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Deforestation caused abrupt shift in Great Lakes nitrogen cycle

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Abstract

Despite the longstanding significance of North America's Great Lakes, little is known about their pre-industrial ecology. Here, we report on when and how humans first became a main driver of Lake Ontario's nutrient dynamics. Nitrogen isotope analyses of archaeological fish show that, prior to the 1830s, Lake Ontario's nitrogen cycle and the trophic ecology of its top predators had remained stable for at least 800 years, despite Indigenous and historical European agricultural land management across the region. An abrupt shift in the nitrogen isotope composition of Lake Ontario's fish community is evident in the early to mid-nineteenth century and reflects the initiation of industrial-scale forest clearance. These data show how the nitrogenous nutrient regimes of even the world's largest freshwater ecosystems can be highly sensitive to short-term watershed forest cover disturbances and indicate a profound shift in the relationship between humans and their environment.

Keywords

Historical Ecology, Archaeology, Great Lakes, Nitrogen Cycle, Fish, Deforestation

37 Introduction

38 Throughout the twentieth century, increased anthropogenic contributions of reactive nitrogen to
39 freshwater environments have profoundly altered aquatic biogeochemical cycles and broader
40 nutrient dynamics, resulting in widespread ecosystem dysfunction, loss of biodiversity, and
41 degraded water quality (Smith and Schindler 2009). While numerous studies have documented
42 how increasing anthropogenic nitrogen inputs are incorporated into aquatic food webs,
43 promoting cultural eutrophication and altering trophic dynamics in the recent past (Schindler, et
44 al. 2006), less consideration has been given to how historical and preindustrial populations have
45 impacted aquatic ecosystems. In the context of recent debate about the timing of the origins of
46 the Anthropocene (Lewis and Maslin 2015), in which past societies are considered as potential
47 architects of the first broad-scale environmental changes, the role of humans as drivers of
48 biogeochemical processes such as the nitrogen cycle is becoming increasingly important (Guiry,
49 et al. 2018; Hadley, et al. 2010; Kintigh, et al. 2014).

50 For conservation and restoration efforts in freshwater ecosystems, where cultural eutrophication
51 caused by increased nutrient loading poses one of the most significant threats globally (Smith
52 and Schindler 2009), long-term retrospectives that document when and how human activities
53 first began to alter natural ecosystem processes, and the nitrogen cycle in particular, could
54 provide a valuable framework for evaluating which modern human activities pose the greatest
55 risks (Canfield, et al. 2010; Humphries and Winemiller 2009). Paleolimnological proxy
56 indicators based on the physical, biological, chemical, and isotopic compositions of sediments
57 have long been used to establish trends in past aquatic productivity and water quality (Beeton
58 1965; Hodell and Schelske 1998; Jeffers, et al. 2015; Schelske, et al. 1983), but, because of
59 potential taphonomic issues (Anderson 2014; Lu, et al. 2014) and because these approaches

60 usually do not incorporate consumers, they are unable to measure the direct impacts of
61 anthropogenic nitrogen loading on nutrient dynamics in the wider biotic community (e.g.,
62 invertebrates, fish, birds; although for a growing literature on invertebrate analyses see Anas, et
63 al. 2019; Frossard, et al. 2014; Perga 2010; Perga, et al. 2010; Schilder, et al. 2017; van
64 Hardenbroek, et al. 2010). Moreover, while many studies using isotopic analyses of museum-
65 archived vertebrate tissues have been able to make important contributions to understanding how
66 biotic communities have responded to human-caused environmental change during the twentieth
67 century (Fera, et al. 2017), a lack of suitable specimens from early historical and pre-industrial
68 time periods has, in most cases, prevented analyses of longer-term environmental variation in
69 vertebrate taxa (Szpak, et al. 2018). In this context, stable nitrogen isotope and other analyses of
70 ancient fish remains from archaeological repositories can provide an invaluable opportunity to
71 gain direct insight into how food web and nutrient dynamics within pre-industrial ecosystems
72 functioned and responded to impacts from human activities at varying temporal and spatial
73 scales (Guiry, et al. 2016a).

74 Using isotopic compositions of fish bone or scales has some interpretive advantages relative to
75 more commonly analyzed materials, such as bulk sedimentary organic matter, because fish stable
76 nitrogen isotope composition is linked firmly with a known ecological point (i.e. a particular
77 species with well-understood ecology and trophic affinity) and spatiotemporal framework
78 (Trueman and Moore 2007) and can be easily parsed to remove diagenetically altered samples
79 using robust quality-control criteria (DeNiro 1985; Szpak 2011). This is important because, in
80 contrast to analyses of bulk sedimentary organic matter, which is composed of a spatially,
81 temporally, and biologically heterogeneous mixture of sources of detritus (Lu, et al. 2014), with
82 fish bone or scales, the ability to compare “taxonomically anchored” isotopic patterns across

83 different parts of a food web will enable much higher-resolution interpretations of environmental
84 change at both the level of nutrient dynamics and the level of broader food web structures.
85 Moreover, with respect to measuring the isotopic compositions of ancient organic materials, the
86 ability to purify collagen extracted from bone (through pretreatment steps outlined below) and
87 assess the extent to which its constituent carbon and nitrogen are biogenic (e.g., using well-
88 established and precise $C:N_{\text{Atomic}}$ criteria (DeNiro 1985; Szpak 2011) provides an additional
89 advantage relative to other important paleontological materials (e.g., chitinous remains from sub
90 fossil invertebrates; Anas, et al. 2019) and makes bone and scale collagen particularly well suited
91 for inter-site comparisons of isotopic variation in biota across time and space. Moreover, because
92 collagen from bones and scales is constructed and remodeled slowly throughout the life of an
93 organism, the isotopic composition of collagen can provide a lifetime average record of dietary
94 intake and environmental conditions that is less susceptible to seasonal or short-term
95 idiosyncratic shifts in behavior (Hobson and Clark 1992). Bone collagen is therefore well suited
96 for reconstructing long-term trends in biology and ecology (Bump, et al. 2007) and can provide
97 an additional perspective with potential to complement and build on the success of previous,
98 well-established paleolimnological proxy indicators (e.g., physical, chemical, and isotopic
99 compositions of sediments and invertebrates; Beeton 1965; Hodell and Schelske 1998; Jeffers, et
100 al. 2015; Perga 2010; Schelske, et al. 1983).

101 We use stable nitrogen isotope compositions and peptide mass fingerprinting (Zooarchaeology
102 by Mass Spectrometry, hereafter, ZooMS) of late Holocene (900-2015 CE) fish bone collagen
103 from a large number of sites associated with Lake Ontario (Figure 1), the most easterly of the
104 Laurentian Great Lakes (hereafter, Great Lakes), to document long-term trends in regional
105 biogeochemical cycles. Results show that, in comparison with the industrial era, the nitrogen

106 cycle and trophic structure of this Great Lakes ecosystem remained remarkably stable until the
107 1830s, despite millennia of Indigenous agricultural and other land management, decades of
108 European settlement, and climatic fluctuations. After this time, increased N-loading from
109 forestry- and agriculture-induced soil erosion caused an unprecedented and abrupt bottom-up
110 shift throughout the entire aquatic ecosystem of Lake Ontario.

111 Context

112 With more than 20% of the world's surface freshwater, North America's Great Lakes are of
113 tremendous ecological, economic, and social interest (Sturner, et al. 2017). Since monitoring
114 programs began, the Great Lakes have experienced significant ecological and chemical changes
115 in response to pollution, hydrological controls, and species invasion (Allan, et al. 2013; Stewart,
116 et al. 2016). Paleolimnological studies show that recent industrial activities have had profound
117 impacts on Great Lakes productivity (Beeton 1965; Hodell and Schelske 1998; Jeffers, et al.
118 2015; Meyers 2003; Schelske and Hodell 1991b; Schelske and Hodell 1995; Schelske, et al.
119 1988; Schelske, et al. 1983), especially through increased phosphorus and nitrogen loading that
120 led to eutrophication of Lake Ontario and Lake Erie by the twentieth century (Hodell and
121 Schelske 1998; Lu, et al. 2010). With respect to nutrient dynamics, particularly the nitrogen
122 cycle, which plays a vital role in primary production in naturally oligotrophic ecosystems such as
123 Lake Ontario (Lean and Knowles 1987; Leggett, et al. 2000), the extent to which these changes
124 represent a departure from longer-term trends in broader food web structure and nutrient
125 dynamics remains unclear (Sturner, et al. 2017; Supplementary Information [SI] 1). But this
126 extent could be determined through isotopic reconstructions of past nutrient and food web
127 dynamics using historical or archaeological fish specimens, as these can provide a
128 complementary isotopic record for ancient environmental variation.

129 Human activities have modified the cycling of nitrogen in aquatic environments across the globe,
130 contributing to the ongoing threat of cultural eutrophication (Gruber and Galloway 2008; Smith
131 and Schindler 2009). The majority of these impacts occur when anthropogenic nitrogen inputs
132 and/or chemical or physical changes to an environment alter the conditions regulating the
133 nitrogen cycle, thereby disrupting the flow of nitrogenous nutrients through an ecosystem
134 (Kendall, et al. 2007). Modeling how nitrogen moves into and through aquatic ecosystems is,
135 therefore, of considerable importance for conservation efforts seeking to restore heavily
136 impacted watersheds (Denk, et al. 2017). With respect to broad-scale human impacts, such
137 factors as climate change, acid rain, nutrient loading, and invasive species introductions can
138 significantly alter the biogeochemical processes that balance an aquatic ecosystem's nitrogen
139 cycle and often, in turn, cause detectable changes in the isotopic composition of nitrogen pools
140 that are available to support aquatic life (Anderson and Cabana 2005; Botrel, et al. 2014; Lake, et
141 al. 2001). Stable nitrogen isotopic compositions of consumer tissues show a stepwise increase,
142 by roughly 3–4‰, in $\delta^{15}\text{N}$ values with each trophic level step up in a food web and have
143 therefore traditionally provided a powerful indicator for food web interactions in ecology and
144 archaeology (Post 2002). However, because any impact on the $\delta^{15}\text{N}$ of aquatic nitrogen pools at
145 the base of a food web (i.e., in various pools of dissolved inorganic nitrogen [DIN]) is passed up
146 the trophic ladder to producers (phytoplankton) and their consumers (invertebrates and fish),
147 isotopic analyses of tissues from aquatic producers and consumers can also serve as a highly
148 integrative indicator for anthropogenic impacts on nutrient dynamics, in addition to food web
149 structure, in aquatic environments (Hoffman, et al. 2012; Morrissey, et al. 2013). While a
150 majority of isotopic research has approached the question of changing freshwater nutrient
151 dynamics through analyses of sedimentary organic matter (e.g., Dubois, et al. 2018; Talbot

152 2001), a growing number of studies are demonstrating that $\delta^{15}\text{N}$ compositions of invertebrates
153 and fish can provide a highly sensitive record for environmental change (e.g., Anas, et al. 2019;
154 Fera, et al. 2017; Lumb and Johnson 2012; Perga, et al. 2010).

155 The natural abundance of ^{15}N in DIN is largely controlled by three factors (for reviews see,
156 Finlay and Kendall 2007; Guiry 2019; Kendall 1998; Kendall, et al. 2007; Talbot 2001):
157 productivity (Hodell and Schelske 1998; Schelske and Hodell 1991a), environmental conditions
158 (temperature, oxygenation, and pH; Finlay and Kendall 2007; Knowles 1982), and nitrogen
159 inputs (quantity and isotopic composition; Heaton 1986; Lake, et al. 2001). These factors are
160 necessarily interrelated because, for instance, increasing the input of reactive nitrogen can
161 increase productivity, and this can affect environmental conditions that regulate other parts of the
162 nitrogen cycle (Finlay and Kendall 2007). Moreover, nitrogen-limitation rates can influence
163 plankton community composition, which is an integral component of determining the presence
164 of nitrogen-fixing bacteria (Gu 2009) that can further modify the isotopic composition of an
165 aquatic ecosystem by introducing isotopically distinctive atmospheric nitrogen. Human activity
166 should affect those factors controlling the nitrogen cycle and its isotopic composition (for review
167 see Guiry 2019) by: 1) contributing reactive nitrogen (i.e., through inputs directly from sewage,
168 agriculture, and soil erosion – typically, but not always, leading to elevated biota $\delta^{15}\text{N}$ (Anderson
169 and Cabana 2005; Morrissey, et al. 2013)), which itself leads to increasing productivity (creating
170 greater nitrogen demand and thereby possibly promoting the importance of N-fixing algae –
171 leading to higher and lower biota $\delta^{15}\text{N}$, respectively (Gu 2009; Pennock, et al. 1996)), and 2)
172 changing chemical conditions, such as pH and oxygen levels, that regulate transformations
173 between important forms of reactive nitrogen (altering the dynamics for nitrification and

174 denitrification – with potential to push biota $\delta^{15}\text{N}$ higher or lower (Collister and Hayes 1991;
175 Sebito, et al. 2006)).

176 With the longest history of European and Indigenous agricultural land management and
177 settlement among the Great Lakes, Lake Ontario and its watershed are an ideal context in which
178 to investigate how large-scale human activities can impact major freshwater environments over
179 time (Hodell and Schelske 1998). Because Lake Ontario receives a large amount of its water
180 from land runoff, it is particularly sensitive to broader ecological processes and human
181 disturbances occurring across the watershed, particularly in terrestrial and riparian zones.
182 Moreover, the significant and highly visible environmental degradation of Lake Ontario's
183 watershed throughout the twentieth century, as well as its close proximity to major population
184 centers, has spurred considerable research into its ecological structure (e.g., Bogue 2001; Estep
185 and Reavie 2015; Meyers 2003; Stewart, et al. 2016). For this reason, the lake has a relatively
186 well-documented ecological and biogeochemical history, with numerous isotopic studies of
187 modern fish identifying recent trends in community structure, trophic dynamics, and energy
188 flows (Colborne, et al. 2016; Fera, et al. 2017; Kiriluk, et al. 1995; Lumb and Johnson 2012;
189 Rush, et al. 2012; Yuille, et al. 2015). The longest-term of these studies, however, only offers a
190 70-year retrospective, one that postdates intensification of European settlement of Lake Ontario's
191 watershed by more than 150 years.

192 While a number of important isotopic studies have helped to document longer-term variability in
193 Lake Ontario's nitrogenous nutrient dynamics (SI 1), these have focused on isotopic
194 compositions of organic matter in lake sediments rather than specific taxa. With respect to the
195 last millennium, the timeframe over which human land management would have intensified,
196 most (95%) of these analyses focus on the period after the mid-nineteenth century and show

197 substantial change in the nitrogen isotopic composition of organic matter settling out of Lake
198 Ontario's water column (SI 1; Figure S1, Figure 2; Hodell and Schelske 1998; McFadden, et al.
199 2004). Only a small fraction (5%; $n=8$ of 178) of analyses cover the early historical period
200 (1600s-1850 CE), when European activities would have started impacting the region, and pre-
201 contact period, when Indigenous farming and other land management activities were accelerating
202 across the watershed (i.e., starting with the Late Woodland, c. 900 CE). Moreover, only one lake
203 sediment isotopic dataset spans both time frames and is not consistent with the others (SI 1,
204 Figure 2). While these data provide an invaluable macro-scale perspective on general patterns in
205 nutrient dynamics in Lake Ontario, they cannot be used to assess the consequences of shifting
206 nitrogenous nutrient regimes for the broader food web (invertebrates and fish). Moreover, the
207 coarse temporal resolution of the lake sediment isotopic record prior to 1850 prevents detailed
208 analyses of when the human activities began to impact the nitrogen cycle and to what extent
209 early historical and Indigenous activities may have been important drivers for change. In this
210 context, isotopic analyses of archaeological fish can provide a complementary line of evidence
211 for nutrient dynamics and food web structure.

212 Materials and methods

213 *Sample description:* Sampling focused on three taxa in the family Salmonidae: lake trout
214 (*Salvelinus namaycush*; $n=222$), Atlantic salmon (*Salmo salar*; $n=158$), and whitefishes
215 (*Coregonus* spp.; primarily *C. clupeaformis* and *C. artedi*, see below; $n=277$; SI 2, Table S1).
216 These taxa were selected to provide a highly integrated record of changes to Lake Ontario's
217 nitrogen cycle and trophic structure. The upper trophic position and longer lifespan (5–15 years)
218 as well as pelagic foraging and highly mobile behavior of these species means that their diets
219 will incorporate a broad range of resources from across the lake (Fera, et al. 2017; Guiry, et al.

220 2016a; Rush, et al. 2012), providing a time-averaged perspective on processes affecting the
221 nitrogen cycle at a lake-wide, multi-seasonal scale. A comparison of isotopic variation among
222 three taxa, each with a different behavioral strategy and trophic position (Fera, et al. 2017; Holm,
223 et al. 2009; Mumby, et al. 2018), should make isotopic trends associated with dietary shifts at the
224 species level (as opposed to changes at the base of the food web) clearly discernible and,
225 therefore, these data also give a long-term perspective on the stability of Lake Ontario's food
226 web structure.

227 With the exception of nine Atlantic salmon samples, all archaeological specimens are from sites
228 within the Lake Ontario watershed. As Lake Ontario is the only known source of Atlantic
229 salmon, specimens from this species at sites outside the watershed were also included because
230 they represent individuals originating from Lake Ontario. Where possible, archaeological fish
231 bone specimens were selected based on minimum number of individual counts per
232 archaeological context in order to avoid sampling the same individual multiple times. For all
233 species, archaeological sampling efforts targeted bone specimens from adult-sized fish in order
234 to exclude juveniles, who may feed at a lower trophic level. All specimens were examined by
235 one or more of three zooarchaeologists (S.N.-H., A.H., T.O.) with specific expertise in the
236 identification through comparative osteology of archaeological fish bones from relevant taxa in
237 the Great Lakes region (Hawkins, et al. 2019). The archaeological samples are largely derived
238 from fish vertebra, because they are more ubiquitous than fish cranial and pectoral girdle bones
239 in the archaeological collections. Zooarchaeologists determined species for each specimen where
240 possible, based on clear morphological differences between taxa. Due to overlaps in osteology
241 and the possibility of hybrids, morphological examination of *Coregonus* vertebrae and many
242 *Coregonus* cranial and pectoral girdle bones cannot be confidently used to assign species-level

243 identification in the Lake Ontario watershed. However, of the five *Coregonus* species that are
244 native to Lake Ontario, only two, lake whitefish (*C. clupeaformis*) and cisco (*C. artedi*), inhabit
245 waters shallow enough to have been broadly accessible using fishing technologies (birch bark
246 canoes and gill nets) available in the pre-contact past, and it is therefore highly likely that
247 archaeological *Coregonus* of that time are composed of these taxa. Some bones in the genera
248 *Salvelinus* and *Salmo* are also morphologically similar to each other, resulting in additional
249 higher-level taxonomic identifications (including to *Salmo salar*/*Salvelinus namaycush* and
250 Salmonidae). In cases where a species-level taxonomic identification was not achieved with a
251 high degree of certainty, we undertook ZooMS analyses ($n=292$) to confirm identifications,
252 comparing the archaeological samples with reference sequences we generated for *S. salar*, *S.*
253 *namaycush*, brook trout (*Salvelinus fontinalis*), round whitefish (*Prosopium cylindraceum*), *C.*
254 *clupeaformis*, and *C. artedi* (Supporting Information 4 Table S1).

255 Dating for pre-contact specimens is based on published (SI 2, Table S2) radiocarbon dates,
256 ceramic seriations, and reconstructed village occupation sequences (e.g., Williamson 2014), to
257 occupations within 100-year bins. Excellent chronological control for data from Indigenous
258 archaeological sites reflects the highly standardized lifecycle in which settlements were built,
259 occupied, decommissioned, and left behind in favor of relocation every 10–30 years (Warrick
260 2008). The relative dearth of data from the seventeenth and eighteenth centuries reflects changes
261 in Indigenous settlement patterns during this period. Reduced sample sizes in the nineteenth
262 century reflect lower frequencies for fish bone at Euro-Canadian archaeological sites. Historical
263 fish scale samples of adult specimens of known date were taken from museum-archived
264 collections with assistance from an Ontario fish identification expert (E.H.). Analyses of scales
265 from museum-archived *Coregonus* specimens include both *C. clupeaformis* and *C. artedi*

266 because: 1) these two species likely account for the majority of archaeological specimens and
267 will therefore be directly comparable; and 2), based on $\delta^{15}\text{N}$ values, they appear to have fed at
268 the same trophic level in early twentieth-century Lake Ontario. For data sourced from the
269 literature ($n=834$; (Colborne, et al. 2016; Fera, et al. 2017; Guiry, et al. 2016a; Kiriluk, et al.
270 1995; Rush, et al. 2012; van der Merwe, et al. 2003; Yuille, et al. 2015)), fork length, when
271 available, was used to exclude juveniles, which may feed at a lower trophic level.

272 *Sample preparation:* Samples weighing between 30 and 300 mg were cut from bone specimens,
273 residual bone lipids were removed with a series of 2:1 chloroform:methanol ultrasonic baths
274 (solution refreshed every 10 minutes until visible signs of reaction ceased), and the samples were
275 then left to air dry (Guiry, et al. 2016b). Bone samples were then demineralized in 0.5 M
276 hydrochloric acid (HCl) for several days at 4°C and then rinsed to neutrality in Milli-Q water.
277 Humic acids and other base soluble contaminants from the burial environment were removed
278 from demineralized bone samples with successive treatments in 0.1 M sodium hydroxide in an
279 ultrasonic bath (solution refreshed every 15 minutes until visible signs of reactions ceased) and
280 then rinsed to neutrality in Milli-Q water (Szpak, et al. 2017a). Bone samples were then
281 solubilized in 10^{-3} M HCl in an oven at 65°C for 48 hours. The resulting solutions were
282 centrifuged, after which the solubilized collagen fraction was transferred to a new tube and then
283 frozen and lyophilized.

284 The protein fraction of fish scale is also composed primarily of collagen, which is
285 compositionally and isotopically comparable to fish bone collagen (Guiry, et al. 2016a; Trueman
286 and Moore 2007). Fish scales samples underwent a succession of three rinses in Milli-Q water in
287 an ultrasonic bath for five minutes each (Guiry, et al. 2016a). Scale samples were then soaked for
288 five minutes in 1.0 M HCl in an ultrasonic bath to remove the mineral phase of the scale's

289 external plate as well as any microscopic debris that may have become adhered to the external
290 surfaces of the fish specimens during preservation and storage. Scale samples were then rinsed to
291 neutrality in Milli-Q water in an ultrasonic bath and left to air dry. The historical fish specimens
292 had been preserved using formalin fixation and then stored in ethanol. While this form of
293 preservation is known to introduce small quantities of carbon, which can slightly alter the stable
294 carbon isotope composition of biological tissues, formalin and ethanol provide no new sources of
295 nitrogen and therefore have a minimal influence the nitrogen isotopic composition of preserved
296 fish scales. To confirm this, we also compared the percent carbon-to-nitrogen ratio (C:N) of
297 historical scales with that of modern fish scales (see below; (Szpak 2011)) to establish that
298 diagenetic nitrogen is not present.

299 *Isotopic analyses:* The stable nitrogen isotope composition as well as percent carbon and
300 nitrogen were measured on 0.5 mg samples of bone and scale collagen using an Elementar Vario
301 MICRO cube elemental analyzer coupled via continuous flow to an Isoprime isotope ratio mass
302 spectrometer in the Archaeology Chemistry Laboratory at the University of British Columbia,
303 Canada. Duplicate or triplicate analyses were performed on 14% of samples. For a full account
304 of procedures used for calibration of isotopic values as well as isotopic uncertainty calculations
305 see SI 3. Analytical uncertainty for $\delta^{15}\text{N}$ measurements was +0.20‰ (Szpak, et al. 2017b; SI 3).
306 The following data quality criteria were applied for establishing the integrity of $\delta^{15}\text{N}$
307 measurements from bone and scale collagen (DeNiro 1985): 1) % carbon and % nitrogen values
308 above 13% and 4.8%, respectively; and 2) atomic C:N values falling between 2.9 and 3.6.

309 *ZooMS analyses:* Collagen was rehydrated with approximately 100 μL 50 mM ammonium
310 bicarbonate per milligram and each digested with 0.4 μg sequencing grade trypsin (Promega,
311 UK) overnight at 37°C. Initial attempts were made at directly spotting these digests through

312 dilution in 0.1% trifluoroacetic acid (TFA; Buckley, et al. 2018), but many required ZipTip
313 purification with C18 solid phase extraction and rehydration in 0.1% TFA (Buckley, et al. 2009).
314 Samples were spotted onto a stainless-steel Matrix Assisted Laser Desorption Ionization
315 (MALDI) target plate with equal volume of 10 mg/mL alpha-cyano hydroxycinnamic acid in
316 50% acetonitrile/0.1% TFA and allowed to air dry. Dried spots were then analyzed using a
317 Bruker Ultraflex II MALDI mass spectrometer over the m/z range 700-3700 with up to 2,000
318 laser acquisitions for each sample. Peptide mass fingerprints were then compared with those of
319 standard reference material via the identification of genus- or species-specific biomarkers (SI 4).

320 *Statistical analyses:* For statistical comparisons of fish $\delta^{15}\text{N}$ values, each taxon (i.e., lake trout,
321 Atlantic salmon, whitefishes) was compared separately by time periods binned by 100 (pre-1925
322 CE) and 25 (post-1925 to present) year groups (SI 5, Table S1) using PAST version 3.22. Bin
323 timeframes were selected with a view to maximizing sample size per bin. To assess whether fish
324 $\delta^{15}\text{N}$ values changed over time we first compared pre-1800 bins to one another to establish
325 whether significant differences occur within each fish taxon between sequential time frames.
326 After finding that no significant differences occur (SI 5, Table S2), we then grouped all pre-1775
327 (preindustrial) bins and compared this individually to all succeeding time periods (SI 5, Table
328 S3). For each bin, normality of distribution was first tested using a Shapiro–Wilk’s test (SI 5,
329 Table S1). For datasets where all time periods were distributed normally, a one-way ANOVA
330 was used, and homogeneity of variance was assessed using a Levene’s test. A post hoc Dunnett’s
331 T3 test or a post hoc Tukey’s HSD test was used to compare groups with variances that were
332 equal or unequal, respectively. For data sets where not all time periods were distributed
333 normally, Mann-Whitney U tests with Bonferroni corrected p values were used to compare
334 groups.

335 Results

336 Of the samples with sufficient collagen ($n=607$ of 662) for isotopic analyses, 96% (total $n=582$)
337 produced acceptable C:N, %C, and %N values satisfying quality control criteria. Interpretations
338 also include previously published data from archaeological ($n=68$; (Guiry, et al. 2016a; van der
339 Merwe, et al. 2003)) and modern (post-1958 CE; $n=819$; (Colborne, et al. 2016; Fera, et al. 2017;
340 Kiriluk, et al. 1995; Lumb and Johnson 2012; Mumby, et al. 2018; Rush, et al. 2012; Yuille, et
341 al. 2015)) specimens. ZooMS analyses of 290 samples confirmed or helped to refine 244 non-
342 species taxonomic identifications provided through bone morphology analyses by
343 zooarchaeologists (it confirmed the identification for 225 of these 244 specimens and refined it
344 to the level of species for the remaining 19). In a further 20 cases, specimens were reassigned to
345 another taxon, namely, 12 to lake trout (*Salvelinus namaycush*) (previously assigned with
346 varying degrees of confidence to the species or genus level in *Salmo* [$n=8$], *Salvelinus* [$n=2$], or
347 *Coregonus* [$n=2$]); four to Atlantic salmon (*Salmo salar*) (previously identified with varying
348 degrees of confidence as *Salvelinus namaycush*); four to whitefish (*C. clupeaformis* or *C. artedi*)
349 (previously assigned with varying degrees of confidence to the species or genus level in *Salmo*
350 *salar* [$n = 1$], *Salvelinus fontinalis* [$n = 2$], or *Salvelinus namaycush* [$n = 1$]). In 15 other cases,
351 ZooMS could not reassign specimens (previously assigned with varying degrees of confidence to
352 the Salmonidae family level [$n = 3$] or to the species or genus level in *Salmo* [$n=4$], *Salvelinus*
353 [$n=1$], or *Coregonus* [$n=4$]) to one of the four species and one genus used in this study. Based on
354 variation in peptide mass fingerprints, these samples likely derive from at least eleven other
355 species.

356 Stable nitrogen isotope compositions of bone and scale collagen from archaeological and
357 historical lake trout ($n=196$), Atlantic salmon ($n=147$), and whitefishes ($n=239$) are summarized

358 in the SI 5, Table S1 and presented in full in Figure 2 and SI 2. Given the large and well-dated
359 sample ($n=1469$) from archaeological sites ($n=42$) and from museum-archived specimens and
360 contemporary fisheries research ($n<50$ locations) across a broad geographical range, we believe
361 that the temporal trends observed in fish bone and scale collagen $\delta^{15}\text{N}$ are representative of
362 processes occurring in Lake Ontario's open water ecosystem at a regional scale.

363 Discussion and conclusion

364 *1200–1830 CE*: Fish $\delta^{15}\text{N}$ shows remarkably little variation between 1200 CE and 1830 CE
365 (Figure 2), despite centuries of Indigenous and European land management in the Lake Ontario
366 watershed, and provides a unique perspective on the long-term stability of a Great Lakes
367 ecosystem. Prior to the nineteenth century, mean $\delta^{15}\text{N}$ values binned by taxon and at 100-year
368 intervals vary by less than 0.4‰ and show no statistically significant differences (SI 5, Tables S1
369 and S2). This long-term stasis, over at least 600 years, suggests that the nitrogen cycle and
370 nitrogenous nutrient inputs for Lake Ontario were highly stable during the past millennium.
371 Analyses of two additional Atlantic salmon specimens (not shown in Figure 2) from earlier
372 archaeological deposits suggests that this stability extends further back in time to at least 900 CE.
373 However, due to a paucity of samples from the earlier time period, discussion will focus on the
374 post-1200 CE time frame. A comparison of mean $\delta^{15}\text{N}$ for pre-1800 bins ($n=4$ see SI 5, Table
375 S1) shows steady offsets between species: lake trout $+1.6\pm 0.2\text{‰}$ > Atlantic salmon $+1.3\pm 0.2\text{‰}$ >
376 whitefishes. Replication of this pattern across all three taxa, encompassing multiple trophic
377 levels, suggests that stability was also characteristic of interspecific trophic relationships
378 throughout the broader pelagic food web during this period.

379 A wealth of historical and paleoenvironmental research on the Lake Ontario watershed (Beeton
380 1965; Bogue 2001; Estep and Reavie 2015; Hodell and Schelske 1998; Jeffers, et al. 2015;

381 Meyers 2003; Schelske and Hodell 1991b; Schelske and Hodell 1995; Schelske, et al. 1988;
382 Schelske, et al. 1983; Wood 2000) provides an excellent opportunity to contextualize and
383 consider the implications of this long-term stability. Proxies for past environmental conditions in
384 other, smaller lakes in the same region document anthropogenic disturbances, at least as early as
385 the thirteenth century CE, resulting from Indigenous land clearance and farming activities (e.g.,
386 Ekdahl, et al. 2004). These studies suggest that Indigenous land management practices, including
387 the use of fire for forest clearance followed by crop cultivation (Munoz and Gajewski 2010), had
388 a significant impact on soil erosion, sedimentation, and nutrient loading, causing early cultural
389 eutrophication of aquatic environments in the region. European settlement of the Lake Ontario
390 watershed began in the seventeenth century (Gentilcore 1984). Settlement expansion in the
391 nineteenth century was accompanied by intensive commercial fishing (1800 onward; Bogue
392 2001), diversion of the Niagara River and work on the Erie and Welland canals (1820s;
393 McIlwraith 1976; Wood 2000), broad-scale land clearance for lumber and agriculture (1840s–
394 1850s; Head 1975; Lower, et al. 1938), and wetland loss due to mining and other activities
395 (1880s; Bogue 2001). Paleolimnological studies suggest that productivity of Lake Ontario’s
396 phytoplankton community began to grow slowly with the first European settlement, with more
397 substantial impacts occurring in the mid-nineteenth century, probably in response to nutrient
398 loading from soil erosion following deforestation (e.g., Estep and Reavie 2015; Hodell, et al.
399 1998; Schelske 1991; SI 1). Despite clear historical and paleolimnological evidence for earlier
400 anthropogenic impacts on nutrient dynamics (e.g., nitrogen and phosphorus loading, pollen
401 species change) in other areas of the watershed (Bunting, et al. 1998; Duthie and Sreenivasa
402 1971; Ekdahl, et al. 2004; Ekdahl, et al. 2007; Munoz and Gajewski 2010; Schelske, et al. 1983;
403 Yang, et al. 1993), archaeological fish stable nitrogen isotopic compositions demonstrate that

404 Lake Ontario's nitrogen cycle and food web structure remained unchanged until the 1830s. In
405 that context, and given the sensitivity of aquatic nitrogen cycles to human disturbances
406 (Anderson and Cabana 2005; Botrel, et al. 2014; Lake, et al. 2001), these data underscore Lake
407 Ontario's stability throughout much of the last millennium and demonstrate a degree of
408 resilience at the level of the nitrogen cycle in large lake ecosystems to a variety of human
409 impacts, even when sustained over long time spans.

410 The resilience of Lake Ontario's nitrogen cycle has important implications for understanding
411 human impacts on freshwater ecosystems in the recent and more ancient past. It is now well
412 known that industrial activities over the past 150 years have fundamentally altered global
413 nutrient distributions and biogeochemical cycles (Galloway and Cowling 2002). Within the
414 context of debate about the onset of the Anthropocene (Lewis and Maslin 2015), there is a
415 growing consensus that "pristine environments" probably did not exist where humans were
416 present, because there was always some form of impact from human activities (Heckenberger, et
417 al. 2003). This perspective has been supported by archaeological research showing that ancient
418 societies, particularly those that used agriculture, had significant impacts at both local and
419 regional scales on terrestrial and aquatic nutrient cycles for millennia (Curtis, et al. 1998; Guiry,
420 et al. 2018; Hadley, et al. 2010). Evaluating the role of past human activities as drivers of
421 ecological change has been a priority for archaeologists (Kintigh, et al. 2014), but is complicated
422 by the fact that ancient anthropogenic impacts on the environment are spatiotemporally
423 heterogeneous and occur along a continuum of intensity (from very significant to ephemeral).
424 For ecologists, this issue represents a source of uncertainty for retrospective studies where
425 archaeological or paleobiological information is not available to establish when and to what
426 extent past human activities have impacted a particular environment or biotic community

427 (Holmes 2006; Lyman and Cannon 2004). In contrast to the cautionary tone of the debate about
428 the existence of “pristine” environments, our data demonstrate that, in fact, nutrient dynamics
429 and trophic relationships in larger aquatic environments can remain unchanged despite centuries
430 of moderate human land use.

431 *Post-1830:* Our analyses of archaeological and historical fish also show that this long-term
432 stability was profoundly altered when Lake Ontario’s nutrient pools experienced an abrupt shift
433 in stable nitrogen isotope composition, becoming enriched in ^{15}N throughout the pelagic
434 ecosystem. During the early to mid-nineteenth-century, fish $\delta^{15}\text{N}$ values underwent a
435 simultaneous, statistically significant elevation across multiple trophic levels, a difference that
436 has been sustained throughout the twentieth century and to this day (SI 4, Table S3). Atlantic
437 salmon, for instance, which had previously had a highly conserved distribution of $\delta^{15}\text{N}$ values,
438 show an elevation of +3‰ in well-dated mid-nineteenth-century museum specimens. Although a
439 paucity of fish samples from the later nineteenth and early to mid-twentieth century precludes
440 detailed comparison, these data show similar overall trends (positive shifts between intraspecific
441 $\delta^{15}\text{N}$ means for pre-1800 and post-1990, ranging from +5.3‰ to +6.2‰, for all fish species) to
442 those observed in Lake Ontario’s isotopic records from sediment cores (+5.5‰ to +5.6‰ for c.
443 1850 to 1990s; Hodell and Schelske 1998; Figure 2). This further indicates that the observed
444 upward pattern in fish $\delta^{15}\text{N}$ predominately reflects a bottom-up shift in baseline $\delta^{15}\text{N}$ rather than
445 changes to fish trophic behavior.

446 It is nonetheless apparent from these data that human activities have resulted in meaningful
447 behavioral changes, including increased niche overlap between Lake Ontario’s salmonid
448 communities. Prior to 1800, there is minimal overlap between the $\delta^{15}\text{N}$ ranges for each species,
449 with overlaps of only 0.1‰ between Atlantic salmon and whitefishes and only 0.9‰ between

450 Atlantic salmon and lake trout over a 600-year period. In contrast, in the twentieth century, there
451 is substantial overlap between $\delta^{15}\text{N}$ ranges for all species in all 25-year temporal bins (overlap
452 ranges from 1.7 to 4.7‰). However, caution is required when interpreting trends in variation
453 across time in this dataset. Differences in tissue turnover rates for sample materials between pre-
454 nineteenth- (mainly collagen from bones – slow turnover) and twentieth-century (mainly
455 previously published muscle from modern fish – fast turnover) time frames make detailed
456 analysis of these trends difficult because tissues with faster turnover rates are inherently more
457 likely to produce greater isotopic variation reflecting seasonal or other short-term dietary
458 aberrations. With that caveat in mind, we argue that it is likely that the greater isotopic variation
459 in twentieth-century specimens reflects changes in energy pathways and foraging behavior in
460 response to major anthropogenic environmental disruptions (e.g., impacts from invasive species,
461 extirpations, or over fishing; Bogue 2001; Colborne, et al. 2016; Dymond, et al. 2019; Fera, et al.
462 2017; Mills, et al. 2003).

463 The interspecific synchronicity and magnitude of this shift could have been caused by multiple
464 related processes (Figure 3), including changes in the nitrogen cycle (at the phytoplankton-DIN
465 pool level) favoring nitrogen transformations that retain ^{15}N and changes in nitrogen inputs to the
466 system that include ^{15}N -enriched nutrients (Finlay and Kendall 2007; Kendall 1998; Kendall, et
467 al. 2007; Talbot 2001). Detailed historical and archaeological dating provides a robust temporal
468 framework to evaluate which of these variables, and therefore what type of human activity, drove
469 this initial ecosystem-wide ^{15}N enrichment in Lake Ontario's food web. At archaeological sites
470 with occupation dates as late as 1832, many fish show no sign of $\delta^{15}\text{N}$ elevation for their
471 respective taxon, whereas a museum specimen dating from no later than 1857 shows
472 significantly elevated $\delta^{15}\text{N}$ relative to the preceding centuries. In addition, many specimens from

473 archaeological contexts with date ranges spanning this *terminus ante quem* (1832) and *terminus*
474 *post quem* (1857) show elevated $\delta^{15}\text{N}$ values. These data therefore provide a temporal bracket for
475 the initiation of change in Lake Ontario's nitrogen cycle, demonstrating that the shift occurred
476 abruptly (within the context of the stability of the preceding 900 years), within a relatively short,
477 15–25-year window (1830s–1850s). This timeframe coincides precisely with the emergence of
478 broad-scale land clearance for agriculture and timber harvesting in the Lake Ontario watershed.
479 From the 1830s onward, especially around the middle of the century, historical analyses
480 document how forestry activities, namely severe cutting as well as widespread burning of the
481 remaining slash, radiated outward from the Lake Ontario waterfront to encompass much of the
482 watershed. These forestry activities accelerated over time due to increased demand for timber
483 (Head 1975) and cleared land, improved export capacity (canals and railroad; Lower, et al. 1938;
484 McIlwraith 1976), and growing sawmill infrastructure (Head 1975; Wood 2000).

485 During this period, between the 1820s and 1850s, the then-province of Upper Canada (which
486 included the Ontario portion of the Lake Ontario watershed) had the fastest growing population
487 in all of North America (Lewis 2001). Prior to 1830, newly settled farmers of European heritage
488 focused on subsistence (Kelly 1973; Kelly 1975) and put considerable effort into clearing land
489 for cultivation through cutting and burning (Ball 1979). However, relative to the mid-nineteenth
490 century, these early farming activities were patchy, small in scale, and diversified in purpose,
491 with most of the land granted to settlers remaining forested (Kelly 1975). Moreover, much of the
492 land that was cleared retained stabilizing features of the previous forest, including stumps, root
493 mats, and stones, for up to a decade, and sometimes much longer, while under cultivation and
494 these would have mitigated issues with soil erosion and nutrient loss (Ball 1979; Kelly 1975).
495 During this early period, in which the farmers were themselves clearing their land, care was

496 often taken throughout the processes, from cutting trees to sowing crops (delaying ploughing), to
497 further minimize soil nutrient loss (Ball 1979; Kelly 1971). As farms became established,
498 however, stumps were removed, ploughs were more widely employed to break up the soil, and
499 more land came under cultivation. Throughout the 1830s and 1840s, the agricultural system of
500 Upper Canada was dominated by plough-intensive (three or more times per biennial fallow) and
501 highly profitable monocropping of wheat (Kelly 1971; Kelly 1973). The focus on a wheat-
502 fallow-wheat agriculture system, as well as labor-saving furrow drainage and general avoidance
503 of manure-based crop fertilization (until the 1850s) were widely blamed for rapid depletion of
504 soil nutrients as well as erosion across the region (Kelly 1971; Kelly 1975). In this context, prior
505 to the 1880s, when a wider appreciation developed among farmers in the Lake Ontario watershed
506 of the risks of higher runoff-to-percolation ratios that come with overly cleared lands, farmland
507 was subjected to a damaging seasonal cycle of spring floods, washing away nutrient-rich organic
508 materials, and summer droughts, parching what soil remained (Kelly 1975).

509 The isotopic composition of reactive forms of nitrogen in terrestrial ecosystems (i.e., soil
510 nutrients including ammonia and nitrate), particularly those used for agriculture, is typically
511 elevated relative to aquatic nitrogen sources (i.e., DIN) due to intense soil denitrification (Botrel,
512 et al. 2014; Heaton 1986; Kendall 1998). In this context, the most parsimonious explanation for a
513 broad-scale isotopic shift in Lake Ontario's nitrogen cycle is that there was an increase in ¹⁵N-
514 enriched nutrient inputs coming directly from soil erosion following deforestation during the
515 1840s in Lake Ontario's watershed area (Figure 3). This interpretation is supported by
516 paleolimnological data (diatoms, sediment accumulation rates, and elemental and isotopic
517 compositions of organic matter) from across the Lake Ontario watershed that records an uptick in
518 nutrient loading from soil erosion runoff over the course of the mid-nineteenth century (Hodell

519 and Schelske 1998). The relatively early timeframe for the observed shift in fish nitrogen
520 isotopic compositions also suggests that its cause lay in new nutrient inputs originating primarily
521 within the local watershed area, rather than from other Great Lakes, because upstream
522 watersheds such as Lake Erie had not been as extensively settled by this time.

523 While inputs of ^{15}N -enriched terrestrial nutrients, released from eroding soils across the
524 watershed, could easily result in the observed isotopic pattern, additional related processes,
525 occurring both on land and in the water, could further contribute to the elevation of the stable
526 nitrogen isotopic composition of Lake Ontario's biota (Figure 3). Higher soil erosion rates would
527 increase mobilization of terrestrial nitrate across oxygen-poor ecotones (i.e., riparian/wetland
528 zones), promoting bacterial denitrification, leaving the residual nutrients that are eventually
529 delivered to Lake Ontario relatively enriched in ^{15}N (Bowden 1987; Knowles 1982; McClain, et
530 al. 2003). Increases in anthropogenic soil disturbance from farming and use of fertilizers
531 (pushing the nitrogen cycle to be more open; Szpak 2014), as well as changes to pH in aquatic
532 environments, may promote volatilization of ammonia before or during nutrient transport to
533 Lake Ontario's food web (Kendall 1998). Increased phosphorous loading from soil runoff would
534 also increase primary productivity, and the resulting bottom-up shift would have increased
535 demand on Lake Ontario's pool of available reactive N, thereby reducing discrimination against
536 ^{15}N during assimilation of DIN by phytoplankton (Hodell and Schelske 1998). Regardless of
537 which of these processes were most facilitative of the nitrogen isotopic patterning in Lake
538 Ontario's fish community, the timing (between 1832 and 1857) and abruptness of the ecosystem-
539 wide shift indicates that the first human activity to have broad-scale impacts on nitrogenous
540 nutrient pathways in the Great Lakes was likely deforestation.

541 It is also worth noting that the time period covered by this study includes significant global
542 climatic fluctuations, including the Medieval Climate Anomaly (MCA; 950–1250 CE) and the
543 Little Ice Age (LIA; 1300–1850 CE). Paleolimnological studies of Lake Ontario (McFadden, et
544 al. 2005), Lake Erie (Finkelstein and Davis 2006), and other lakes in the local (Keizer, et al.
545 2015; Mullins, et al. 2011; Paquette and Gajewski 2013; Stager, et al. 2017) and broader region
546 (Laird, et al. 2012) show that climatic variability associated with both the MCA and LIA did
547 have some impact on aquatic and terrestrial environments across the Great Lakes and St.
548 Lawrence River region. For instance, evidence from flux in biogenic silica sedimentation rates
549 and changes in diatom (e.g., *Fragelaria* spp.) communities (Finkelstein, et al. 2005; McFadden,
550 et al. 2005) demonstrates that these periods of climatic change were felt in the Lake Ontario
551 watershed over the past 1000 years. Climatic variability could affect Lake Ontario’s nitrogen
552 cycle and the nitrogen isotopic composition of fish through changes in adjacent wetland size
553 (important sites for denitrification; Bowden 1987; McClain, et al. 2003), terrestrial runoff
554 impacts (Anderson and Cabana 2005; Heaton 1986), fluctuating dissolved oxygen levels
555 (controlling nitrification and denitrification rates; Knowles 1982), and changes to productivity
556 and thermally regulated stratification (seasonal draw-down of DIN; Hodell and Schelske 1998).
557 Assuming that the influence of these potential sources of environmental variation did not cancel
558 one another out, our data suggest that late Holocene climatic variation did not influence Lake
559 Ontario’s nitrogen cycle or trophic structure.

560 The temporal as well as taxonomical detail offered by isotopic data from archaeological fish
561 specimens has provided a framework for higher resolution interpretation of food web dynamics
562 as well as the timing and likely cause of the first human impacts on Lake Ontario’s nitrogen
563 cycle. While previous isotopic analyses of organic matter from sediments (SI 1 and Figure 2;

564 Hodell and Schelske 1998; McFadden, et al. 2004) have demonstrated that Lake Ontario's
565 nitrogenous nutrients were becoming ^{15}N enriched during the nineteenth century, temporal
566 differences between datasets as well as a dearth of pre-1850 data prevented detailed analyses of
567 the processes which drove this change in lake-wide nitrogen cycling (SI 1). In addition, although
568 a variety of paleoenvironmental indicators (e.g., biogenic silica; Schelske 1991; Schelske and
569 Hodell 1991b; Schelske, et al. 1988) have shown human activities, such as deforestation, had
570 begun to have major nutrient (particularly P) loading impacts on the Lake Ontario watershed
571 from the mid-nineteenth century onward (although see, McAndrews and Boyko 1972; Schelske,
572 et al. 1983), our results demonstrate that Lake Ontario's nitrogen cycle had already been
573 substantially impacted decades earlier. Finally, by analyzing material from known taxa, we have
574 been able to document the impact that these changes in lake-wide nutrient dynamics had on long-
575 term food web structure.

576 *Anthropological and Ecological Implications:* This study is the first detailed analysis of long-
577 term isotopic variation in freshwater fish. Results suggest that, by altering aquatic nutrient inputs,
578 nineteenth-century European forestry practices in the lower Great Lakes region, whether focused
579 on timber extraction or producing cleared land for agriculture, left a durable isotopic signature on
580 fish in Lake Ontario. While the findings of this research focus on the Great Lakes watershed, the
581 implications are global; human impacts on aquatic nutrient dynamics, particularly the nitrogen
582 cycle, may be detectable in archaeological fish remains wherever land management has
583 extensively modified the balance of nutrient exchange between local terrestrial and aquatic
584 landscapes. Our results are particularly germane to recent discussion in conservation, calling for
585 increased attention to how human impacts on land may be a driver of environmental
586 deterioration in adjacent aquatic ecosystems (Abell 2002). For this reason, our findings have

587 significant potential to serve as a model for research that seeks to understand fundamental
588 changes in the way past societies, through forestry and other land management practices,
589 affected aquatic nutrient dynamics around the globe (Jenny, et al. 2019).

590 With respect to debate on the origins of the Anthropocene, our results provide an important
591 waypoint for the earliest lasting anthropogenic alteration of one of the world’s great freshwater
592 ecosystems. By identifying the tipping point at which human activities began to reorganize
593 nutrient dynamics in the Great Lakes region, our results do not reveal a “golden spike” indicator
594 for the starting point of the Anthropocene (Lewis and Maslin 2015), but, rather, provide a new
595 signpost along the path of the global expansion of industrial impacts. These changes to Lake
596 Ontario nonetheless represent a profound shift in the way humans have impacted the Great Lakes
597 environment and illustrate some important anthropological and ecological considerations for how
598 we can approach aquatic human–environment relationships in archaeology and historical ecology
599 moving forward.

600 From an ecological perspective, an ecosystem-wide, bottom-up alteration of the nitrogen cycle of
601 one of the world’s largest freshwater environments over the course of only one or two decades,
602 as shown here for Lake Ontario, demonstrates an unprecedented shift in the nature and scale of
603 how humans cause environmental change. Understanding how humans have shaped earth’s
604 biological and physical systems remains an important archaeological and ecological challenge
605 because finding lasting evidence of human impacts within complex regional or global systems is
606 rare. This is particularly true for our understanding of ancient environmental nutrient dynamics,
607 such as the cycling of nitrogen throughout an ecosystem, which, although fundamental to the
608 success of human societies both today and in the past, preserves relatively few traces of change.
609 Not only do our findings demonstrate how isotopic analyses of archaeological fauna can provide

610 a clearer overview of ancient nutrient dynamics, they do so in one of the most enigmatic arenas
611 of human–environment interaction, namely, aquatic ecosystems. Even today, in comparison with
612 terrestrial environments, surprisingly little is known about how escalating human exploitation of
613 the world’s aquatic environments will impact these vital ecosystems in the long term (Abell
614 2002). In that context, deeper retrospectives that are based on integrating results of contemporary
615 studies with data from both historical and archaeological resources, as demonstrated by this
616 study, have the potential to significantly enhance our understanding of the sensitivity of different
617 environments to human disturbances (Barak, et al. 2016; Dubois, et al. 2018; Jeffers, et al. 2015;
618 McLauchlan, et al. 2013).

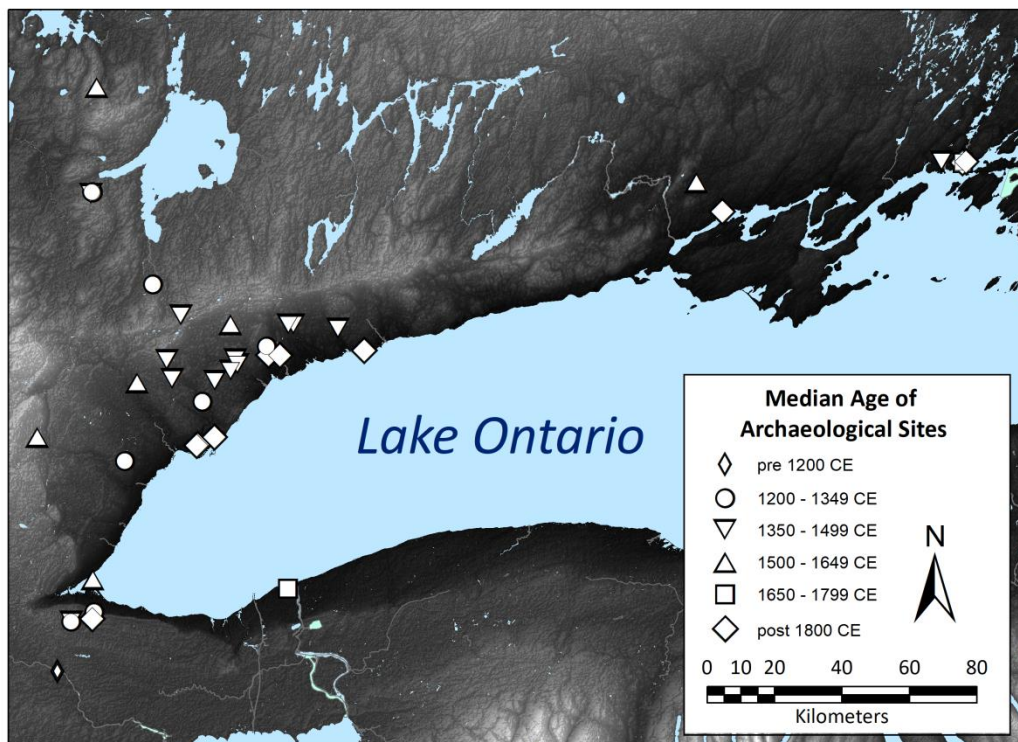
619 From an anthropological perspective, approaching questions of human–environment interactions
620 from the vantage point of nutrient dynamics could provide a novel outlook on the role of humans
621 as drivers of ecological change. While the relationship between humans and their environment
622 has traditionally been framed against a backdrop of such factors as technological change and the
623 development of social complexity as sources of evidence for macro-scale exploitation of
624 environmental resources (Kintigh, et al. 2014), there is an increasing recognition that an
625 understanding of how humans have influenced their landscape at a molecular level can provide
626 powerful new lines of evidence for previously invisible but profound shifts in human–
627 environment relations (Guiry, et al. 2018). In particular, throughout most times and places in
628 human history, the carrying capacity of local terrestrial and aquatic ecosystems, which are the
629 environmental foundations upon which societies are sustained, has hinged on the maintenance of
630 well-balanced biogeochemical cycles and nutrient regimes. With respect to lake and other
631 freshwater ecosystems, which have been and continue to be of pivotal importance to many
632 cultures, it is possible to use isotopic analyses of local fauna to look back in time and pinpoint

633 where, when, and in that context perhaps even how, human activities reached a turning point to
634 become dominant drivers in their ecosystem.

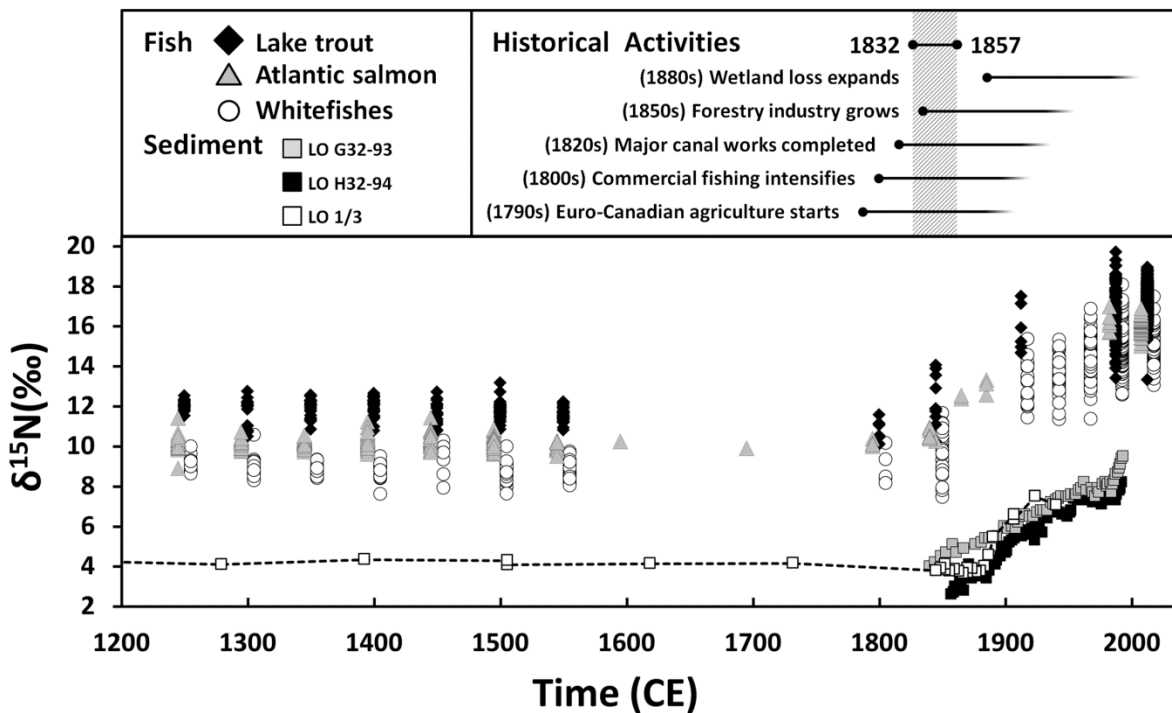
635

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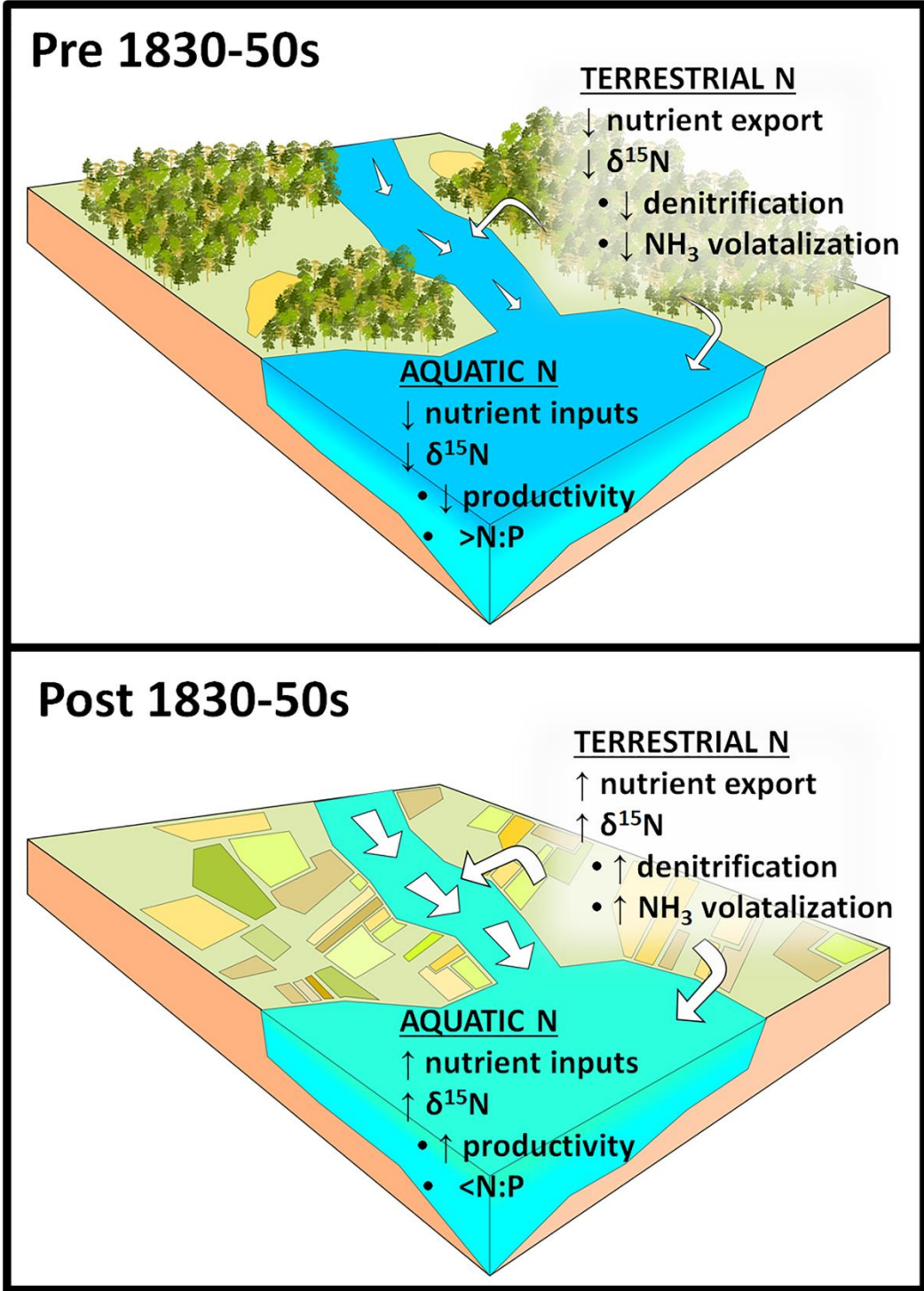


662
663 **Figure 1. Map of study area** showing locations of archaeological sites from which fish bone
664 samples were collected. Map data from NASA and ESRI.
665



666
 667 **Figure 2. Stable nitrogen isotope values for fish ($n=1469$) versus time.** Data are binned by 50-
 668 year (archaeological) and 25-year (historical/modern) intervals. For fish data sourced from the
 669 literature (Colborne, et al. 2016; Fera, et al. 2017; Guiry, et al. 2016a; Kiriluk, et al. 1995; Lumb
 670 and Johnson 2012; Mumby, et al. 2018; Rush, et al. 2012; van der Merwe, et al. 2003; Yuille, et
 671 al. 2015). For sediment data sourced from the literature: LO G32-93, LO H32-94 (Hodell and
 672 Schelske 1998), and LO 1/3 (McFadden, et al. 2004).

673



674

675 **Figure 3. Schematic diagram of nutrient regime shifts** (nitrogen sources and isotopic
 676 compositions) that could be associated with the intensification of deforestation and agriculture in
 677 the Lake Ontario watershed during the early to mid-nineteenth century.

678 Supplementary Information

679 **SI1** Paleoenvironmental context

680 **SI2** Isotopic results

681 **SI3** Calibration and analytical uncertainty

682 **SI4** Collagen peptide finger-printing results

683 **SI5** Statistical results

684

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