### 1 Life histories and niche dynamics in late Quaternary proboscideans

### 2 from Midwestern North America: evidence from stable isotope

### 3 analyses

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#### 20 ABSTRACT

- 21 Stable isotopes of mammoths and mastodons have the potential to illuminate ecological changes
- 22 in late Pleistocene landscapes and megafaunal populations as these species approached
- 23 extinction. The ecological factors at play in this extinction remain unresolved, but isotopes of
- bone collagen ( $\delta^{13}$ C,  $\delta^{15}$ N) and tooth enamel ( $\delta^{13}$ C,  $\delta^{18}$ O,  ${}^{87}$ Sr/ ${}^{86}$ Sr) from the Midwest, USA are
- 25 leveraged to examine ecological and behavioral changes that occurred during the last
- 26 interglacial-glacial cycle. Both species had significant C3 contributions to their diets and
- 27 experienced increasing levels of niche overlap as they approached extinction. A subset of
- 28 mastodons after the last glacial maximum (LGM) exhibit low  $\delta^{15}$ N values that may represent
- 29 expansion into a novel ecological niche, perhaps densely occupied by other herbivores. Stable
- 30 isotopes from serial and micro-sampled enamel show increasing seasonality and decreasing
- temperatures as mammoths transitioned from Marine Isotope Stage (MIS) 5e to glacial
- 32 conditions (MIS 4, MIS 3, MIS 2). Isotopic variability in enamel suggests mobility patterns and
- 33 life histories have potentially large impacts on the interpretation of their stable isotope ecology.
- 34 This study further refines the ecology of midwestern mammoths and mastodons demonstrating
- 35 increasing seasonality and niche overlap as they responded to landscape changes in the final
- 36 millennia before extinction.

#### 37 INTRODUCTION

Historically, the late Quaternary record of mammoths and mastodons in the Midwest has 38 39 played an important role in understanding megafaunal extinctions in North America (e.g., Fisher 40 2008, 2018; Graham et al., 1981; Saunders et al. 2010; Widga et al., 2017a; Yansa and Adams, 41 2012). Whether extinctions are viewed as human-induced (Mosimann and Martin 1975; Alroy 42 2001; Surovell et al. 2005; Fisher 2009; Surovell et al. 2016), the result of late Pleistocene 43 landscape changes (Stuart et al. 2004; Nogués-Bravo et al. 2008; Widga et al. 2017a), a function 44 of other ecological processes (Ripple and Van Valkenburgh 2010), or some combination, this 45 region is unparalleled in its density of large, late Quaternary vertebrates and associated 46 paleoecological localities. For these reasons, the midcontinent has proven critical to understanding ecological dynamics in proboscidean taxa leading up to extinction. In recent years, 47 it has been recognized that multiple proboscidean taxa shared an ecological niche in late 48 49 Pleistocene landscapes (Saunders et al. 2010) despite distinct extinction trajectories (Widga et al. 50 2017a). Chronological studies indicate that proboscideans in this region experienced extinction in situ, rather than mobilizing to follow preferred niche space (Saunders et al. 2010). Despite 51 52 refinements in our understanding of mammoths and mastodons in the region, many significant 53 questions remain. Modern megafauna have a profound impact on local vegetation (Guldemond and Van Aarde 2008; Valeix et al. 2011), and it is unclear what effect late Pleistocene 54 55 mammoths and mastodons would have had on canopy cover, nutrient cycling, and fruit dispersal 56 in non-analogue vegetation communities. There are still issues of equifinality in how the 57 presence of human predators (Fisher 2009) or the virtual absence of both humans and large carnivores (Widga et al. 2017a) impacted proboscidean populations in the region as they neared 58 extinction. 59

60	The high profile debate surrounding the cause of late Pleistocene megafaunal extinctions
61	has spurred a number of productive regional studies to address the timing and paleoecology of
62	extinction in megafaunal taxa (Pacher and Stuart 2009; Stuart and Lister 2011, 2012; Stuart
63	2015; Widga et al. 2017a). The results of this research serve to constrain the number of possible
64	extinction scenarios and to highlight the need for regional-scale and taxon-specific analyses.
65	As in modern ecosystems, Proboscidea in late Pleistocene North America were long-lived
66	taxa that likely had a profound impact on the landscape around them. Due to their size and
67	energetic requirements, elephantoids are a disruptive ecological force, promoting open canopies
68	in forests through tree destruction (Chafota and Owen-Smith 2009) and trampling vegetation
69	(Plumptre 1994). Their dung is a key component of soil nutrient cycling (Owen-Smith 1992;
70	Augustine et al. 2003), and was likely even more important in N limited tundra and boreal forest
71	systems of temperate areas during the late Pleistocene. Even in death, mammoths and mastodons
72	probably wrought major changes on the systems within which they were interred (Coe 1978;
73	Keenan et al. 2018) providing valuable nutrient resources to scavengers and microbial
74	communities. Precisely because proboscideans play such varied and important roles in the
75	ecosystems they inhabit, they are good study taxa to better understand Pleistocene ecosystems.
76	They are also central players in many extinction scenarios.
77	One of the major challenges to ecological questions such as these is scale (Delcourt and
78	Delcourt 1991; Denny et al. 2004; Davis and Pineda-Munoz 2016). Processes that are acting at
79	the level of an individual or a locality can vary significantly in space and time (Table 1).
80	Paleoecological data collected from individual animals from local sites is constrained by larger
81	regional patterns that may or may not be apparent (e.g., taphonomic contexts, predator-prey
82	dynamics). Paleoecological studies of vertebrate taxa often begin with the premise that the

- 83 individual is an archive of environmental phenomena experienced in life. In long-lived taxa such
- 84 as proboscideans, this window into past landscapes may span many decades. This longevity is
- both a benefit and a challenge to studies of proboscidean paleoecology.
- 86 Table 1. Scales of paleoecological analysis.

Scale	Method	Phenomena	Selected References
Small	Stomach	Animal health, Landscape (local	Rhodes et al. 1998; Teale and Miller
(Days)	contents,	scale)	2012; Green, DeSantis, and Smith 2017;
	enamel micro-		Smith and DeSantis 2018.
	wear		
Meso	Hair,	Seasonal diets, Reproductive	Fox and Fisher 2001; Hoppe 2004;
(Weeks-	Tusk/Tooth	events (e.g., musth, calving),	Hoppe and Koch 2007; Cerling et al.
Months)	dentin, Tooth	Migration/Dispersal	2009; Metcalfe and Longstaffe 2012;
	enamel		Fisher 2018
Large	Bone	Climate/Vegetation trends,	Bocherens et al. 1996; Iacumin et al.
(Years -	collagen, tooth	Large-scale land-use trends,	2010; Szpak et al. 2010
Decades)	enamel, tooth	Population-level trends in diet	
	mesowear	responses to landscape changes.	

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Some approaches offer relatively high-resolution snapshots of animal ecology at a scale 88 89 that is of a short duration (days). Micro-wear analyses of dentin and enamel (Green et al. 2017; 90 Smith and DeSantis 2018) are increasingly sophisticated, and have the potential to track short-91 term dietary trends. The remains of stomach contents also provide ecological information at this 92 scale (Lepper et al. 1991; Newsom and Mihlbachler 2006; van Geel et al. 2011; Fisher et al. 93 2012; Teale and Miller 2012; Birks et al. 2019), which are essentially the 'last meal' representing 94 a few hours of individual browsing. These techniques offer paleoecological insights that are 95 minimally time-averaged, and at a timescale that may be comparable to modern observations of 96 animal behavior. 97 Other approaches resolve time periods that are weeks to months in duration. Fisher's 98 (Fisher and Fox 2006; Fisher 2009; 2018) work on incremental growth structures in 99 proboscidean tusk and molar dentin reliably record weekly to monthly behaviors. The resolution

of these methods may even include short-term, often periodic, life history events, such as

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101 reproductive competition (musth) and calving. Other researchers (Hoppe et al. 1999; Metcalfe 102 and Longstaffe 2012; 2014; Pérez-Crespo et al. 2016) have explored incremental growth trends 103 in proboscidean tooth enamel. Adult molars form over the course of 10-12 years with an enamel extension rate of ~1 cm/year in both modern elephants (Uno et al. 2013) and mammoths (Dirks 104 et al. 2012; Metcalfe and Longstaffe 2012) providing the opportunity to understand meso-scale 105 106 (potentially monthly) changes in diet and behavior. 107 Finally, some techniques measure animal diet and behavior over much longer scales (years-decades). In humans, bone collagen is replaced at a rate of 1.5-4% per year (Hedges et al. 108 109 2007). For equally long-lived proboscidean taxa, this means that stable isotope analyses of bone collagen is essentially sampling a moving average of ~20 years of animal growth. For younger 110 111 age groups, this average will be weighted towards time periods of accelerated maturation 112 (adolescence), when collagen is replaced at a much greater rate (Hedges et al. 2007). Tooth enamel can also be sampled at a resolution (i.e., "Bulk" enamel) that averages a year (or more) of 113 114 growth (Hoppe 2004; Baumann and Crowley 2015), and it is likely that this is the approximate temporal scale that is controlling tooth mesowear (Fortelius and Solounias 2000). 115 116 Stable isotope studies are an important part of the paleoecological toolkit for 117 understanding Quaternary proboscideans and are capable of resolving animal behavior at multiple timescales. Progressively larger, more complete datasets characterize isotopic studies of 118 119 Beringian mammoths, where stable carbon and oxygen isotopes in bone collagen and tooth 120 enamel reliably track climate and landscape changes over the late Pleistocene (Bocherens et al.

121 1996; Iacumin et al. 2010; Szpak et al. 2010; Arppe et al. 2019), the place of mammoths in

regional food webs (Fox-Dobbs et al. 2007, 2008), and characteristics of animal growth and
maturation (Metcalfe and Longstaffe 2012; Rountrey et al. 2012; El Adli et al. 2017).

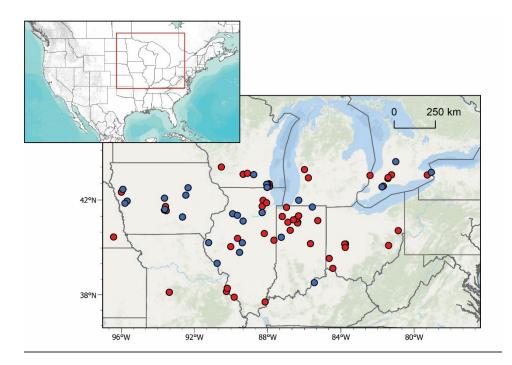
124 These studies have also been important to understanding both lineages of proboscideans 125 in temperate North America. From the West Coast (Coltrain et al. 2004; El Adli et al. 2015) to 126 the southwestern (Metcalfe et al. 2011), and eastern US (Koch et al. 1998; Hoppe and Koch 2007), isotopic approaches have been very successful in understanding local to regional scale 127 128 behavior in mammoths and mastodons. Although there have been efforts to understand 129 mammoth and mastodon behaviors in the Midwest at relatively limited geographic scales 130 (Saunders et al. 2010; Baumann and Crowley 2015), there is a need to systematically address long-term isotopic trends throughout the region. In this paper, we approach this problem from a 131 broad regional perspective, leveraging a recently reported radiocarbon (<sup>14</sup>C) dataset with 132 133 associated isotopic data on bone collagen. We also utilize an enamel dataset consisting of both 134 serial bulk and micro-milled mammoth molar enamel samples (C, O, Sr isotope systems). Together, the results of these analyses offer a picture of mammoth and mastodon diets ( $\delta^{13}$ C, 135  $\delta^{15}$ N), late Quaternary paleoclimate ( $\delta^{18}$ O), and animal mobility ( ${}^{87}$ Sr/ ${}^{86}$ Sr) that is geographically 136 137 comprehensive and spans the past 50,000 years.

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#### **MATERIALS AND METHODS**

The proboscidean dataset in this study (Figure 1; SM Table 1, SM Table 2, SM Table 3) was acquired with the goal of understanding mammoth and mastodon population dynamics during the late Pleistocene as these taxa approach extinction. Chronological and broad-scale paleoecological implications for this dataset were explored in Widga et al. (2017a) and more recently in Broughton and Weiztel (2018). In this paper we focus on the implications of these

data for the stable isotope ecology of midwestern Proboscidea. We also discuss annual patterns 144 145 in five serially sampled mammoth teeth spanning the last glacial-interglacial cycle. Finally, we micro-sampled two mammoths from the Jones Spring locality in Hickory Co., MO. Although 146 147 beyond the range of radiocarbon dating, both samples are associated with well-dated stratigraphic contexts (Haynes 1985). Specimen 305JS77 is an enamel ridge-plate recovered 148 from unit d1 (spring feeder), refitting to an M3 from unit c2 (lower peat). Unit c2 is part of the 149 150 lower Trolinger formation (Haynes 1985) and can be assigned to Marine Isotope Stage (MIS) 4. 151 Specimen 64JS73 is an enamel ridge-plate from unit e2 (sandy peat) in the upper Trolinger formation (Haynes 1985) and can be assigned to MIS 3. Together, these samples provide dozens 152 of seasonally-calibrated isotopic snapshots representing mammoth behavior from individuals that 153 154 pre-date the last glacial maximum (LGM).



156Figure 1. Map of dated midwestern mammoths (blue) and mastodons (red) with associated157 $\delta^{13}C_{coll}$  and  $\delta^{15}N_{coll}$  data. See SM Table 2 for details.

159	Mammoth and Mastodon bone collagen. Proboscidean samples were selected to widely
160	sample midwestern Proboscidea, both stratigraphically and geographically (Widga et al. 2017a).
161	Due to extensive late Pleistocene glaciation in the region, this dataset is dominated by samples
162	dating to the LGM or younger (<22 ka). Only 14 out of 93 (15%) localities predate the LGM.
163	All samples were removed from dense bone, tooth or tusk dentin and submitted to the
164	University of Arizona AMS laboratory. Collagen was prepared using standard acid-base-acid
165	techniques (Brock et al. 2010), its quality evaluated visually, and through ancillary
166	Carbon:Nitrogen (C:N) analyses. Visually, well-preserved collagen had a white, fluffy
167	appearance and C:N ratios within the range of modern bones (2.9-3.6) (Tuross et al. 1988).
168	Samples outside of this range were not included in the study. Samples that had the potential to be
169	terminal ages were subjected to additional analyses where the ABA-extracted gelatin was ultra-
170	filtered (UF) through >30 kD syringe filters to isolate relatively undegraded protein chains
171	(Higham et al. 2006). This fraction was also dated. All radiocarbon ages in this dataset are on
172	collagen from proboscidean bone or tooth dentin and available in Widga et al. (2017a), through
173	the Neotoma Paleoecology Database (www.neotomadb.org) or in SM Table 1. Measured
174	radiocarbon ages were calibrated in Oxcal v4.3 (Bronk Ramsey 2009) using the Intcal13 dataset
175	(Reimer et al. 2013). All stable isotope samples were analyzed on a Finnigan Delta PlusXL
176	continuous-flow gas-ratio mass spectrometer coupled to a Costech elemental analyzer at the
177	University of Arizona. Standardization is based on acetanilide for elemental concentration, NBS-
178	22 and USGS-24 for $\delta^{13}$ C, and IAEA-N-1 and IAEA-N-2 for $\delta^{15}$ N. Isotopic corrections were
179	done using a regression method based on two isotopic standards. The long-term analytical
180	precision (at 1 $\sigma$ ) is better than $\pm 0.1\%$ for $\delta^{13}C$ and $\pm 0.2\%$ for $\delta^{15}N$ . All $\delta^{13}C$ results are

reported relative to Vienna Pee Dee Belemnite (VPDB) and all  $\delta^{15}$ N results are reported relative to N-Air.

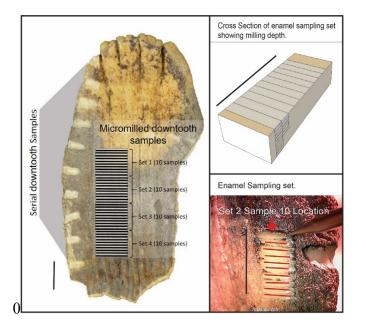
Serial and Micro-sampling of Mammoth tooth enamel. Mammoth enamel ridge-plates
were sampled at two different scales. Serial sampling consisted of milling a series of 5-10 mg
samples of enamel powder with a handheld rotary tool equipped with a 1.5 mm diameter carbide
bit along the axis of growth. Sample spacing was ~1 sample per centimeter of tooth growth.
However, given the geometry and timing of enamel maturation (Dirks et al. 2012), these samples
at best, approximate an annual scale of dietary and water inputs.

189 Micro-sampling however, has the potential to record sub-annual patterns in animal 190 movement and behavior (Metcalfe and Longstaffe 2012). For this project, we built a custom 191 micromill capable of *in situ*, micron-resolution, vertical sampling of a complete mammoth molar. 192 This micromill setup consisted of two Newmark linear stages coupled to a Newmark vertical 193 stage to allow movement in 3-dimensions. These stages were controlled by a Newmark NSC-G 194 3-axis motion controller using GalilTools on a PC. A 4-cm diameter ball joint allowed levelling of a metal (Version 1) or acrylic (Version 2) plate for holding a specimen. The armature for 195 196 Version 1 consisted of a 1971 Olympus Vanox microscope retrofitted with a stationary Proxxon 197 50/E rotary tool using a 0.5mm end mill. Version 2 has replaced this setup with a U-strut 198 armature using a 3D printed drill mount to allow for greater vertical and horizontal movement to 199 accommodate large, organically-shaped specimens. Specimens were stabilized on the mounting 200 plate using heat-flexible thermoplastic cradle affixed to a metal plate with machine screws 201 (Version 1). However, we later developed an acrylic mounting plate method where a mammoth tooth could be sufficiently stabilized using zip ties (Version 2). This micromill was developed to 202 203 address the challenges of accurately micro-milling large specimens with minimal instrumentation

204 costs. Complete plans for this micromill are available under an open hardware license at

#### 205 <u>https://osf.io/8uhqd/?view\_only=43b4242623a94a529e4c6ef2396345e9</u>.

206 Micro-mill sampling resolution was 1 sample per millimeter along the growth axis of the 207 tooth plate (Figure 2). Each sample was milled in 100µm-deep passes through the entire 208 thickness of the enamel. The lowest enamel sample (i.e., closest to the enamel-dentin boundary) in the series was used for isotopic analyses to minimize the effects of mineralization and 209 210 diagenesis on the biological signal (Zazzo et al. 2006). Enamel powder was collected in de-211 ionized water to; 1) maximize sample recovery, and 2) lubricate the mill. These samples were 212 too small for standard pretreatment of tooth enamel  $CO_3$  (Koch et al. 1997). However, paired bulk enamel samples treated with 0.1 N acetic acid and 2.5% NaOCl show results that are the 213 same as untreated bulk samples. Although this technique is both time- and labor- intensive, it is 214 215 minimally invasive and is capable of sampling enamel growth structures at high resolution.



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Figure 2. Schematic illustration of serial and micro-sampling strategies. Black bar is equal to 1 cm.

219	All enamel powder samples were measured in a Finnigan Delta Plus XL mass
220	spectrometer in continuous flow mode connected to a Gas Bench with a CombiPAL autosampler
221	at the Iowa State University Stable Isotope lab, Department of Geological and Atmospheric
222	Sciences. Reference standards (NBS-18, NBS-19) were used for isotopic corrections, and to
223	assign the data to the appropriate isotopic scale. Corrections were done using a regression
224	method using NBS-18 and NBS-19. Isotope results are reported in per mil (‰). The long-term
225	precision (at 1 $\sigma$ ) of the mass spectrometer is ±0.09‰ for $\delta^{18}$ O and ±0.06‰ for $\delta^{13}$ C,
226	respectively, and precision is not compromised with small carbonate samples (~150
227	micrograms). Both $\delta^{13}$ C and $\delta^{18}$ O are reported VPDB.
228	Enamel $\delta^{13}$ C results are corrected -14.1‰ to approximate the $\delta^{13}$ C of dietary input
229	$(\delta^{13}C_{diet})$ (Daniel Bryant and Froelich 1995).
230	Enamel $\delta^{18}$ O results were converted from VPDB to SMOW using the equation:
231	$\delta^{18}$ O SMOW = (1.03086* $\delta^{18}$ O VPDB)+30.86
232	The $\delta^{18}O$ of enamel phosphate ( $\delta^{18}O_p$ ) for these samples was calculated from the $\delta^{18}O$ of enamel
233	carbonate ( $\delta^{18}O_c$ ) (Fox and Fisher 2001)
234	$\delta^{18}O_{p} = (\delta^{18}O_{c}/1.106) - 4.7288$
235	Estimates of body water $\delta^{18}O(\delta^{18}O_w)$ were calculated following Dauxe et al. (2008)
236	$\delta^{18}O_w = (1.54^* \ \delta^{18}O_p)$ -33.72 (Dauxe et al. 2008)
237	The <sup>87</sup> Sr/ <sup>86</sup> Sr component of enamel bioapatite reflects changes in the geochemical
238	makeup of the surface an animal grazed across during tooth formation. In serial-scale analyses,

239	enamel powder samples were split from the light isotope samples described above, and represent
240	the same portions of mammoth teeth. Each ${\sim}5$ mg sample of powder was leached in 500 ${\mu}L$ 0.1
241	N acetic acid for four hours to remove diagenetic calcite and rinsed 3 times with deionized water
242	(centrifuging between each rinse). In the micromilled series, small sample sizes prevented Sr
243	analyses from being performed on the same samples as $\delta^{13}C$ and $\delta^{18}O$ analyses. Therefore, Sr
244	from these growth series was analyzed opportunistically, or at the scale of 1 sample every 2mm.
245	All enamel samples were then dissolved in 7.5 N HNO <sub>3</sub> and the Sr eluted through ion-
246	exchange columns filled with strontium-spec resin at the University of Kansas Isotope
247	Geochemistry Laboratory. <sup>87</sup> Sr/ <sup>86</sup> Sr ratios were measured on a Thermal Ionization Mass
248	Spectrometer (TIMS), an automated VG Sector 54, 8-collector system with a 20-sample turret, at
249	the University of Kansas Isotope Geochemistry laboratory. Isotope ratios were adjusted to
250	correspond to a value of 0.71250 on NBS-987 for ${}^{87}$ Sr/ ${}^{86}$ Sr. We also assumed a value of ${}^{86}$ Sr/ ${}^{88}$ Sr
251	of 0.1194 to correct for fractionation.

The distribution of <sup>87</sup>Sr/<sup>86</sup>Sr values in vegetation across the surface of the midcontinent is 252 determined by the values of soil parent material. In a large part of this region, surface materials 253 254 are composed of allochthonous Quaternary deposits such as loess, alluvium, and glacial debris. 255 Therefore continent-scale Sr isoscape models derived from bedrock or water (Bataille and Bowen 2012) are not ideal for understanding first order variability in midwestern <sup>87</sup>Sr/<sup>86</sup>Sr. For 256 257 these reasons, Widga et al. (2017) proposed a Sr isoscape for the Midwest based on surface vegetation. At a regional scale, these trends in vegetation <sup>87</sup>Sr/<sup>86</sup>Sr reflect the Quaternary history 258 of the region, and are consistent with other, empirically derived, patterns in Sr isotope 259 260 distribution from the area (Slater et al. 2014; Hedman et al. 2018). Sr isotope values in mammoth

261 enamel are compared to a <sup>87</sup>Sr/<sup>86</sup>Sr isoscape constructed from the combined datasets of Widga et
262 al. (2017b) and Hedman et al. (2009), and Hedman et al. (2018).

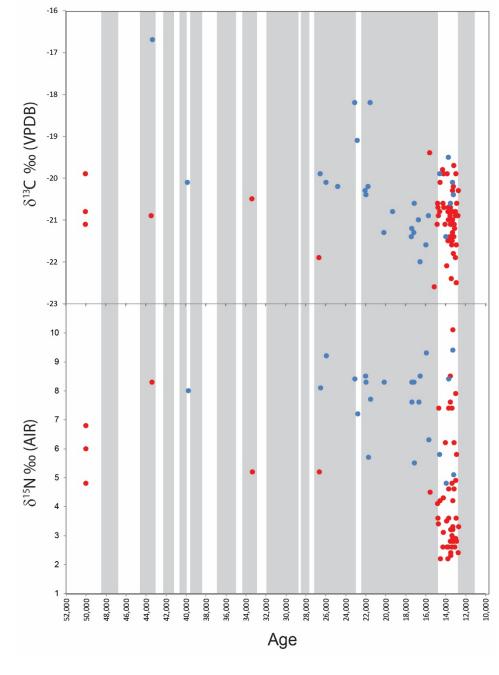
#### 263 **RESULTS**

The collagen of 54 mastodons and 22 mammoths was analyzed for  $\delta^{13}C_{coll}$  and  $\delta^{15}N_{coll}$ (Table 2). Three mastodons have <sup>14</sup>C ages that place them beyond the range of radiocarbon dating, and the youngest mastodons date to the early part of the Younger Dryas, shortly before extinction. Despite being well-represented prior to the LGM and during deglaciation (i.e., Oldest Dryas, Bølling, Allerød, Younger Dryas), this dataset lacks mastodons from the study region during the coldest parts of the LGM. Mammoths are present in this dataset from 40 ka until their extinction in the region during the late Allerød.

271 Visually, there is substantial amounts of overlap in the  $\delta^{13}C_{coll}$  and  $\delta^{15}N_{coll}$  of mammoths 272 and mastodons through the duration of the dataset (Figure 3, Table 2). Both taxa show average 273  $\delta^{13}C_{coll}$  values around -20‰, consistent with a diet dominated by C3 trees, shrubs, and/or cool-274 season grasses. This is broadly consistent with variable, but shared diets during time periods 275 when both taxa occupied the region. The average  $\delta^{13}C_{coll}$  values for mammoths (-20.4‰) is 276 similar to that of mastodons (-21.0‰). The average  $\delta^{15}N_{coll}$  of mammoths (7.5‰) is elevated 277 compared to mastodons (4.4‰).

However, dietary relationships between taxa are not static through time. Prior to the LGM, both  $\delta^{13}C_{coll}$  and  $\delta^{15}N_{coll}$  are significantly different between mammoths and mastodons ( $\delta^{13}C$  t-test; p=0.013;  $\delta^{15}N$  t-test; p=0.010). During the Oldest Dryas the  $\delta^{13}C_{coll}$  of both taxa is very similar, although  $\delta^{15}N_{coll}$  between mammoths and mastodons remains distinct (t-test; p=0.012). During the Allerød,  $\delta^{13}C_{coll}$  and  $\delta^{15}N_{coll}$  between taxa are indistinguishable.

283	The $\delta^{13}C_{coll}$ signature of mastodon diets changes little throughout the last 50 ka. Despite a
284	noticeable absence of mastodon material during the height of the LGM, the only significant shift
285	in the $\delta^{13}C_{coll}$ of mastodon diets occurs between the Bølling and the Allerød (t-test; p=0.008).
286	Mastodon $\delta^{15}N_{coll}$ values fall clearly into two groups, those that date prior to the LGM,
287	and those that post-date the LGM. Mastodon $\delta^{15}N_{coll}$ values are significantly higher in pre-LGM
288	samples than in samples dating to the Younger Dryas (t-test; p=0.010), Allerød (t-test; p=0.017),
289	and Bølling (t-test; p<0.000). Of note is a group of mastodons that show lower $\delta^{15}N_{coll}$ values
290	during the Oldest Dryas, Bølling, Allerød, and Younger Dryas, when the average $\delta^{15}N_{coll}$ values
291	decrease to values <5%. A similar shift is not evident in mammoths at this time.
292	Mammoth $\delta^{13}C_{coll}$ during MIS 3 is significantly different from mammoths dating to LGM
293	II (t-test; p=0.008) or the Oldest Dryas (t-test; p=0.005). There are no significant differences in
294	mammoth $\delta^{15}N_{coll}$ between different time periods.
295	



297Figure 3. Changes in midwestern proboscidean collagen  $\delta^{13}$ C and  $\delta^{15}$ N through time.298Mastodons=red, Mammoths=blue, glacial stadials = grey bars. Stadial chronology from

Rasmussen et al., 2014.

Taxon	Chronozone	Ν	$\overline{\mathbf{x}}\delta^{13}\mathbf{C}$ (s.d)	$\overline{x}  \delta^{15} N (s.d)$
Mammut	Younger Dryas (12.9-11.5 ka)	2	-20.6 (0.4)	2.9 (0.6)
	Allerød (14.0-12.9 ka)	35	-21.1 (0.6)	4.2 (2.1)
	Bølling (14.6-14.0 ka)	6	-20.4 (0.5)	3.8 (1.5)
	Oldest Dryas (14.6-17.4 ka)	5	-20.5 (0.7)	4.6 (1.6)
	LGM II (17.4-20.9 ka)	0		
	LGM I (20.9-22.9 ka)	0		
	Pre-LGM (>22.9 ka)	6	-20.9 (0.7)	6.1 (1.3)
	Average	54	-21.0 (0.7)	4.4 (2.0)
Mammuthus	Allerød (14.0-12.9 ka)	3	-20.6 (0.9)	6.1 (2.0)
	Bølling (14.6-14.0 ka)	1	-19.9	5.8
	Oldest Dryas (14.6-17.4 ka)	4	-21.4 (0.5)	7.9 (1.3)
	LGM II (17.4-20.9 ka)	4	-21.1 (0.4)	7.4 (1.3)
	LGM I (20.9-22.9 ka)	5	-20.1 (1.1)	7.6 (1.2)
	Pre-LGM (>22.9 ka)	5	-19.6 (0.8)	8.2 (0.7)
	Average	22	-20.4 (1.0)	7.5 (1.3)

Table 2. Mastodon and Mammoth collagen stable isotope values, by chronozone.  $\delta^{13}$ C values reported as % relative to VPDB.  $\delta^{15}$ N values reported as % relative to Air.

304  $\overline{\mathbf{x}} =$  mean; s.d. = standard deviation

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306 All serial enamel series were between 9 and 16 cm in length and represent multiple years. 307 Stable oxygen isotopes of three MIS 2 mammoths (Principia College, Brookings, Schaeffer) overlap significantly, while MIS 4 (234JS75) and MIS 5e (232JS77) mammoths from Jones 308 309 Spring show elevated values indicative of warmer conditions (Table 3; Figure 4). The variance in serially sampled MIS 2 mammoths (average amp. = 2.5%) was not significantly different from 310 the series from Jones Spring, dating to MIS 4 (amp. = 2.9‰)(F-test; p=0.320). However, the MIS 311 312 5e mammoth from Jones Spring has a smaller amplitude (amp. = 1.1%) compared to both the 313 MIS 4 mammoth (F-test; p=0.003) and the three MIS 2 mammoths (F-test; p<0.000). Strontium isotope ratios of this animal indicates a home range on older surfaces of the Ozarks, throughout 314 tooth formation, with a period of relatively elevated <sup>87</sup>Sr/<sup>86</sup>Sr values also corresponding to low 315  $\delta^{18}$ O values. 316

ID $\delta^{13}C$			$\delta^{18}O$			<sup>87</sup> Sr/ <sup>86</sup> Sr						
Serial Analyses	Х	Min	Max	Amp	Х	Min	Max	Amp	Х	Min	Max	Amp
Principia College (MIS 2)	-10.9	-11.3	-10.0	1.4	-7.9	-8.7	-6.7	2.0				
Brookings Mammoth (MIS 2)	-9.9	-10.5	-9.4	1.1	-9.2	-10.0	-7.8	2.2				
Schaefer Mammoth (MIS 2)	-9.0	-9.9	-7.0	3.0	-8.3	-9.9	-6.6	3.3				
Jones Sp. Mammoth, 243JS75 (MIS 4)	-1.7	-2.3	-0.2	2.1	-3.4	-4.5	-1.6	2.9				
Jones Sp. Mammoth, 232JS77 (MIS 5e)	-2.8	-3.8	-1.3	2.5	-0.5	-4.6	3.6	1.1	0.7162	0.7146	0.7177	-0.0031
Micro-Analyses					1			1				
305JS77 (MIS 5e)	-2.4	-3.5	-0.2	3.3	-3.9	-4.9	-2.9	2.0	0.7109	0.7106	0.7118	-0.0012
64JS73 (MIS 3)	-2.8	-4.3	-1.7	2.6	-4.9	-12.0	-0.4	11.7	0.7152	0.7140	0.7164	-0.0025

Table 3. Summary of serial and micro-sampled *Mammuthus* tooth enamel:  $\delta^{13}$ C,  $\delta^{18}$ O, and <sup>87</sup>Sr/<sup>86</sup>Sr.

319

320 Two mammoth teeth from the Jones Spring locality in Hickory Co., MO were micro-

321 sampled (Figure 5). These specimens are from beds representative of MIS 5e and MIS 3

322 deposits.

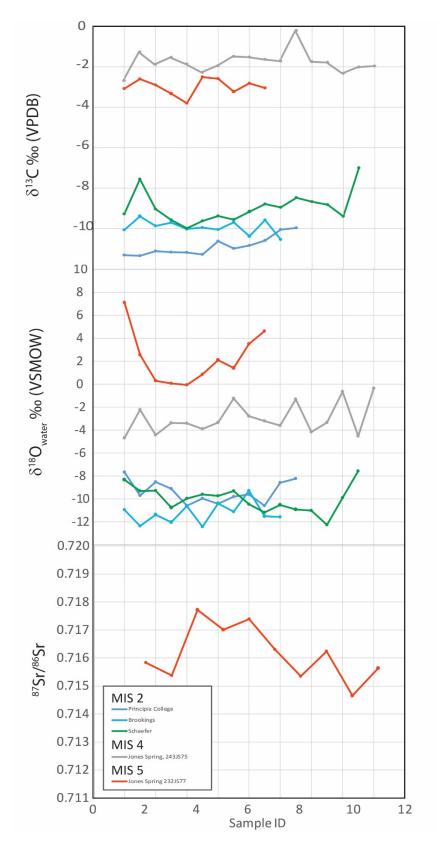
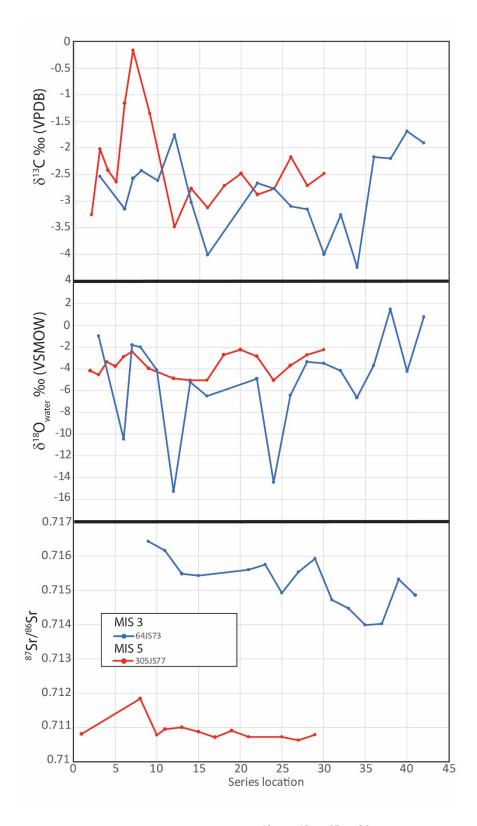


Figure 4. Time series, serial enamel  $\delta^{13}$ C,  $\delta^{18}$ O,  $^{87}$ Sr/ $^{86}$ Sr.







326

Figure 5. Time series, micro-sampled enamel δ<sup>13</sup>C, δ<sup>18</sup>O, <sup>87</sup>Sr/<sup>86</sup>Sr. Both specimens are from Jones Spring, Hickory County, Missouri.

329	The MIS 3 molar (64JS73) shows regular negative excursions in $\delta^{18}$ O values suggestive
330	of seasonal temperature changes in ingested water. These excursions are relatively short-lived
331	and extreme. Seasonal variation in enamel growth rate and maturation may account for the
332	perceived short length of these periods. Maximum $\delta^{18}O$ values in both mammoths are similar
333	however; the MIS 5e mammoth lacks these negative excursions and exhibits values that are more
334	complacent through the length of the tooth.
335	The $\delta^{13}$ C series from both molars indicate these mammoths experienced a C4 diet
336	throughout the year. However, deep dips in the $\delta^{18}$ O series of the MIS 3 molar also correspond to
337	at least two temporary peaks in the $\delta^{13}$ C series.
338	The <sup>87</sup> Sr/ <sup>86</sup> Sr isotope series from both animals demonstrate an adherence to separate
339	home ranges. The MIS 5e mammoth shows values similar to bedrock units outcropping locally
340	in central-western Missouri and neighboring areas of Kansas and Oklahoma. The MIS 3
341	mammoth, however, had a home range across rock units with much higher <sup>87</sup> Sr/ <sup>86</sup> Sr values. The
342	home ranges for these animals do not overlap, despite their recovery from different strata within
343	the same locality.

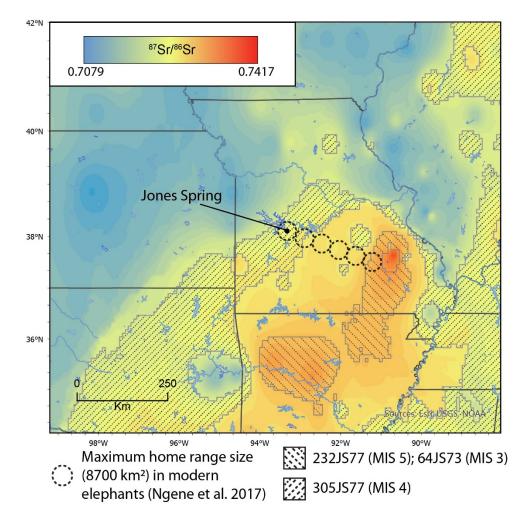


Figure 6. Mobility in mammoths from Jones Spring, Hickory Co., MO. Hatched areas indicate
the range of <sup>87</sup>Sr/<sup>86</sup>Sr values from mammoth molar ridge-plates. One mammoth (305JS77)
exhibits local values. Two separate mammoths (232JS77, 64JS73) exhibit Sr values suggesting
>200 km movement from the central core of the Ozark uplift. This implies movement over
multiple years that is at least six times larger than the maximum home range size documented in
modern elephants (Netosha National Park, Namibia). Basemap isoscape data from Widga et al.
(2017b), Hedman et al. (2009, 2018).

#### 352 **DISCUSSION**

Despite overall similar isotopic values in mammoths and mastodons throughout the period of this study, underlying nuances are informative to regional changes in niche structure and climate. The absence of mastodons during the coldest parts of the LGM suggests that *Mammut* were, at least to some degree, sensitive to colder climates and adjusted its range accordingly.

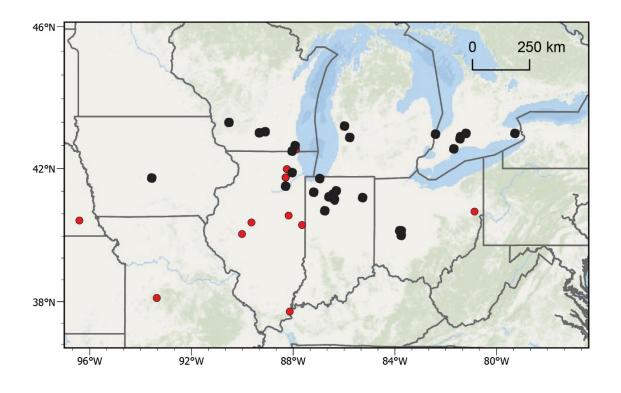
358 Comparisons between co-eval mammoths and mastodons indicate a gradual collapse of 359 niche structure. Prior to MIS 2,  $\delta^{13}C_{coll}$  and  $\delta^{15}N_{coll}$  values between taxa were significantly 360 different. By the end of the Allerød, mammoth and mastodon diets were isotopically 361 indistinguishable.

Through time, there was no significant change in the  $\delta^{13}C_{coll}$  of mastodons. Although average mammoth  $\delta^{13}C_{coll}$  in the region during the latter part of the LGM was slightly more negative than other time periods, this is not a marked shift and may be a function of decreased pCO2 during that time (Schubert and Jahren 2015). Globally, Siberian and European mammoths exhibit a similar range of  $\delta^{13}C_{coll}$  values (Iacumin et al. 2010; Szpak et al. 2010; Arppe et al. 2019; Schwartz-Narbonne et al. 2019).

In both taxa, mean  $\delta^{15}N_{coll}$  decreases slightly throughout the sequence, however, the minimum  $\delta^{15}N$  values for mastodons during the Bølling, Allerød, and Younger Dryas are significantly lower than earlier mastodons. The anomalously low values of these late mastodons are also shared with other published mastodon values in the Great Lakes region (Metcalfe et al. 2013). They are also significantly lower than contemporary midwestern, Eurasian or Beringean mammoths (with some exceptions, Drucker et al. 2018). The timing of these low, mastodon  $\delta^{15}N_{coll}$  values correspond to regionally low  $\delta^{15}N_{coll}$  values in the bone collagen of non-

375 proboscidean taxa from European late Quaternary contexts (Drucker et al., 2009; Richards and 376 Hedges, 2003; Rabanus-Wallace et al., 2017; Stevens et al., 2008) suggesting broad changes in global climate may have had cascading impacts in the N budget of local ecosystems. 377 378 However, understanding in more detail how climate might have affected N cycling in midwestern ecosystems remains unclear. Flux in soil and plant N may be a function of plant-379 based N<sub>2</sub> fixation (Shearer and Kohl 1993), rooting depth (Schulze et al. 1994), N loss related to 380 climate factors (Austin and Vitousek 1998; Handley and Raven 1992), microbial activity, or 381 382 mycorrhizal colonization (Hobbie et al. 2000; Michelsen et al. 1998;). Furthermore, the role of 383 large herbivore populations in N flux may be significant (Frank et al. 2004). In N limited 384 environments such as tundra, terrestrial plants may receive relatively more N from inorganic sources. Boreal forests like those of the Midwest during the Bølling and Allerød, however, 385 exhibit relatively greater biological productivity, so plant shoots are more likely to take in 386 volatized ammonia (low  $\delta^{15}$ N) from organic sources such as urea (Fujiyoshi et al. 2017). 387 388 At this time, it is difficult to distinguish which (if any) of these factors had an impact on late Quaternary mastodon  $\delta^{15}$ N<sub>coll</sub> values. Although there is a wide range in  $\delta^{15}$ N<sub>coll</sub> of mammoths 389 and mastodons throughout the sequence,  $\delta^{15}N_{coll}$  values below 5% are only present in mastodons 390 391 during post-LGM deglaciation in the lower Great Lakes (39-43 deg. latitude), throughout an area 392 suggested to be vegetated by a disharmonius flora dominated by Black Ash and Spruce (Gonzales and Grimm 2009; Gill et al. 2009) (Figure 7). The low  $\delta^{15}N_{coll}$  values suggest that 393 394 these mastodons occupied an undefined, local-scale, dietary niche that was not shared by 395 contemporary mammoths, or by earlier mastodons. Previous research has suggested that the Bølling-Allerød may have been a time of high mastodon populations in the Great Lakes region 396 397 (Widga et al. 2017a). The concentration of mastodons around water sources may have had an

- impact on the  $\delta^{15}$ N of browsed plants due to increased contributions of herbivore urea.
- 399 Isotopically light plant shoots can occur with increased utilization of volatized NH<sub>4</sub> from organic
- 400 sources, including urea from herbivores.



401

402Figure 7. Mastodon distribution during the Bølling-Allerød. Black circles indicate mastodon403 $\delta^{15}N_{coll} < 5\%$ . Red circles represent mastodons with  $\delta^{15}N_{coll} > 5\%$ .

404

#### 405 <u>Life Histories of midwestern Mammoths</u>

Through analyses of incremental growth structures in mammoths, such as molar enamel or tusk dentin, we can reconstruct longitudinal life histories that reflect the landscape experienced by an animal over multiple years. Late Pleistocene mammoths from the study area exhibit relatively consistent, down-tooth patterns in  $\delta^{13}$ C and  $\delta^{18}$ O. Similar bulk enamel isotope values in South Dakota, Wisconsin, and Illinois mammoths suggest access to broadly similar resources, and relatively stable access to these resources across multiple years of the life of an

individual (Figure 4). Less negative  $\delta^{13}$ C and  $\delta^{18}$ O values in the last forming samples of the 412 413 Schaeffer mammoth suggest a change in that animal's life history in the year before death. This 414 change could be explained by local environmental changes resulting in nutritional stress. 415 Two mammoths from the Jones Spring site in southwest Missouri provide a pre-LGM perspective on landscapes that mammoths occupied (Figure 4). A tooth plate dating to MIS 5e 416 from Jones Spring has less negative  $\delta^{18}$ O and  $\delta^{13}$ C values compared to the MIS 2 samples. This 417 indicates warmer overall conditions and a diet that incorporated more C4 grasses throughout 418 419 multiple years. Pollen from the same stratigraphic units further suggest that southwestern Missouri during MIS 5e was dominated by *Pinus* (no *Picea*) with a significant non-arboreal 420 421 component (King 1973). Importantly, Sr isotopes from this tooth indicate that while this tooth 422 was forming, the animal was foraging across surfaces that are more radiogenic than local values. The nearest area with  ${}^{87}$ Sr/ ${}^{86}$ Sr values >0.7140 is the central Ozark uplift to the east of Jones 423 Spring (Figure 6). The wide amplitude of  $\delta^{18}$ O values throughout the length of this tooth, 424 425 combined with the Sr isotope data suggesting adherence to the central Ozark uplift suggest 426 relatively broad shifts in annual water availability and/or that this animal utilized a variety of 427 water sources, including surface sources and freshwater springs. The MIS 4 molar from Jones Spring has  $\delta^{18}$ O values intermediate between the MIS 2 428 samples and the MIS 5e sample, along with  $\delta^{13}$ C values that indicate more C4 consumption than 429 430 MIS 2 samples (Figure 4). This mammoth occupied a cooler environment than the MIS 5e 431 mammoth, but significantly warmer than the late Pleistocene mammoths in the MIS 2 group. In the case of the two, microsampled mammoths from Jones Spring, MO, isotopic trends 432 in each animal illustrates different life histories (Figure 5). The MIS 4 mammoth occupied the 433 western Ozarks as indicated by <sup>87</sup>Sr/<sup>86</sup>Sr values deposited throughout the development of the 434

435 sampled portion of the tooth. It occupied a landscape where C4 vegetation was common, with 436 mild winter temperatures. The MIS 3 mammoth from Jones Spring occupied the central core of the Ozark uplift indicated by <sup>87</sup>Sr/<sup>86</sup>Sr values deposited during the formation of the sampled 437 438 tooth. However, between the cessation of enamel formation and death, this animal moved  $\sim 200$ km to the western Ozarks where it was preserved in the Jones Spring deposits. The  $\delta^{18}$ O series of 439 this animal suggests greater seasonality during MIS 3 compared to MIS 5e, with deep negative 440 excursions during the cold season. It also had a diet composed primarily of C4 vegetation. 441 These results suggest a broadly similar geographic scale of landscape-use in late 442 443 Quaternary mammoths in the Midwest to mammoths in the Great Plains and Florida (Esker et al. 2019; Hoppe et al. 1999; Hoppe 2004). Overall, mammoths from both regions do not engage in 444

significant seasonal migrations, but can move greater distances at annual to decadal time-scales.

Further, although  $\delta^{13}C_{coll}$  values in mammoths clearly suggest a niche that, at times, included C3rich diets in a region that was dominated by forest, not grasslands, both Jones Spring mammoths were mixed feeders with C4 grasses making up a significant part of their diet. The individual life histories of these mammoths are highly variable, underscoring the need to control for individual mobility in stable isotope studies of large herbivore taxa.

451

#### Testing scenarios of late Pleistocene Proboscidean population dynamics

452 Can this stable isotope dataset illuminate trends in late Proboscidean population 453 dynamics during deglaciation in the Midwest? It is possible that the shift in  $\delta^{15}$ N represents the 454 colonization of a novel ecological niche that also coincides with climate and vegetational 455 changes at the beginning of the Bølling-Allerød. However, it is unclear what this niche might be. 456 Stable isotopes alone do not adequately define this niche space, and further paleobotanical work 457 is necessary. Drucker et al. (2018) also noted anomalously low  $\delta^{15}$ N values in LGM mammoths

from Mezyhrich in central Europe, and attributed this pattern to a large, but unspecific, shift in dietary niche. This is counter-intuitive in warming landscapes of the Midwest since boreal forests typically have increased amounts of N fixing microbes relative to tundra environments, thus increasing the  $\delta^{15}$ N in forage, overall.

Some studies have suggested that the growth rate of an individual is inversely correlated 462 with  $\delta^{15}N$  (Warinner and Tuross 2010). This would be consistent with some scenarios of late 463 Pleistocene mastodon population dynamics. Fisher (Fisher 2009; 2018) suggests that predator 464 465 pressure from Paleoindian groups who were megafaunal specialists would have caused 466 mastodons to mature at a younger age. A decrease in the age at weaning would mean a shorter period of nursing-related, elevated dietary  $\delta^{15}$ N in young mastodons. If this were the case, we 467 would expect an overall decrease in  $\delta^{15}$ N of bone collagen in animals in their first and second 468 decade of life. In our dataset, there is no significant change in maximum or mean  $\delta^{15}$ N in 469 mastodon bone collagen, despite a subset of mastodons that have lower  $\delta^{15}$ N values. If predator 470 471 pressure is contributing to faster maturation and shortening the time of nursing, then it is only 472 occurring in some areas. However, even if this were the case in these areas, it is still uncertain what ecological processes might drive an increase in growth rate. Depending on local forage 473 conditions experienced by an animal, an increase in growth rate may be caused by increased 474 475 predator pressure (Fisher 2009) or a decrease in population density (Wolverton et al. 2009). It is also possible that low  $\delta^{15}$ N values in post-LGM mastodons represents a systematic 476 change in predator avoidance strategies among some mastodon populations (Fiedel et al. 2019). 477 Mastodons may have been attracted to low  $\delta^{15}N$  areas such as marshes and wetlands as a 478 response to new predators on the landscape (i.e., humans). However, taphonomic modification of 479 late Pleistocene proboscidean materials by humans or other predators is extremely rare in this 480

dataset (Widga et al. 2017a), so we see this scenario as an unlikely driver of mastodon landscape
use.

If proboscidean populations are dense on the landscape (Widga et al. 2017a) there may 483 484 also be rapid, significant, and long-term changes to N cycle. An influx of N via urea as would be expected in high-density areas, would cause isotopically heavy plant roots and isotopically light 485 shoots. Under intense grazing pressure, changes to N contributions change quickly from 486 inorganic soil mineral N reservoirs to N from urea (McNaughton et al. 1988; Knapp et al. 1999). 487 Individuals with lower  $\delta^{15}N_{coll}$  values during the Allerød lived in areas with high mastodon 488 489 populations, which may have contributed to isotopically light forage (due to ammonia 490 volitization, Knapp et al. 1999). However, this scenario does not explain why this shift is absent 491 in mammoths during this time.

492

#### 493 **CONCLUSIONS**

494 Stable isotopes of proboscidean tissues in the midwestern US illustrate the variability 495 inherent in modern paleoecological analyses. Traditional approaches to understanding animal 496 diets and landscape use through morphology and a reliance on modern analogues is inadequate 497 for understanding paleoecological changes within megafaunal populations, or during times of 498 rapid ecological change such as the late Quaternary.

499 Stable isotopes from bone collagen of midwestern proboscideans suggest consistently
500 C3-dominated diets over the last 50 ka. During the LGM, mammoth diets may have included C3
501 grasses, but the prevalence of C3 flora during the post-LGM period is likely due to a landscape

shift to more forest (Gonzales and Grimm 2009; Saunders et al. 2010) with grasses making up
very little of the floral communities in the southern Great Lakes region.

504 Despite the strong C3-signal in mammoth and mastodon diets during this period, the 505 range of dietary flexibility and the degree of overlap between these two taxa is striking. The 506 isotopically defined dietary niche of mammoths and mastodons show increasing overlap as they 507 approach extinction. Niche overlap is also supported by assemblages where both taxa co-occur 508 (Saunders et al. 2010; Widga et al. 2017a). However, there are exceptions to this pattern and some late mastodons exhibit low  $\delta^{15}$ N values that may indicate the evolution and occupation of a 509 510 new dietary niche, physiological responses to late Pleistocene ecological changes, or some other 511 process acting on late mastodon populations.

Further resolution of mammoth and mastodon life histories are gleaned from stable 512 isotopes in tooth enamel.  $\delta^{13}$ C and  $\delta^{18}$ O generally track climate and landscape changes 513 experienced during tooth formation. MIS 2 mammoths are broadly similar in their isotopic life 514 515 histories and illustrate relative homogeneity of landscape conditions across the Midwest at this 516 time. Micro-sampled mammoth molars from Jones Spring, MO, indicate a significant increase in 517 MIS 3 seasonality, compared to the last interglacial period (MIS 5e). Finally, some mammoths in 518 this study died >200 km away from where they lived during the formation of sampled molars 519 suggesting lifetime mobility patterns in mammoths could have a significant effect on presumed 520 'local' stable isotope values.

521

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531	

### 533 Figures

534	Figure 1. Map of dated midwestern mammoths (blue) and mastodons (red) with
535	associated $\delta^{13}C_{coll}$ and $\delta^{15}N_{coll}$ data. See SM Table 2 for details.
536	Figure 2. Schematic illustration of serial and micro-sampling strategies.
537	Figure 3. Changes in midwestern proboscidean collagen $\delta^{13}C$ and $\delta^{15}N$ through time.
538	Figure 4. Time series, serial enamel $\delta^{13}$ C, $\delta^{18}$ O, $^{87}$ Sr/ $^{86}$ Sr.
539	Figure 5. Time series, micro-sampled enamel $\delta^{13}$ C, $\delta^{18}$ O, $^{87}$ Sr/ $^{86}$ Sr. Both specimens are
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546	home range size documented in modern elephants (Netosha National Park, Namibia).
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549	mastodon $\delta^{15}N_{coll} < 5\%$ . Red circles represent mastodons with $\delta^{15}N_{coll} > 5\%$ .
550	
551	Tables
552	Table 1. Temporal Scale in Paleoecological Analyses.
553	Table 2. Mammoth and mastodon collagen stable isotope values, by chronzone.
554	Table 3. Summary of serial and micro-sampled tooth enamel series: $\delta^{13}$ C, $\delta^{18}$ O, ${}^{87}$ Sr/ ${}^{86}$ Sr.

#### **Supplemental information** 556 SM Supplemental Material. Word File 557 SM Table 1. Mammoth and Mastodon $\delta^{13}$ C, $\delta^{15}$ N data. 558 559 SM Table 2. Mammoth enamel isotope data. SM Table 3. Radiocarbon dated mammoths and mastodons from the Midwest. 560 561 References 562 563 564 Alley, Richard B. 2000. "The Younger Dryas Cold Interval as Viewed from Central Greenland." Quaternary Science Reviews 19 (1): 213-26. https://doi.org/10.1016/S0277-565 566 3791(99)00062-1. Alroy, John. 2001. "A Multispecies Overkill Simulation of the End-Pleistocene Megafaunal 567 Mass Extinction." Science 292 (5523): 1893-96. 568 569 https://doi.org/10.1126/science.1059342. 570 Arppe, Laura, Juha A. Karhu, Sergey Vartanyan, Dorothée G. Drucker, Heli Etu-Sihvola, and Hervé Bocherens. 2019. "Thriving or Surviving? The Isotopic Record of the Wrangel 571 572 Island Woolly Mammoth Population." Quaternary Science Reviews 222 (October): 105884. https://doi.org/10.1016/j.quascirev.2019.105884. 573 574 Augustine, David J., Samuel J. McNaughton, and Douglas A. Frank. 2003. "Feedbacks Between Soil Nutrients and Large Herbivores in a Managed Savanna Ecosystem." Ecological 575 Applications 13 (5): 1325–37. https://doi.org/10.1890/02-5283. 576 577 Austin, Amy T., and P. M. Vitousek. 1998. "Nutrient Dynamics on a Precipitation Gradient in Hawai'i." Oecologia 113 (4): 519–29. https://doi.org/10.1007/s004420050405. 578 579 Bataille, Clément P., and Gabriel J. Bowen. 2012. "Mapping 87Sr/86Sr Variations in Bedrock and Water for Large Scale Provenance Studies." Chemical Geology 304-305 (April): 39-580 581 52. https://doi.org/10.1016/j.chemgeo.2012.01.028. Baumann, Eric J., and Brooke E. Crowley. 2015. "Stable Isotopes Reveal Ecological Differences 582 583 amongst Now-Extinct Proboscideans from the Cincinnati Region, USA." Boreas 44 (1): 240-54. https://doi.org/10.1111/bor.12091. 584 Birks, Hilary H., Bas van Geel, Daniel C. Fisher, Eric C. Grimm, Wim J. Kuijper, Jan van Arkel, 585 586 and Guido B. A. van Reenen. 2019. "Evidence for the Diet and Habitat of Two Late Pleistocene Mastodons from the Midwest, USA." Quaternary Research 91 (2): 792-812. 587 https://doi.org/10.1017/qua.2018.100. 588 589 Bocherens, Hervé, Gilles Pacaud, Petr A. Lazarev, and André Mariotti. 1996. "Stable Isotope Abundances (13C, 15N) in Collagen and Soft Tissues from Pleistocene Mammals from 590 Yakutia: Implications for the Palaeobiology of the Mammoth Steppe." Palaeogeography, 591 592 Palaeoclimatology, Palaeoecology, Biogenic Phosphates as Palaeoenvironmental 593 Indicators, 126 (1): 31–44. https://doi.org/10.1016/S0031-0182(96)00068-5.

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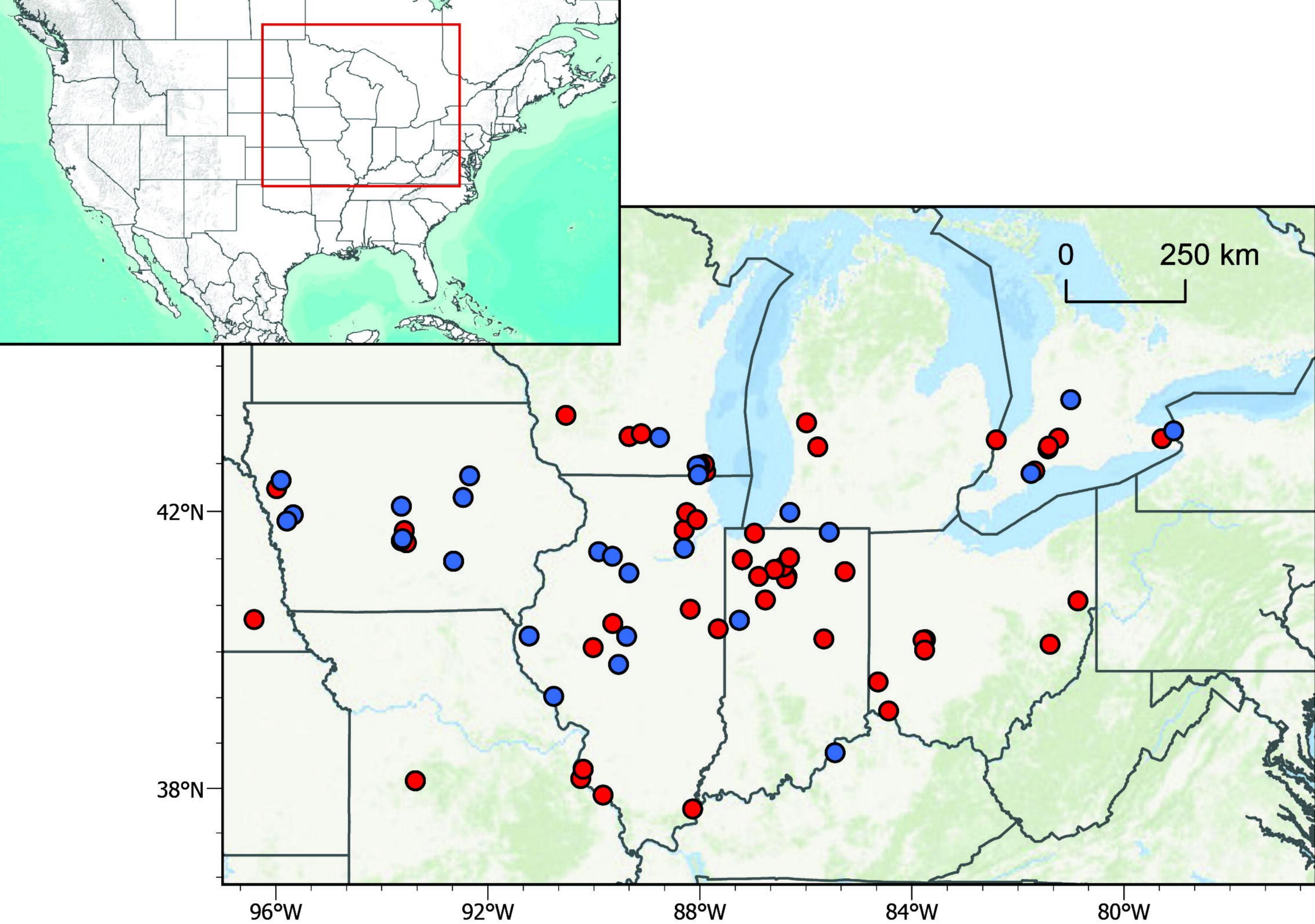
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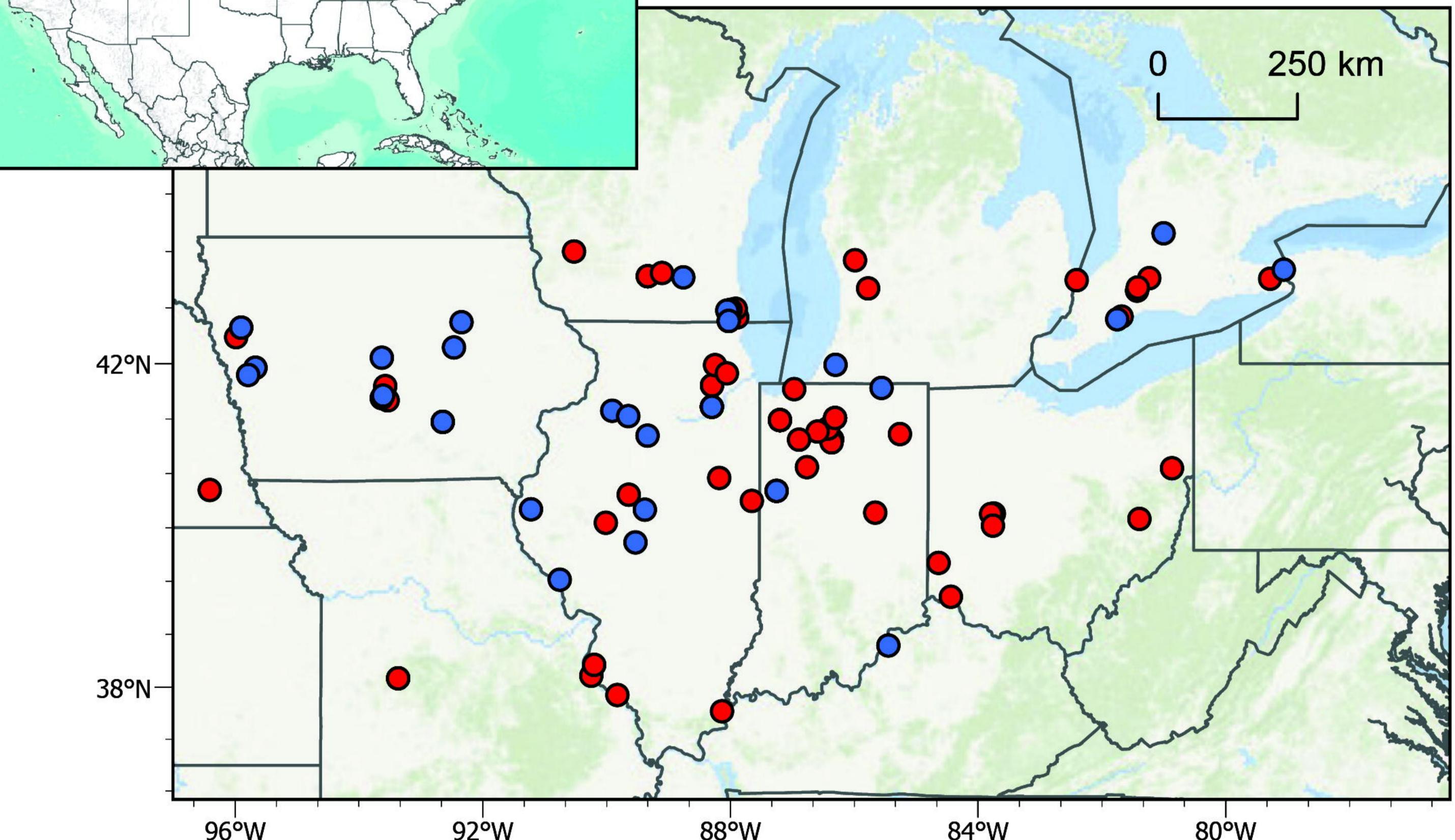
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## **Micromilled downtooth** samples

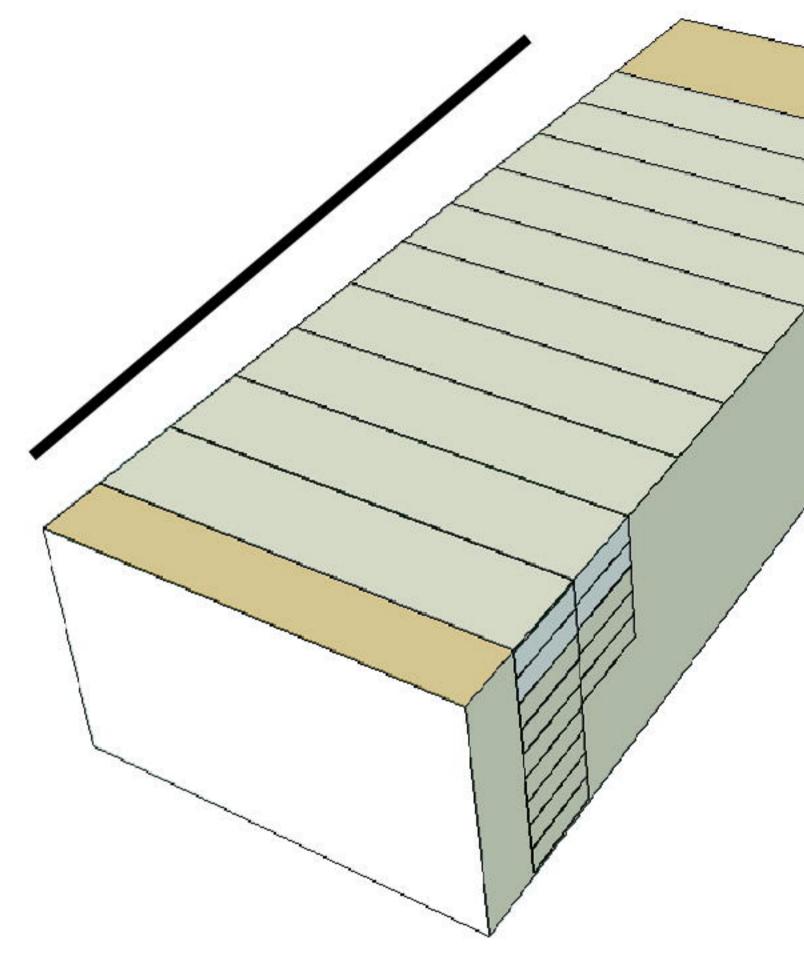
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- Set 2 (10 samples)

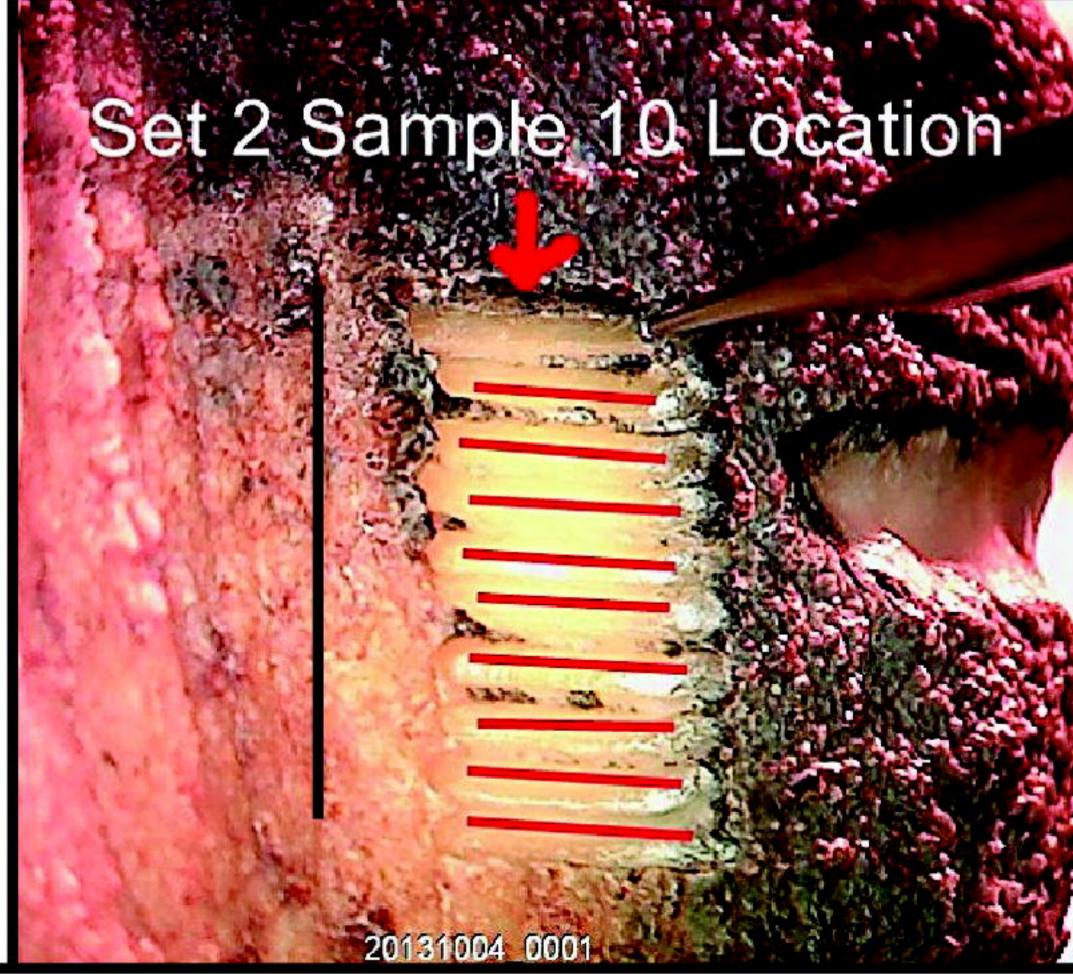
- Set 3 (10 samples)

Set 4 (10 samples)

Cross Section of enamel sampling set showing milling depth.

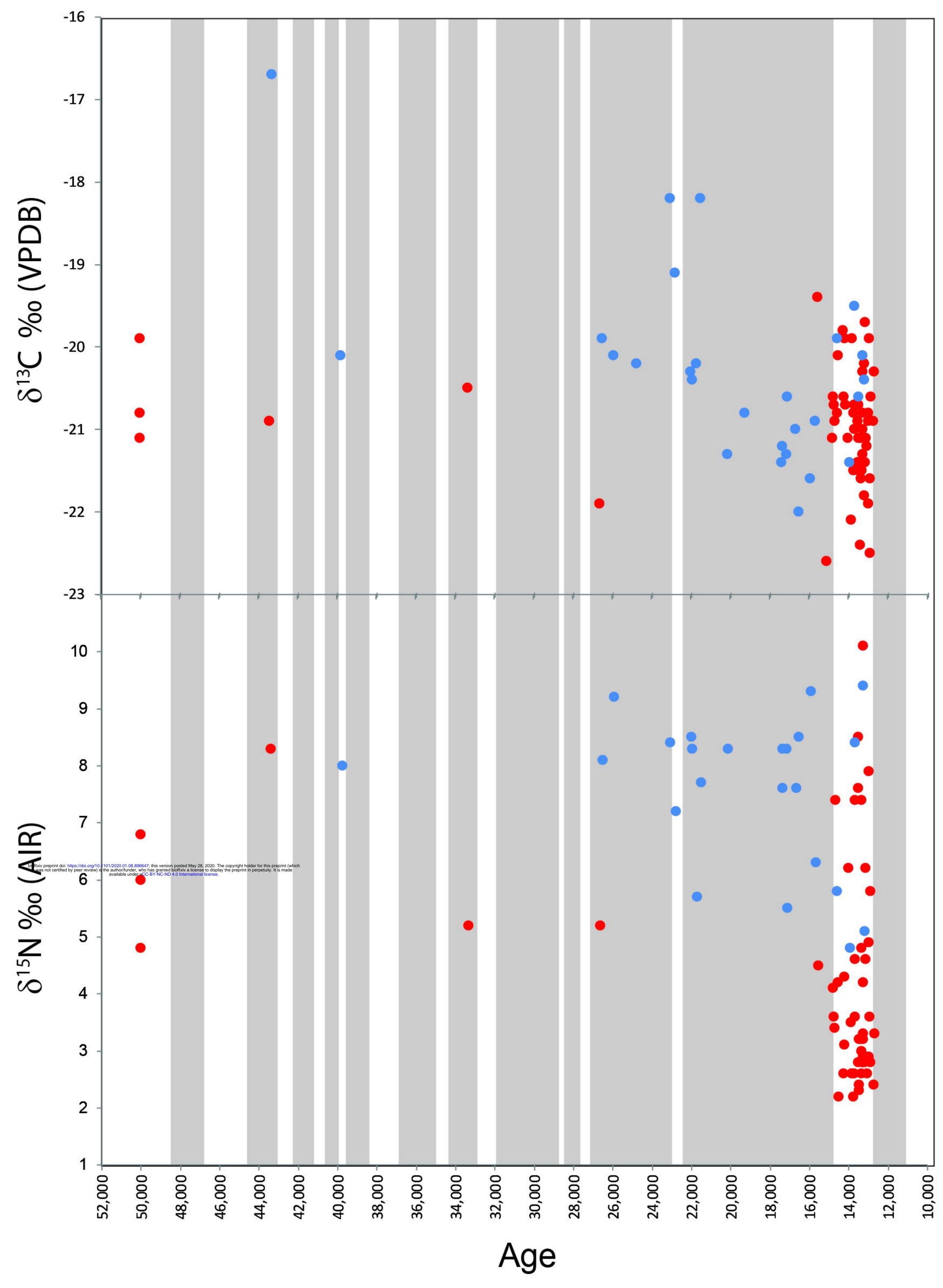


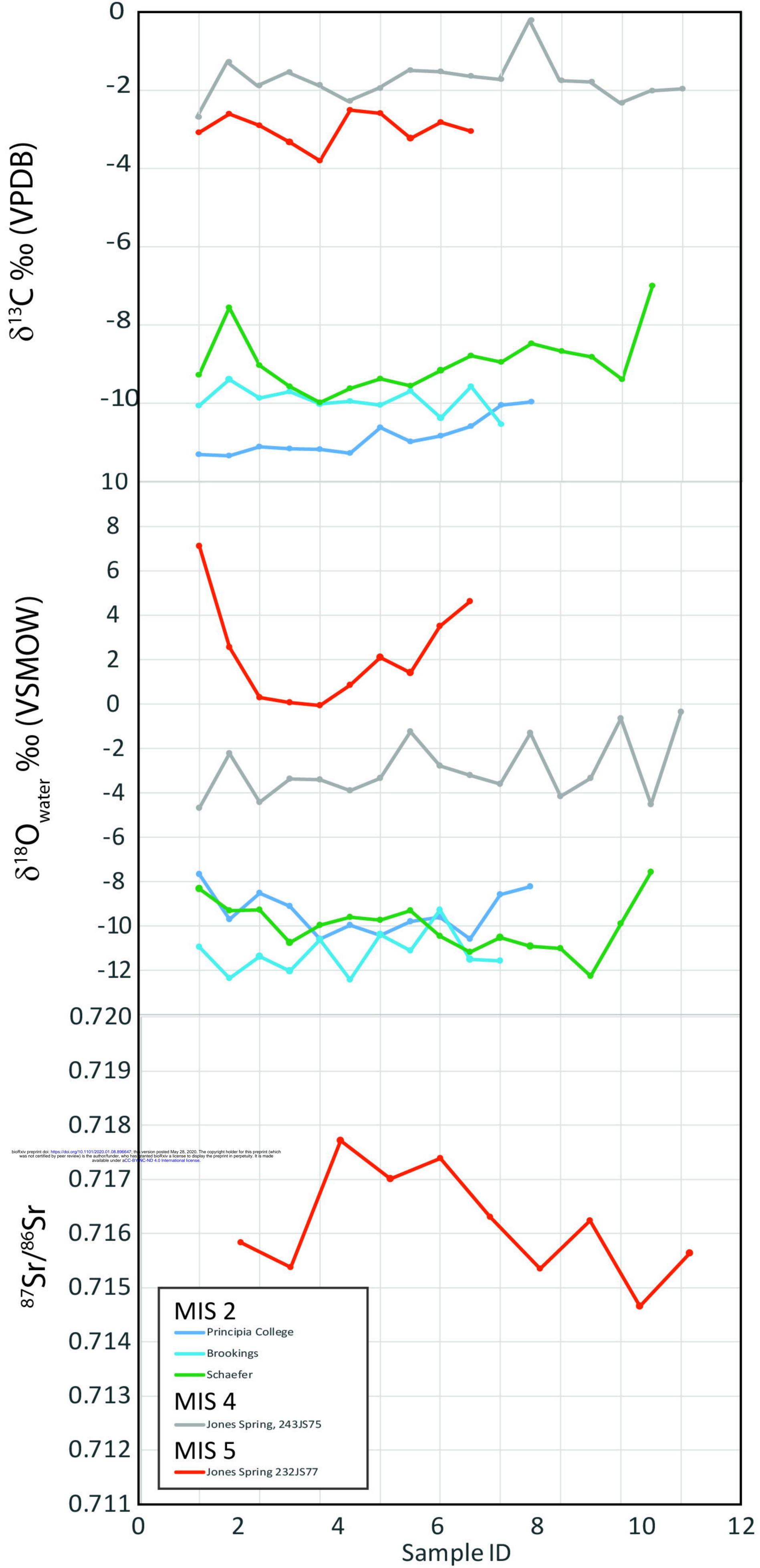
## Enamel Sampling set.



samples)







8<sup>13</sup>C % (VPDB)

