preference.

374 Concerning the hypothesis of continuous increase in HI and reduction in ET, our results
375 generally do not support it. In both biometric characters, no apparent progressive, continuous
376 evolutionary changes were observed in the lineage of *E. recki* from the Shungura Formation.
377 Instead, a significant decrease in enamel thickness was observed between members B and C,
378 followed by little to no change from Member C to upper Member G, then again going through a
379 rapid, significant decrease in the top members (K, L). Statistically, little significant changes
380 between members were perceived in the values of hypsodonty from members B to G, until values
381 display significant increases in K-LL then in UL. These patterns fit better the description of
382 “incremental changes” suggested by Lister (2013), and of stepwise evolution by Saarinen & Lister
383 (2023), to mark a difference with gradual evolution through “infinitesimal gradations”
384 accumulated in a continuous process. Temporal gaps in our sample do not impact this
385 interpretation of successive sudden changes and stasis, but it is expected that future collections in
386 Shungura deposits within non-documented time intervals will help identifying more precisely
387 tipping ages, especially between the lower sequence of Shungura and members K and L.

388 One way to interpret these results is that they would correspond to the lumping of multiple
389 lineages within the “*Elephas recki* Complex” (see Zhang 2020). However, similar stepwise
390 changes in evolutionary rates can be observed in single lineages (Saarinen & Lister 2023) and

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391 should not necessarily seen as evidence for taxonomic comingling. As indicated above, the debate
392 on the taxonomic status of *Elephas recki* has still not reached a conclusion (e.g., Sanders 2024).
393 We considered here that the studied material is more parsimoniously interpreted as belonging to a
394 single lineage (following Beden 1980 among many others). We acknowledge that subspecies were
395 used as a practical rank for chronological stages, and that alternatively at least some of them could
396 be reconsidered as chronospecies, notably *Elephas recki brumpti* in agreement with Sanders
397 (2024). Yet we are convinced that further work is required for finalizing the taxonomic revision of
398 the Shungura material attributed to *Elephas recki*.

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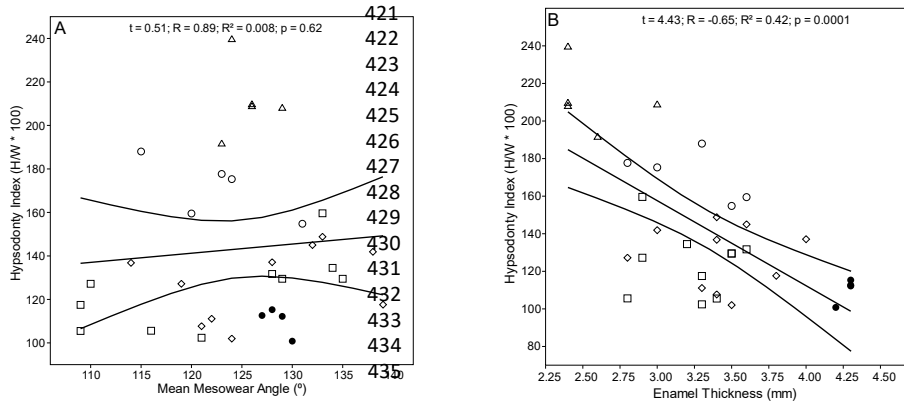
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399 In the meantime, we compared our results with those recently obtained by Saarinen & Lister
400 (2023) on the '*Elephas recki* complex' from sites contemporaneous to the Shungura Formation
401 (notably from eastern and western Turkana, see Supplementary Comparative Data Fig. S2). Their
402 results for HI very well fit are congruent with the trend and values we observed, and to some extent
403 fill the temporal gap between upper G and K, suggesting that the main shift occurred sharply just
404 after 1.5 Ma. However, compared to ours, their ET results display lower values for pre-3.0 Ma
405 specimens (equivalent to Member B in age), no change between these and values for the interval
406 3.0 Ma - 2.5 Ma (Member C age), then again lower values after 2.0 Ma. It remains to be determined
407 if these differences reflect real regional differences, if discrepancies in measurements acquired by
408 different operators could be sufficient to explain these trend and absolute value differences, or if
409 the way of defining the temporal range of specimens could impact the observed distributions.

410 As initially suggested by Maglio (1973) for elephantid molar morphology, a HI increase should
411 be correlated with a decrease in ET in relation to an increasing proportion of abrasive food in diet,
412 mainly grass. To verify these assumptions, after conducting a linear regression test, we evaluated
413 the functional relationship of variables of the two functions for the subset of specimens for which
414 HI, ET and MWA data were available. We then assessed the correlation by comparing the HI and
415 ET for specimens. The results ($R^2 = 0.42$) showed significant links between the biometric
416 characters (Fig. 6B). Regarding correlation between the two independent biometric characters (HI
417 and ET) and behavior (MWA), the results showed $R^2 = 0.008$, $t = 0.51$, $p = 0.62$ and $R^2 = 0.044$, t
418 $= 1.21$, $p = 0.23$, respectively, hence no significant correlation between the HI and MWA (Fig.
419 6A), nor between ET and MWA values (see Fig. S1).

420



436
 437 FIG. 6. – Linear regressions (Ordinary Least Squares Regression): **A**, MWA versus HI; **B**, ET versus HI in
 438 the lineage of *Elephas recki* throughout members of the Shungura Formation. Symbols correspond to
 439 members: filled circles, Member B; open squares, members C to F; open diamonds, Member G; open
 440 circles, Member K and lower Member L; open triangles, upper Member L.
 441

442 Our MWA results indicated a constantly abrasive diet (i.e., consumption incorporating a
 443 significant amount of grasses) showing little variation through time. This is consistent with results
 444 obtained by Saarinen *et al.* (2015) for *Elephas recki* from the Koobi Fora Formation (also
 445 belonging the Omo Group of the Turkana Depression). Carbon isotope contents in dental enamel
 446 of *Elephas* obtained by Cerling *et al.* (2015) between 3.5 Ma and 1.0 Ma for Kenyan formations
 447 of the Omo Group display a signal that can be interpreted as more dominated by grazing but
 448 similarly with no trend through time. Saarinen & Lister (2023) obtained also a range of variation
 449 in their MWA results constant through time (see Supplementary Comparative Data Fig. S2 C),
 450 however these are systematically lower than ours (with a mean difference of 11°). They suggest a
 451 diet dominated by low abrasion to mixed contents, and a third of individuals (N = 24 out of 68)
 452 exhibiting a grazing signal. This is at odd with our findings showing a clear majority of individuals
 453 (N = 63 out of 80) fall in the grazing range few individuals with a fully grazing signal (4.3%). This
 454 is at odd with our results (55.0 % of individual in grazing range, Fig. 5), with above mentioned
 455 isotopic data and with data indicating that the Turkana sites from Kenya sampled landscapes
 456 consistently more arid and open than the Omo (see, e.g., Levin *et al.* 2011). We suggest that these
 457 differences could be linked to different ways of acquiring MWA values, which stresses the need

458 of building comparisons based on measurements performed by a single operator at this stage of
459 methodological development.

460 At the local scale of the Shungura Formation, our results on biometric character changes of
461 *Elephas recki* are decoupled from the tooth mesowear proxy indicating a constantly abrasive diet
462 (i.e., consumption of grasses). This decoupling is congruent with results obtained by Lister (2013)
463 and Saarinen & Lister (2023) using stable isotope data and MWA, respectively, at broader
464 temporal, geographical and taxonomical scales. This suggests that behavioral adaptation, such as
465 diet or feeding preferences, preceded by far what has been depicted as morphological adaptations
466 to these preferences (Maglio 1973). Indeed, enamel isotopic contents of eastern African
467 elephantids displayed C₄ signals interpreted as a dietary behavior dominated by graze from the
468 Late Miocene until 0.2 Ma (Cerling *et al.* 2011; Uno *et al.* 2011). However, Saarinen & Lister
469 (2023) observed that, over long timescales, major proboscidean dental adaptations aligned with
470 periods of aridification, suggesting that increasing aridity and not just grazing was a key driver of
471 these evolutionary changes due to shifts toward tougher, dry-adapted vegetation.

472 Isotopic results obtained by Negash *et al.* (2020) for elephantids from the Shungura Formation
473 are not fully congruent with this view. As for our MWA results, they indicate a graze-dominated
474 mixed diet to strictly grazing diet, as previously for bovids and suids from the same formation
475 (Bibi *et al.* 2013), but also display significant differences between Member B and Member C
476 values (matching an increase in our MWA results between these two members, although non-
477 significant) and an increase in C₄ plants between the lower and the upper part of the sequence.
478 These results could be more in line with the stepwise morphological changes we observed. Yet,
479 these isotopic contents at family level cannot be attributed with certainty to *Elephas recki* and may
480 include different dental positions, and as such are not fully comparable with the dataset used in the
481 present study. In addition, the interpretation of a shift toward a more C₄ signal may not necessarily
482 reflect a significant change of dietary category (see discussion by Blondel *et al.* 2018). Future work
483 on Shungura isotopic ecology will be required to test these results specifically for *E. recki*.

484 ~~The usual~~ A common assumption in evolutionary ecology is that behavior and morphology are
485 tightly connected and that permanent changes in the former would be quickly followed by
486 morphological adaptations. On these premises, many studies suggested that global climatic
487 changes drove vegetation changes resulting in behavioral and/or morphological innovations of
488 African fauna (e.g., Coppens 1975; Vrba 1988, 1995; Bobe & Eck 2001; Negash *et al.* 2015, 2020).

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489 The decoupling between dietary behavior and morphological changes in elephantids raises the
490 possibility of alternate hypotheses for explaining morphological changes through dietary
491 adaptations.

492 The ‘evolutionary lag’ (Lister 2013) between the acquisition of the grass-dominated diets and
493 the morphological changes seen in proboscideans and other taxa could be related to intrinsic
494 evolutionary constraints, such as the slow reproduction rate of K-strategist elephantids (but not
495 explaining for relatively sudden changes in their tooth morphology) or the need to acquire first
496 craniomandibular morphologies that allow accommodating higher-crown teeth with more enamel
497 plates. [We checked the recent work by Zhang \(2020\) focusing on cranio-mandibular morphology
498 for taxonomic and phylogenetic purpose, in order to see if some of the character changes listed in
499 this comprehensive study did not suggest documented](#) particular changes that could have played a
500 role in relaxing constraints on teeth between *E. r. brumpti*, *E. r. shungurensis-atavus*, and *E. r.*
501 *ileretensis-recki*. [Our search was not successful, and this question remains open to future advances.](#)

502 External, alternative factors may also have contributed to the observed morphological changes.
503 For instance, exogenous grit could have increased as the eastern African landscapes dried up
504 during the Plio-Pleistocene time interval. This could have resulted in a higher rate of abrasion of
505 elephantid teeth, even after they transitioned to a grazing diet. Debates about the role of exogenous
506 grits in herbivore tooth wear versus abrasive plant materials are not new (e.g., Strömberg *et al.*
507 2013; Merceron *et al.* 2016). Grits significantly influenced the selective adaptation of a mammal's
508 tooth with hypsodonty, as noted by Ungar (2015). Janis and Fortelius (1988) suggested that
509 ungulates that live in open, dry habitats and feed on ground level often grow hypsodont molars.
510 However, in contrast, mammals that feed aboveground in wet, closed-canopy forests tend to have
511 brachydont (low-crowned molars).

512 Studying the amount of grit and dust in the eastern African rift system can help testing the
513 hypothesis that there is a link between external grit and the rate of morphological evolution.
514 Volcanic ash contains glassy materials that make it highly abrasive (Strömberg *et al.* 2013;
515 Madden 2015). Madden (2015) suggested that the ideal place to test this hypothesis would be the
516 fossiliferous sites in the Plio-Pleistocene geologic sequences of the eastern African rift system
517 such as in Ethiopia and Kenya, where two lineages of suids, australopithecines, and *Theropithecus*
518 all responded similarly to changes in the concentration of external grit. The lineage of *Elephas*
519 *recki* from the Shungura Formation could have responded the same way as the above referred taxa.

520 Saarinen & Lister (2023) reached this conclusion after observing tight correlations between
 521 changes in dental feature changes (including HI and ET) and peaks periods of increased aridity
 522 (based on dust in marine sediments as a proxy of general aridification and dustiness in eastern
 523 Africa). According to them, episodes of increased airborne dust may have resulted in rapid
 524 morphological responses preserved during following milder time intervals, explaining the
 525 observed stepwise evolution through this ratchet effect.

526 This requires considering that increased hypsodonty mostly responded to exogenous particles
 527 and that mesowear mostly responded to abrasion by plant silica contents. Sanson *et al.*, (2017)
 528 concluded, among others, that dental wear in general results from interaction between exogenous
 529 and endogenous (plant-embedded) abrasives. If some studies reported that exogenous particles
 530 (dust, silt, sand) has little influence on mesowear (Kaiser *et al.*, 2013), others demonstrated that it
 531 may not be necessarily the case depending on environmental conditions (e.g., Wronski & Schulz-
 532 Kornas 2015) or on the considered taxa (e.g., Martin *et al.* 2021). Experimental works and
 533 hypotheses based on medium-sized ungulates – prominently ruminants – may not so readily apply
 534 to proboscideans, i.e. hindgut fermenters of particularly large size. The apparent lack of impact of
 535 Early Pleistocene grit abundance trends on mesowear analyses in proboscideans therefore calls for
 536 caution.

537 And yet, the apparent lack of impact of Early Pleistocene grit abundance trends on mesowear
 538 analysis results calls for caution, and for iInvestigating additional ecological and morphological
 539 data from *Elephas* in well-documented, local contexts such as the Shungura Formation should
 540 provide relevant tests of single-factor explanations.

541

542

543 CONCLUSION

544

545 The decoupling we observed between morphology and diet aligned with that observed by Lister
 546 (Lister 2013) at family level, this time at the local scale of the Lower Omo Valley, with a different
 547 dietary proxy and an expanded dataset for what has been described as a single lineage. At the core
 548 of this decoupling are stepwise changes in dental morphology for which there seem to be no fully
 549 satisfying explanations for the time being. Could the abrupt change in HI and ET trends between
 550 Member G and Member K be linked to a cladogenesis resulting in a diversification of what has

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551 been named the ‘*Elephas recki* complex’ (Sanders 2024)? Or can it be explained through changes
552 in evolutionary rates within an anagenetic framework in response to environmental factors?
553 Answering to ~~this~~these questions should require additional work on Shungura elephantid
554 taxonomy, multiproxy paleoecological data focusing at specific level, and an integration of these
555 megaherbivores within the more general picture of local community dynamics. This work is
556 currently in progress and should help taking a better account of Plio-Pleistocene megaherbivores
557 within our explanations of eastern African ecosystem evolution.

558

559

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