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1 **Advantages of zinc isotopes as a new dietary proxy in marine ecology**

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13

14 **Abstract**

15 In marine ecology, dietary interpretations of faunal assemblages often rely on nitrogen isotopes as the
16 main or only applicable trophic level tracer. We investigate geographic variability and trophic level isotopic
17 discrimination factors of a new tracer, bone ⁶⁶Zn/⁶⁴Zn ratios ($\delta^{66}\text{Zn}$ value), and compared it to collagen
18 nitrogen and carbon stable isotope ($\delta^{15}\text{N}$ and $\delta^{13}\text{C}$) values. Focusing on ringed seals (*Pusa hispida*) and
19 polar bears (*Ursus maritimus*) from multiple Arctic archaeological sites, we investigate trophic interactions
20 between predator and prey over a broad geographic area. All proxies show variability among sites,
21 influenced by the regional food web baselines. However, $\delta^{66}\text{Zn}$ shows a significantly higher homogeneity
22 among different sites. We observe a clear trophic spacing for $\delta^{15}\text{N}$ and $\delta^{66}\text{Zn}$ values in all locations, yet
23 $\delta^{66}\text{Zn}$ may more reliably record trophic levels between *U. maritimus* and prey species than $\delta^{15}\text{N}$. $\delta^{66}\text{Zn}$
24 analysis allows a more direct dietary comparability between spatially and temporally distinct locations
25 than what is possible by $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ analysis alone. When combining all three proxies a more detailed
26 and refined dietary analysis is possible.

27 **Keywords**

28 Zinc isotopes, diet, trophic ecology, nitrogen isotopes, Arctic, polar bear, ringed seal

29

30

31 Main

32 In ecology, archaeology and palaeontology, accurately reconstructing trophic levels can be challenging.
33 Among others, these reconstructions are required for effective management and conservation strategies¹,
34 understanding changing predator-prey and foraging ecology related to climate change² and for the
35 comparison of modern to fossil faunal assemblages^{3,4}. Stable isotope analysis is an effective tool for
36 analysing marine food webs, complementary to and often more reliable than non-stable isotope
37 approaches⁵. Bone collagen and soft tissue $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values are the traditional geochemical proxies
38 used for dietary and trophic level reconstructions^{6,7,8}. Only recently, studies of the bone's mineral phase
39 (bioapatite) non-traditional isotope systems such as calcium, magnesium and zinc (Zn) have shown
40 potential as palaeodietary proxies in the terrestrial^{9,10,11} and marine realm^{12,13}. Element and isotope ratios
41 observed in a diagenetically more resistant mineral phase (e.g., enamel) can preserve dietary information
42 beyond the scope of application of collagen, as recently demonstrated for Zn¹⁴. In addition, even when
43 collagen is well preserved, combining traditional collagen $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ analyses with Zn isotope analyses
44 of bioapatite may provide complementary dietary information as proven by the lack of correlation of
45 those tracers within individuals of the same species, implying independent controlling mechanisms¹⁵.

46 Nitrogen-15 becomes relatively enriched in the tissues of aquatic consumers with successive trophic
47 level¹⁶. The $\delta^{13}\text{C}$ values behave much more conservatively with trophic level, increasing typically by less
48 than 1 ‰ with trophic position for most tissues, compared to on average 3.4 ‰ higher $\delta^{15}\text{N}$ values
49 between a predator and its prey^{17,18}. Carbon isotopes are therefore more commonly used to infer the
50 source(s) of primary production at the base of the food web. For Zn, studies have shown a distinct ⁶⁶Zn
51 depletion in carnivore bioapatite relative to that of herbivores^{11,14}. As muscles and most organs are
52 typically ⁶⁶Zn depleted relative to the animal's diet and its bulk body $\delta^{66}\text{Zn}$ composition^{19,20,21}, bones of
53 carnivores (and their bulk body composition) have lower $\delta^{66}\text{Zn}$ values than their prey's. Diet thus exerts
54 control on the $\delta^{66}\text{Zn}$ values of soft tissue and bioapatite. Most non-diet related factors, such as sex and
55 age of an animal, have so far shown no effect on the isotope values and relative isotopic variability in body
56 tissues^{21,22}. Provenance, on the other hand, appears to affect the Zn isotopic compositions of terrestrial
57 vertebrates, though it remains unclear to which extent^{11,14,23}. Provenance is known to play an important
58 role for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ studies in marine food webs²⁴ but has until now not been studied for $\delta^{66}\text{Zn}$.

59 In the marine realm, the $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values of particulate organic matter (POM), consisting of
60 phytoplankton, bacteria, microzooplankton and detritus, show a substantial spatial variation within and
61 among ocean basins^{5,25,26}. Variation in the isotopic composition at the base of the marine food web is
62 passed along to higher trophic levels, limiting the comparability of populations from areas with varying
63 isotopic food web baselines, or migratory animals. A high variability in modern baseline $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$
64 values is documented by the isotopic composition of POM, zooplankton, filter feeders, as well as higher
65 trophic level consumers across the Arctic^{6,27,28,29,30,31} (Supplementary Text). Noteworthy is a west-east ¹³C
66 depletion observed in consumers from the Bering Sea (Bering Strait) through the Chukchi Sea to the
67 Beaufort Sea^{6,27,28,31} and significant spatial variability in zooplankton $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values between the
68 Canadian Arctic Archipelago (CAA), Labrador Sea, and Baffin Bay^{31,32}. Particularly for bone collagen, with
69 its long turnover time, trophic level reconstructions can be compromised when animals frequently

70 migrate between areas of differing baseline isotopic composition, or when comparing animals from
71 spatially and temporally distinct locations.

72 Data on food web baseline $\delta^{66}\text{Zn}$ values and variability is non-existent for both the continental and marine
73 realms. Because of biological uptake, dissolved Zn concentrations are highly depleted in marine surface
74 waters, often much less than 1 nmol kg^{-1} ^{33,34}. The isotopic composition of dissolved Zn below 500 m seems
75 to be globally homogenous with values close to $+0.5 \text{ ‰}$, despite variable Zn concentrations^{35,36}. Bulk
76 marine Zn is enriched in ^{66}Zn relative to its major inputs from rivers and aeolian dust, which centre on the
77 global crustal average of $+0.3 \text{ ‰}$ ³⁷. Although most studies on cultured phytoplankton demonstrate a
78 preferential uptake of light Zn into the cell relative to the bulk growth medium^{38,39}, Atlantic and Pacific
79 vertical Zn isotope profiles generally show lower $\delta^{66}\text{Zn}$ values in surficial waters compared to that of the
80 deep water^{33,36,40,41,42}. These studies demonstrate that the isotopic composition of Zn is most variable
81 within the surface water ($< 500 \text{ m}$), often with higher values in the uppermost surface ($< 20 \text{ m}$).

82 Here, we aim at exploring Zn isotopes as a reliable tracer of marine trophic levels. To do so, we investigate
83 combined bone $\delta^{66}\text{Zn}$, $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ from the same species across 13 locations (17 sites) in a large
84 geographic area stretching across the Arctic from the Hudson Strait in the east to the Bering Strait in the
85 west. We include 5 locations and two single samples with already published $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values^{4,13,43,44},
86 as well as one site with already published $\delta^{66}\text{Zn}$ values¹³. For this study, we analysed 167 archaeological
87 bones, concentrating on polar bears (*Ursus maritimus*) and ringed seals *Pusa hispida* (*Phoca hispida*).
88 Focusing on these species allows us to investigate $\delta^{66}\text{Zn}$ trophic level isotopic discrimination factors
89 between predator and prey geographically. Both species have a circumpolar Arctic distribution and are
90 abundant throughout the Arctic today^{45,46}. Particularly, *P. hispida* remains are frequently found in
91 archaeological assemblages with a large temporal and spatial range^{43,47,48}. Studying the isotopic
92 composition of high trophic level predators such as *U. maritimus* and *P. hispida* has the advantage of their
93 tissues' composition dampening the effects of short-term environmental variation. This effectively leads
94 to less isotopic variability and "noise" in the animal's tissues compared to those of lower trophic levels⁴³.
95 Therefore, these two species are prime targets to investigate geographical variability of dietary proxies
96 and trophic level isotopic discrimination factors.

97 Results

98 Bone collagen $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and bone $\delta^{66}\text{Zn}$ values of *P. hispida*, *U. maritimus*, harp seal (*Pagophilus*
99 *groenlandicus*) and beluga whale (*Delphinapterus leucas*) are reported in table 1 and Supplementary Table
100 1. All collagen samples had yields and elemental (wt% C, wt% N, C:N_{atomic}) compositions characteristic of
101 samples with isotopic compositions not altered by contaminant or degradation in the burial
102 environment^{49,50} (Supplementary Table 1). Likewise, $\delta^{66}\text{Zn}$ values do not indicate a modification due to
103 diagenesis or contamination for the majority of samples, but we cannot exclude it as a possibility for
104 outlier values (Supplementary Discussion). Our results also indicate that the presence of collagen-bound
105 Zn and thereby collagen preservation has no effect on the mineral phase $\delta^{66}\text{Zn}$ values (Supplementary Fig.
106 1-2, Supplementary Table 2, Supplementary Results, Supplementary Discussion).

107 Statistically significant differences between *P. hispida* populations was determined through ANOVA for
108 $\delta^{13}\text{C}$ ($F(12, 91) = 24.4$, $p\text{-value} < 0.05$), $\delta^{15}\text{N}$ ($F(12, 91) = 34.74$, $p\text{-value} < 0.05$) and $\delta^{66}\text{Zn}$ values ($F(12, 91)$
109 $= 5.867$, $p\text{-value} < 0.05$). Post-hoc Tukey pair-wise comparisons draw out the populations from Little
110 Cornwallis (QjJx-1) and the North shore of Devon Island (QkHn-13) both part of the CAA, as well as eastern

111 Ellesmere Island (near Skraeling Island, Sffk-4), linked to the North Water Polynya, as distinct from some
112 of the other sites in regards to their $\delta^{66}\text{Zn}$ values (Extended Data Fig. 1). Every other site, regardless of
113 their broad geographic group, are not significantly different from one another.

114 Results for pair-wise comparisons of sites' $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ *P. hispida* values show a higher degree of
115 heterogeneity (Extended Data Fig. 2-3). However, most of the differences can be linked to geographic
116 groups. Sites from the CAA are being drawn out as different in their $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values to most of the
117 other sites. The western sites of the Amundsen and Coronation Gulf, as well as the Bering Strait, differ in
118 their $\delta^{15}\text{N}$ values, but not for $\delta^{13}\text{C}$ values. Finally, $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values from the Eastern sites of the Hudson
119 Bay and the Labrador Sea are identified as significantly different than those of western sites.

120 Levene's tests for equal variance show that $\delta^{66}\text{Zn}$ values are more homogeneous between *P. hispida* and
121 *U. maritimus* ($F(1, 125) = 3.43$, $p = 0.27$) and across sites ($F(8, 118) = 1.72$, $p = 0.40$) than $\delta^{15}\text{N}$ values
122 (respectively $F(1, 125) = 6.95$, $p = 0.04$; and $F(1,118) = 2.62$, $p = 0.05$).

123

124 Discussion

125 Site-specific isotopic variability

126 The maximum variability for inter-site mean *P. hispida* bone $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values (3.55 and 3.40 ‰)
127 exceeds the maximum intra-site (1.77 and 2.67 ‰, Fig. 1) and typical trophic level variability. The QjJx-1
128 site on Little Cornwallis Island is a notable exception with very high on-site *P. hispida* bone collagen $\delta^{15}\text{N}$
129 variability (3.85 ‰)¹³. Post-hoc Tukey pair-wise comparisons demonstrate a large heterogeneity in $\delta^{15}\text{N}$
130 and $\delta^{13}\text{C}$ values between archaeological populations (Extended Data Fig. 2-3). Isotopic heterogeneity
131 between populations is related to geographic location resulting in $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values from populations
132 of different regions plotting in distinct groups on a $\delta^{15}\text{N}$ versus $\delta^{13}\text{C}$ plot (Fig. 1b). Based on site proximity
133 and sample isotopic composition, we grouped sites from the Bering/Chukchi Sea, Amundsen and
134 Coronation Gulf, CAA, North Water Polynya, Hudson Bay, and sites influenced by the Labrador Sea
135 (Hudson Strait and Frobisher Bay, Fig. 1a). The $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ variability between the archaeological sites
136 is in good agreement in both spacing and amplitude with modern geographical variations observed from
137 zooplankton^{26,27,28} and higher consumer soft tissue^{6,27,28,31} including *P. hispida*^{51,52,53,54,55,56} and *U.*
138 *maritimus*⁴⁶. While dietary differences between populations may have contributed to the geographic
139 spacing of $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values, an integration of regional baseline isotopic patterns is the main factor
140 controlling the observed inter-site isotopic variability in *P. hispida* and *U. maritimus* collagen
141 (Supplementary Discussion).

142 In contrast to *P. hispida* bone $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values, the highest variability for mean $\delta^{66}\text{Zn}$ values between
143 sites (0.23 ‰) does not exceed the maximum variability observed within a single site (0.36 ‰).
144 Additionally, mean $\delta^{66}\text{Zn}$ values between sites never exceeds mean trophic position variability between
145 *U. maritimus* and *P. hispida* (~0.32 ‰). While ANOVA analysis of *P. hispida* $\delta^{66}\text{Zn}$ values did reveal
146 statistically significant differences between *P. hispida* populations, post-hoc pair-wise comparisons tests
147 show considerably more homogeneity in $\delta^{66}\text{Zn}$ values than for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values (Extended Data Fig. 1-
148 3). Similarly, Levene's tests for equal variance shows that for all sites, *P. hispida* and *U. maritimus* $\delta^{66}\text{Zn}$
149 values have an equal variance, whereas $\delta^{15}\text{N}$ values are more heterogeneous. The low geographical $\delta^{66}\text{Zn}$

150 variability in *P. hispida* and *U. maritimus* bones implies that Arctic food-web baseline and/or low trophic
151 level consumer $\delta^{66}\text{Zn}$ values are more homogenous than for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values. This is remarkable
152 considering the large surface water's isotopic variability observed for dissolved Zn across the Atlantic and
153 Pacific of -1.1 to $+0.9$ ‰ and -0.9 to $+0.2$ ‰, respectively^{33,42}.

154 Based on post-hoc Tukey pair-wise comparison, $\delta^{66}\text{Zn}$ values from *P. hispida* populations from the sites
155 QjJx-1 (Little Cornwallis Island), QkHn-13 (Devon Island) and Sffk-4 (eastern Ellesmere Island) were
156 identified as statistically different from other populations (Fig. 2, Extended Data Fig. 1). QjJx-1 and QkHn-
157 13 are located within the CAA. The CAA is composed of multiple channels and interconnected basins, in
158 which water mass modification and transport are governed by its complex topography and shelf exchange
159 processes⁵⁷. Within this setting, baseline $\delta^{66}\text{Zn}$ values may be more variable on a regional scale than for
160 the rest of the Arctic. For the Sffk-4 site, we observe in all three species analysed lower mean $\delta^{66}\text{Zn}$ values
161 compared to other sites indicating a regionally lower baseline $\delta^{66}\text{Zn}$ value (Fig. 2, Fig. 3). The Sffk-4 site is
162 located at the biologically highly productive⁵⁸ northern edge of the North Water Polynya, a region in which
163 the reduced ice-cover or ice-free conditions influence biological processes (e.g., by upwelling, increased
164 nutrient renewal)⁵⁹, which in turn may modify the $\delta^{66}\text{Zn}$ baseline.

165

166 **Species-specific isotopic variability**

167 Both $\delta^{66}\text{Zn}$ and $\delta^{15}\text{N}$ values are controlled by diet but show a better correlation for *U. maritimus* samples
168 across all sites than for *P. hispida* (Fig. 1 c, e), perhaps related to the more specialised diet of *U.*
169 *maritimus*^{60,61}. As with $\delta^{15}\text{N}$ values, bone $\delta^{66}\text{Zn}$ values clearly demonstrate a trophic spacing between *U.*
170 *maritimus* and *P. hispida* in all locations analysed (Fig. 2). The KkJg-1 site in Hudson Bay is an exception
171 with two *U. maritimus* samples showing anomalously high $\delta^{66}\text{Zn}$ values which may relate to non-dietary
172 factors such as contamination, misidentification, diagenesis or physiological effects (Supplementary
173 Discussion). Even when including the KkJg-1 site, Levene's tests for equal variance demonstrate an equal
174 variance between *P. hispida* and *U. maritimus* $\delta^{66}\text{Zn}$ values ($F(1, 125) = 3.43$, $p = 0.27$), whereas their $\delta^{15}\text{N}$
175 values demonstrate heterogeneity ($F(1, 125) = 6.95$, $p = 0.04$). Because $\delta^{66}\text{Zn}$ is more homogenous in its
176 value for a specific taxon (and diet?), $\delta^{66}\text{Zn}$ may more reliably reflect trophic positions than bulk $\delta^{15}\text{N}$
177 values, when investigating multiple species across multiple sites, proving a better inter-site comparability.
178 *U. maritimus* bones are on average 0.32 ‰ lower than their primary prey species *P. hispida* (mean $\Delta^{66}\text{Zn}_{U.$
179 *maritimus} - P. hispida} = -0.32 ‰). Previously, estimations of trophic discrimination factors between enamel and
180 bones of terrestrial mammalian carnivores and herbivores were between -0.6 and -0.4 ‰ for the Tham
181 Hay Marklot (THM) cave¹⁴ and the modern Koobi Fora region¹¹. These studies, however, had a lower
182 sample size and compared multiple carnivores and herbivores with varying diets. Predicted bone $\delta^{66}\text{Zn}$
183 trophic level discrimination factors are between -0.36 and -0.38 ‰ when calculated using individual $\delta^{15}\text{N}$
184 trophic levels⁸ from all marine mammal taxa with available $\delta^{66}\text{Zn}$ data (Supplementary Discussion,
185 Supplementary Equation 1-3). These values are close to the mean $\Delta^{66}\text{Zn}_{U. maritimus} - P. hispida}$ value.*

186 Particularly for archaeological material, assigning a trophic level to multiple species when utilising $\delta^{15}\text{N}$
187 values alone can be challenging, as shown by the large differences in mean $\delta^{15}\text{N}$ trophic discrimination
188 factors between *U. maritimus* and *P. hispida* ($\Delta^{15}\text{N}_{U. maritimus} - P. hispida}$) for individual sites ($+2.2$ to $+7.0$ ‰,
189 Fig. 2 b). Besides diet, $\Delta^{15}\text{N}_{U. maritimus} - P. hispida}$ variability may be influenced by physiological effects or

190 unknown archaeological assemblage effects related to human hunting and/or scavenging (Supplementary
191 Discussion). For $\delta^{66}\text{Zn}$, two sites from the same geographic area close to the Labrador Sea (KkDo-1 and
192 JfEI-4), have a markedly lower trophic spacing between *P. hispida* and *U. maritimus* $\delta^{66}\text{Zn}$ values of -0.22
193 and -0.24 ‰ (Fig. 2). Modern *U. maritimus* individuals from the area belong to the Davis Strait
194 population⁶². In addition to *P. hispida* and contrary to most other *U. maritimus* populations, this one
195 obtains a large percentage of its biomass from the consumption of harp seals (*Pagophilus*
196 *groenlandicus*)^{60,61}. However, bone $\delta^{66}\text{Zn}$ of *P. hispida* and *P. groenlandicus* from the same site are
197 indistinguishable (Fig. 3). Instead of the consumption of *P. groenlandicus*, the lower trophic discrimination
198 factor for these sites may arise from unknown dietary contributions, population-specific physiological
199 effects or unknown archaeological assemblage effects.

200 Unlike *P. hispida*, most *P. groenlandicus* leave their Canadian Arctic summering grounds, ahead of the
201 formation of local pack ice in autumn^{63,64}. However, when sympatric with *P. hispida*, *P. groenlandicus*
202 feeds at a similar trophic level, consuming many of the same prey species, and both species show no
203 statistical difference in muscle and liver $\delta^{15}\text{N}$ values⁶⁵. Indeed, bones of both seal species cover the same
204 $\delta^{66}\text{Zn}$ range for the same location (Fig. 3). However, bones of *P. groenlandicus* from Hudson Strait (KkDo-
205 1) have almost 1 ‰ lower mean $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values than those of *P. hispida*, perhaps related to this
206 species' seasonal southwards migration (Fig. 3 b, d). In contrast, some *P. groenlandicus* individuals remain
207 in west Greenland waters during winter^{66,67}, which may explain why bones of both seal species from
208 eastern Ellesmere Island show a similar $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and $\delta^{66}\text{Zn}$ range (Fig. 3 a, c). Due to its long turnover
209 time, the bone collagen isotopic composition of *P. groenlandicus* likely represents an amalgamation of
210 different food sources and local isotopic baseline values along their migration route and within their
211 seasonal feeding grounds. *P. groenlandicus* bone $\delta^{66}\text{Zn}$ values do not seem to record migratory signals,
212 again arguing for lower baseline variability or Zn isotopic homogenisation within low trophic level
213 organisms. Despite a very low samples size ($n = 2$) beluga whale (*Delphinapterus leucas*) $\delta^{66}\text{Zn}$ values fall
214 within the same range as *P. hispida* and *P. groenlandicus* with slightly higher mean values (Fig. 3b, d).
215 Indeed, all three species occupy a similar trophic level^{8,65}. When sympatric with *P. hispida*, *D. leucas*
216 typically has slightly lower soft tissue $\delta^{15}\text{N}$ values likely due to migrating between areas with differing
217 baselines or a more offshore/pelagic foraging^{8,54}. Here, *D. leucas* $\delta^{15}\text{N}$ values are higher than those of *P.*
218 *hispida* and their $\delta^{13}\text{C}$ values are highly variable (Fig. 3). Instead of only reflecting this species' trophic
219 position relative to *P. hispida*, their collagen $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values are likely influenced by the high mobility
220 of this species⁶⁸ and its foraging in locations with different isotopic baselines.

221 The trophic levels of *U. maritimus*, *P. hispida* and *P. groenlandicus* are reflected by their bone collagen
222 $\delta^{15}\text{N}$ and bone $\delta^{66}\text{Zn}$ values across the Arctic. The analysis of Zn isotopes, however, offers additional
223 advantages for studying marine trophic ecology, not only due to its greater preservation potential in fossil
224 material¹⁴, but also due to the lower baseline controlled species-specific spatial isotopic gradients. The
225 inclusion of $\delta^{66}\text{Zn}$ analysis in ecological, archaeological and palaeontological studies may thus allow more
226 robust interpretations of spatial and temporal trophic interactions. Additionally, while both $\delta^{66}\text{Zn}$ and $\delta^{15}\text{N}$
227 generally record trophic levels, they do not record physiological and/or dietary effects equally, thus
228 providing a strong incentive to combine $\delta^{66}\text{Zn}$ with $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ analyses where possible.

229

230

231 Conclusion

232 This study presents the first comparison of archaeological $\delta^{66}\text{Zn}$ bone compared with traditional collagen
233 $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values for the same species across a large geographic area. Focussing on prey (*P. hispida*)
234 and predator (*U. maritimus*) we investigate the baseline variability and trophic spacing of these dietary
235 proxies. Our results show that:

- 236 1) Overall, $\delta^{66}\text{Zn}$ values shows less site-specific variability within a species, likely due to a lower
237 baseline variability than for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$. As such, $\delta^{66}\text{Zn}$ values are particularly valuable for
238 dietary studies on highly mobile species (or consumers thereof) and for comparing geographically
239 and temporally distinct populations.
- 240 2) We observe the expected trophic level spacing for collagen $\delta^{15}\text{N}$ and bone $\delta^{66}\text{Zn}$ values between
241 *U. maritimus* and *P. hispida*. *U. maritimus* bone $\delta^{66}\text{Zn}$ values are on average 0.32 ‰ lower than of
242 its primary prey *P. hispida*.
- 243 3) Bone $\delta^{66}\text{Zn}$ values of the migratory species *P. groenlandicus* and *D. leucas* are consistent with
244 respect to their known trophic positions when compared with *P. hispida* and *U. maritimus* values.
245 In contrast, their collagen $\delta^{15}\text{N}$ (and $\delta^{13}\text{C}$) values appear to document their relative trophic
246 positions less precisely, likely influenced by variations in baseline isotopic compositions along
247 their migration routes.

248 In ecological, archaeological and palaeontological research, trophic level estimations often rely exclusively
249 on the $\delta^{15}\text{N}$ tracer, sometimes biased by physiological, habitat and baseline effects. We demonstrate that
250 the inclusion of $\delta^{66}\text{Zn}$ analysis can provide otherwise inaccessible supplementary dietary information and
251 more robust trophic level estimations.

252

253 Material and Methods

254 For this study, we compare $\delta^{66}\text{Zn}$, $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values of 104 *P. hispida*, 47 *U. maritimus*, 11 *P.*
255 *groenlandicus* and 2 *D. leucas* archaeological bone samples from across the Arctic (Supplementary
256 Table 1). The data presented herein includes already-published $\delta^{66}\text{Zn}$ values from an archaeological site
257 (QjJx-1) on Little Cornwallis Island¹³. Additional $\delta^{66}\text{Zn}$ values analysed for this study comprise 93 *P. hispida*
258 bone samples from 13 archaeological sites (12 locations) and 37 *U. maritimus* bone samples from 11
259 archaeological sites (8 locations) as well as *P. groenlandicus* and *D. leucas* samples (2 sites, 1 site,
260 respectively). For 6 of the 17 sites analysed here (RbJu-1, PaJs-13, QkHn-13, QjJx-1, KTZ sites), $\delta^{15}\text{N}$ and
261 $\delta^{13}\text{C}$ values were already published elsewhere^{4,13,43}. Additionally, $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values from one *P. hispida*
262 sample from the NkRi-3 and OkRn-1 sites were already published elsewhere (Sample Nr. 4945, 9535)⁴⁴.
263 For the sites JfEl-4, KcFs-2, NkRi-3, seal bones analysed are identified as most likely *P. hispida*, but we
264 cannot completely rule out that some samples may also come from other Phocidea (Supplementary
265 Discussion). A single walrus (*Odobenus rosmarus*) bone and a potentially misidentified *D. leucas* bone from
266 the JfEl-4 site were also measured and are discussed compared to previously measured *O. rosmarus* bones
267 from the QjJx-1 site¹³ and our $\delta^{66}\text{Zn}$ data from other species and sites in the Supplementary Discussion.
268 Additional information and references regarding the archaeological context of the samples and sites are
269 provided in the Supplementary Text.

270 $\delta^{66}\text{Zn}$ analysis

271 All samples' surfaces were mechanically abraded (cleaned) to avoid sediment contamination, using a
272 dental drill equipped with a diamond-tipped burr. Approximately 10 to 50 mg chunks were then sampled
273 using a diamond-tipped cutting wheel. The chunks were then ultrasonicated in ultrapure water (Milli-Q
274 water) for 5 min and dried in a drying chamber for a few days at 50 °C. Bone samples and reference
275 materials NIST SRM 1400 and NIST SRM 1486 were subjected to different dissolution methods (HCl and
276 HNO_3) to investigate the impact of the organic bone phases on its Zn isotope signal (Supplementary
277 Methods, Supplementary Discussion). The column chromatography steps (3.1.2) for a quantitative
278 recovery of sample $\text{Zn}^{69,70}$ were the same for all samples regardless of the dissolution methods used. Each
279 column chromatography batch ($n = 15$) included up to 13 samples, one chemistry blank and at least one
280 reference standard (SRM 1400 and/or 1486).

281 Zn purification was performed in two steps, following the modified ion exchange method adapted from
282 Moynier et al.⁷⁰, first described in Jaouen et al.¹¹. Each step included AG-1x8 resin that was cleaned and
283 conditioned prior to sample loading. One ml of AG-1x8 resin (200–400 mesh) was placed in 10 ml
284 hydrophobic interaction columns (Macro-Prep® Methyl HIC). Resin cleaning involved 5 ml 3 % HNO_3
285 followed by 5 ml ultrapure water. These cleaning steps were repeated. The resin was then conditioned
286 with 3 ml 1.5M HBr. After loading, 2 ml HBr were added for matrix residue elution followed by Zn elution
287 with 5 ml HNO_3 . Following the second column step, the solution was evaporated for 13 h at 100 °C and
288 the residue re-dissolved in 1 ml 3 % HNO_3 .

289 Zn isotope ratios were measured using a Thermo Fisher Neptune MC-ICP-MS at the Max Planck Institute
290 for Evolutionary Anthropology (Leipzig, Germany) and a Thermo Fisher Neptune Plus MC-ICP-MS at the
291 Géosciences Environnement Toulouse - Observatoire Midi-Pyrénées (Toulouse, France). Instrumental
292 mass fractionation was corrected by Cu doping following the protocol of Maréchal et al.⁶⁹ and Toutain et
293 al.⁷¹. The in-house reference material Zn AA-MPI was used for standard bracketing. $\delta^{66}\text{Zn}$ values are
294 expressed relative to the JMC-Lyon reference material. Analysed sample solution Zn concentrations were
295 close to 300 ppb as was the Zn concentration used for the standard mixture solution. Zn concentrations
296 in the respective samples were estimated following a protocol adapted from one used for Sr by Copeland
297 et al.⁷², applying a regression equation based on the ^{64}Zn signal intensity (V) of three solutions with known
298 Zn concentrations (150, 300 and 600 ppb). $\delta^{66}\text{Zn}$ uncertainties were estimated from standard replicate
299 analyses and ranged between ± 0.01 ‰ and ± 0.03 ‰ (1SD). Additional reference materials SRM 1486
300 and SRM 1400 were analysed alongside the samples. SRM reference materials and samples show a normal
301 Zn mass dependent isotopic fractionation, i.e. the absence of isobaric interferences, as the $\delta^{66}\text{Zn}$ vs. $\delta^{67}\text{Zn}$
302 and $\delta^{66}\text{Zn}$ vs. $\delta^{68}\text{Zn}$ values fall onto lines with slopes close to the theoretic mass fractionation values of 1.5
303 and 2, respectively.

304

305 $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ analysis

306 Bone surfaces were cleaned with a dental drill equipped with a diamond-cutting wheel. Subsamples of
307 bone chunks (100-200 mg) were demineralised in 0.5 M HCl at 4°C. After demineralisation, samples were
308 rinsed to neutrality with Type I water (resistivity >18.2 M Ω -cm). Any bone samples with dark colouration

309 were treated with 0.1 M NaOH for successive 30 min treatments under sonication at room temperature
310 until the solution no longer changed colour. The samples rinsed to neutrality with Type I water and then
311 the insoluble collagen residue was solubilised in ~8 ml of 0.01 M HCl at 75°C for 48 h. The resulting solution
312 containing the solubilised collagen was filtered through a 5–8 µm filter and then filtered using a Microsep®
313 30 kDa molecular weight cut-off (MWCO) ultrafilter (Pall Corporation, Port Washington, NY) to remove
314 low molecular weight compounds⁷³. The >30 kDa fraction was freeze-dried, and the collagen yield was
315 calculated.

316 Carbon and nitrogen isotopic and elemental compositions were determined using an IsoPrime continuous
317 flow isotope-ratio mass spectrometer (CF-IRMS) coupled to a Vario Micro elemental analyser (Elementar,
318 Hanau, Germany). Carbon and nitrogen isotopic compositions were calibrated relative to the VPDB and
319 AIR scales, respectively, using a two-point calibration anchored by USGS40 (accepted $\delta^{13}\text{C}$ -26.39 ± 0.04
320 ‰, $\delta^{15}\text{N}$ -4.52 ± 0.06 ‰) and USGS41 (accepted $\delta^{13}\text{C}$ $+37.63\pm 0.05$ ‰, $\delta^{15}\text{N}$ $+47.57\pm 0.11$ ‰)⁷⁴. Standard
321 uncertainty was determined to be ± 0.20 ‰ for $\delta^{13}\text{C}$ and ± 0.25 ‰ for $\delta^{15}\text{N}$ ⁷⁵. Additional details are provided
322 in the Supplementary Methods.

323

324 **Statistical analyses**

325 Analysis of variance (ANOVA) were performed across the dataset in order to determine statistical
326 differences in $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and $\delta^{66}\text{Zn}$ values between *P. hispida* populations. A single *P. hispida* specimen
327 from Little Cornwallis Island was excluded from the statistical analysis ($\delta^{66}\text{Zn} = 1.00$ ‰, from Jaouen et
328 al.¹³), as it could disproportionately influence the analysis (see Extended Data Fig. 1-3 versus 4-6). It was
329 singled-out as an extreme outlier lying more than 3 times the interquartile range above the third quartile,
330 both within-site and for the whole *P. hispida* dataset. Where variance was found to be significant, post-
331 hoc Tukey pair-wise comparisons were carried out to determine which populations were significantly
332 different from each other in terms of their $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and $\delta^{66}\text{Zn}$ values. To adhere to ANOVA's assumptions,
333 each *P. hispida* populations' $\delta^{13}\text{C}$, $\delta^{15}\text{N}$ and $\delta^{66}\text{Zn}$ datasets underwent visual inspection to check for
334 normally distributed and homogeneous residuals, as well as tested for equal variance using Levene's test.
335 Accordingly, we report the results of ANOVAs and post-hoc Tukey pair-wise comparisons (Extended Data
336 Fig. 1-3). In order to investigate the homogeneity of $\delta^{66}\text{Zn}$ values within the Arctic relative to $\delta^{15}\text{N}$ values,
337 a series of Levene's test for equal variance (with Bonferroni correction) was performed on Zn and N
338 isotopic values between *P. hispida* and *U. maritimus*, as well as between sites for which data is available
339 for both species. All statistical analyses were conducted using the free program R software⁷⁶.

340

341 **References**

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- 517 76. R Core Team, *R version 3.6.1* (R Foundation for Statistical Computing, Vienna, Austria, 2018).

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520

521

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528

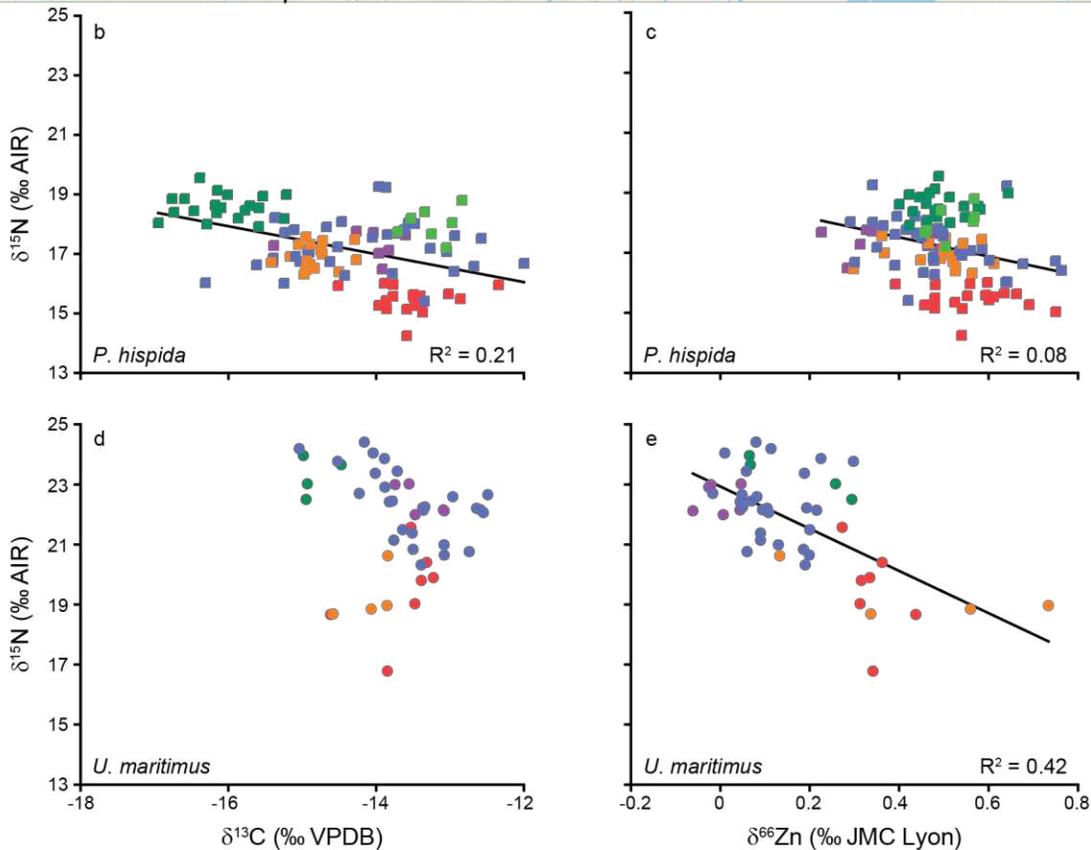
529 **Author contributions**

530 K.J. and J.M. designed the study. P.S., K.J. and J.M. selected the sample material. P.S., M.R. and C.H.
531 performed the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ analyses. J.M. and N.B. performed the $\delta^{66}\text{Zn}$ analyses at the MPI EVA Leipzig.
532 J.M., K.J. and P.M. performed the $\delta^{66}\text{Zn}$ analyses at the CNRS in Toulouse. N.B. performed the statistical
533 analysis. J.M., K.J., N.B., P.S., M.R. and J.-J.H., analysed and interpreted the data. J.M. wrote the initial
534 manuscript. All authors contributed to editing the final version of the manuscript.

535

536 **Competing interests**

537 The authors declare no competing interests



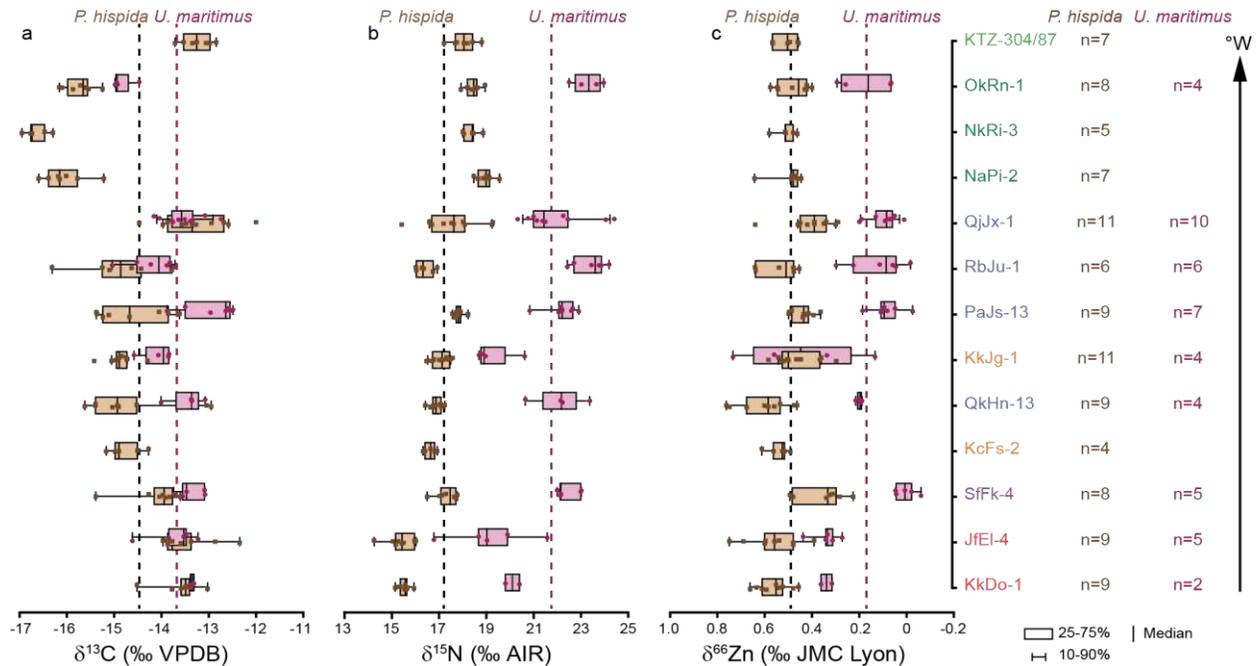
538

539 **Figure 1:** Isotopic composition of *P. hispida* (squares, n = 104) and *U. maritimus* (dots, n = 47) samples
 540 from Arctic archaeological sites colour coded as geographic groups. a) Map indicating the archaeological
 541 sites analysed and geographic colour coding: Light green for the Bering Strait; dark green for the
 542 Amundsen and Coronation Gulf; blue for the CAA; orange for the Hudson Bay; purple for the North Water
 543 Polynya; and red for sites influenced by the Labrador Sea in the Hudson Strait and Frobisher Bay. b) $\delta^{15}\text{N}$

544 versus $\delta^{13}\text{C}$ plot for *P. hispidus* samples (p -value < 0.05; $R^2 = 0.21$; $n = 104$). c) $\delta^{15}\text{N}$ versus $\delta^{66}\text{Zn}$ plot for *P.*
 545 *hispidus* samples (p -value < 0.05; $R^2 = 0.08$; $n = 104$). d) $\delta^{15}\text{N}$ versus $\delta^{13}\text{C}$ plot for *U. maritimus* samples (no
 546 correlation, p -value > 0.05; $n = 47$). e) $\delta^{15}\text{N}$ versus $\delta^{66}\text{Zn}$ plot for *U. maritimus* samples (p -value < 0.05; R^2
 547 = 0.42; $n = 47$). We included already published $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values^{4,13,43,44} and already published $\delta^{66}\text{Zn}$
 548 values from QjJx-1¹³. The map is redrawn after www.google.com/maps.

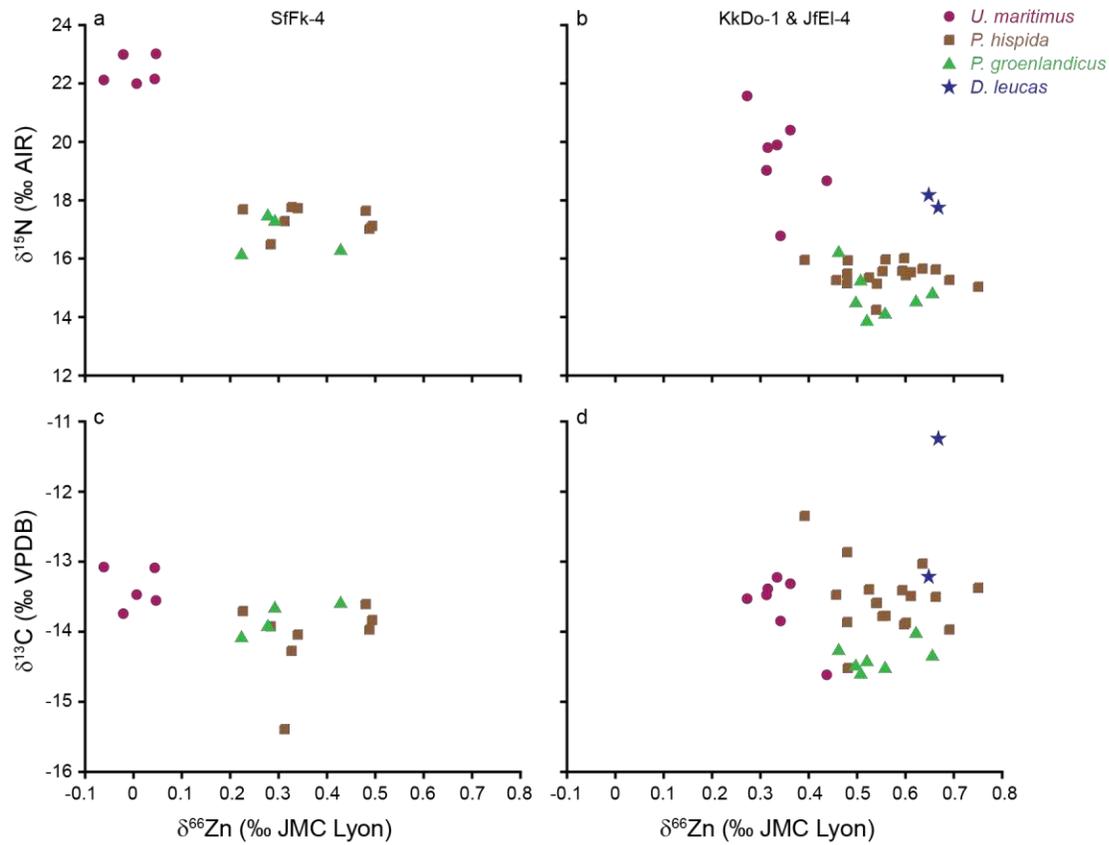
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552 **Figure 2:** Range of $\delta^{13}\text{C}$ (a), $\delta^{15}\text{N}$ (b) and $\delta^{66}\text{Zn}$ (c) values for *P. hispidus* and *U. maritimus* bones for all
 553 locations. Site names are colour coded following figure 1. Dashed lines represent mean values when
 554 including all sites. We included already published $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values from sites RbJu-1, PaJs-13, QkHn-
 555 13, KTZ and QjJx-1 sites^{4,13,43,44} and already published $\delta^{66}\text{Zn}$ values from QjJx-1¹³.



556

557 **Figure 3:** Isotopic composition (δ¹⁵N, δ¹³C versus δ⁶⁶Zn) of *U. maritimus* (dots), *P. hispida* (squares), *P.*
 558 *groenlandicus* (triangle) and *D. leucas* (star) bones for the SfFk-4 (a, c) and combined KkDo-1 and JfEI-4
 559 sites (b, d).

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570 **Table 1:** Isotopic range ($\delta^{13}\text{C}$, $\delta^{15}\text{N}$, $\delta^{66}\text{Zn}$) for all bone samples discussed in this study for which all three
 571 elements were analysed. Max. = maximum value, min. = minimum value, SD = standard deviation, n =
 572 number of individuals/bone samples, [Zn] Zn concentration.

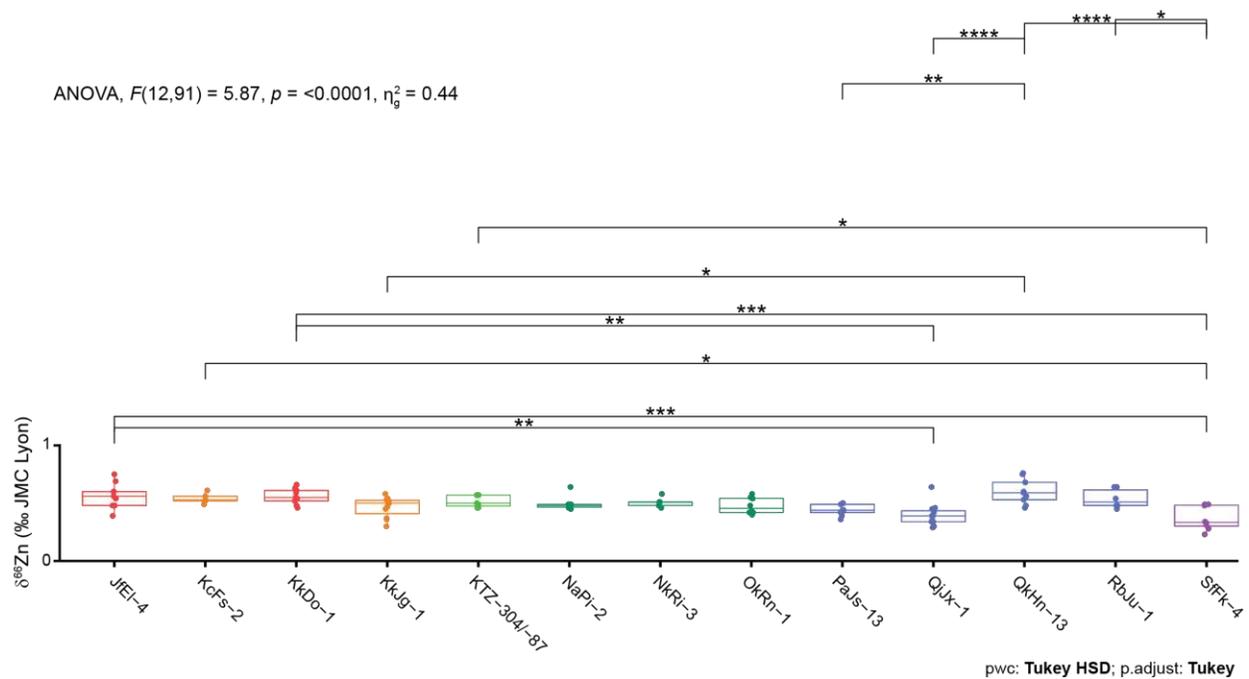
Species	n		$\delta^{13}\text{C}$ (‰, VPDB)	$\delta^{15}\text{N}$ (‰, AIR)	$\delta^{66}\text{Zn}$ (‰, JMC Lyon)	[Zn] ($\mu\text{g/g}$)
<i>U. maritimus</i>	47	max.	-12.49	+24.41	+0.73	901
		min.	-15.04	+16.78	-0.06	80
		mean	-13.68	+21.75	+0.17	276
		SD	0.65	1.72	0.16	155
<i>P. hispida</i>	104	max.	-12.00	+19.56	+0.76	878
		min.	-16.95	+14.25	+0.23	79
		mean	-14.48	+17.22	+0.49	167
		SD	1.13	1.15	0.10	105
<i>P. groenlandicus</i>	11	max.	-13.60	+17.45	+0.66	273
		min.	-14.61	+13.84	+0.22	82
		mean	-14.18	+15.47	+0.46	135
		SD	0.35	1.26	0.14	57
<i>D. leucas</i>	2	max.	-11.24	+18.18	+0.67	1025
		min.	-13.21	+17.75	+0.65	381
		mean	-12.23	+17.97	+0.66	703
		SD	1.39	0.30	0.01	455

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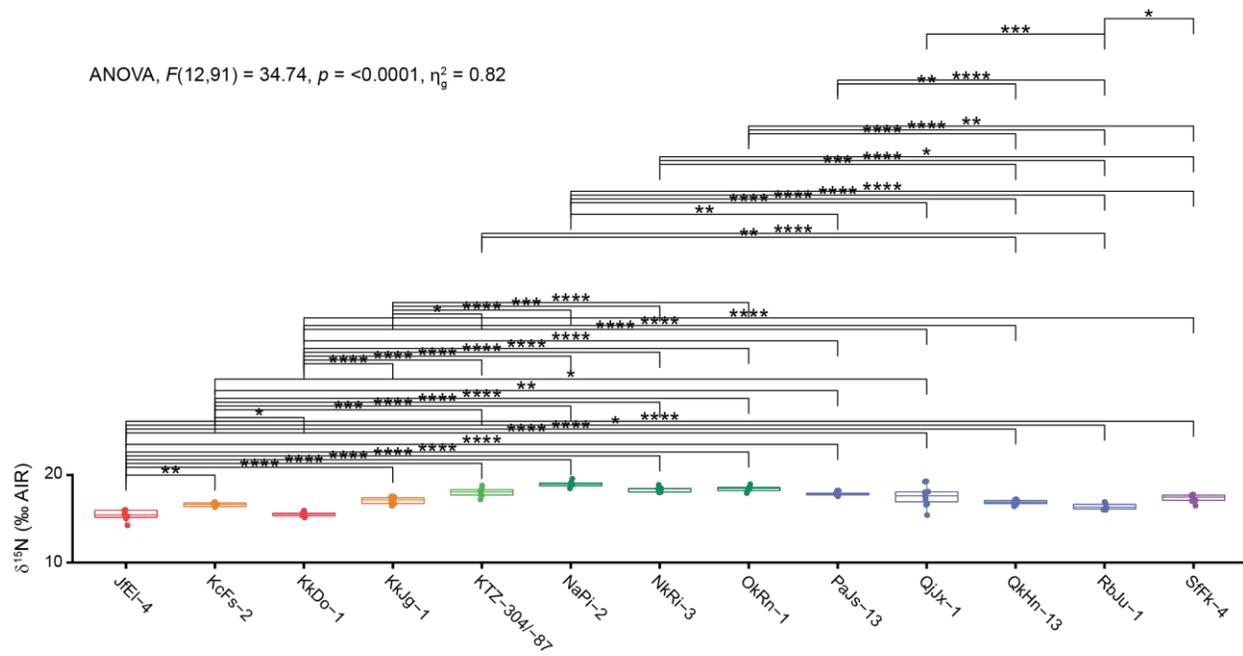
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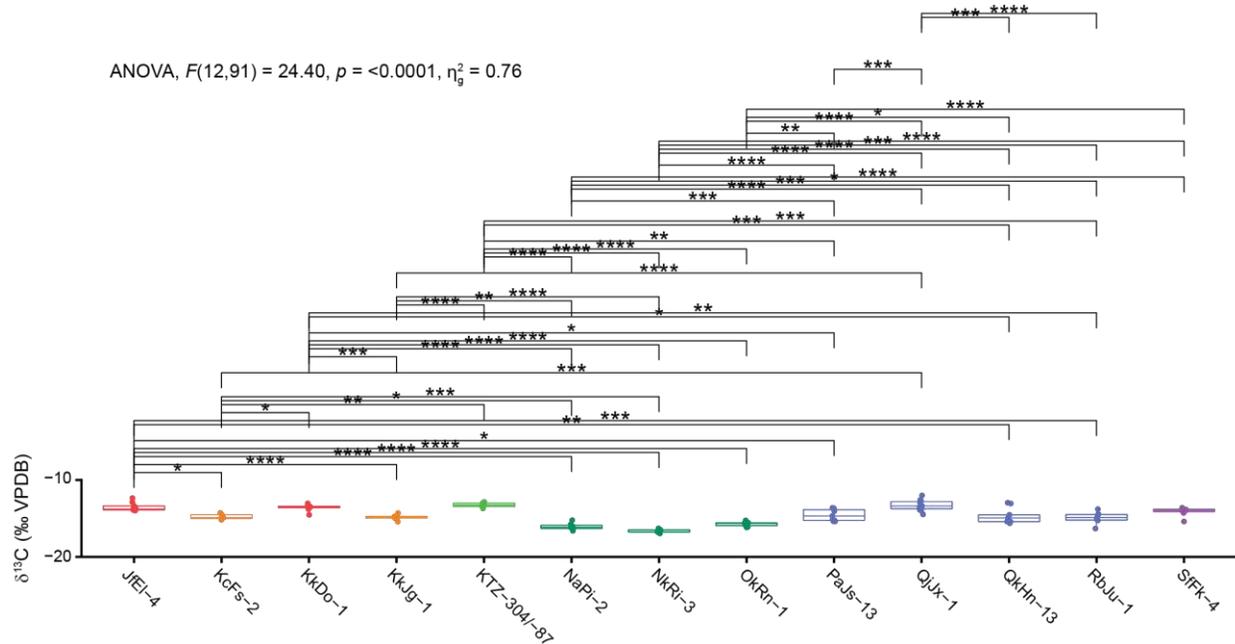
577
 578 **Extended Data Figure 1:** Results for ANOVA testing and post-hoc Tukey pair-wise comparisons between
 579 sites (indicated by their Borden code) for $\delta^{66}\text{Zn}$ values of *P. hispidus*. A single *P. hispidus* specimen from Little
 580 Cornwallis Island¹³ was excluded from statistical analysis as it was singled-out as an extreme outlier ($\delta^{66}\text{Zn}$
 581 = 1.00 ‰). The boxes from the box and whisker plots represent the 25th–75th percentiles, with the
 582 median as a bold horizontal line. Significance level is indicated by “*” (p -value < 0.05), “***” (p -value <
 583 0.005), “****” (p -value < 0.0005) and “*****” (p -value < 0.00005).

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 586 **Extended Data Figure 2:** Results for ANOVA testing and post-hoc Tukey pair-wise comparisons between
 587 sites (indicated by their Borden code) for $\delta^{15}\text{N}$ values of *P. hispidia*. A single *P. hispidia* specimen from Little
 588 Cornwallis Island¹³ was excluded from statistical analysis as it was singled-out as an extreme outlier ($\delta^{66}\text{Zn}$
 589 = 1.00 ‰). The boxes from the box and whisker plots represent the 25th–75th percentiles, with the
 590 median as a bold horizontal line. Significance level is indicated by “*” (p -value < 0.05), “**” (p -value <
 591 0.005), “***” (p -value < 0.0005) and “****” (p -value < 0.00005).

592



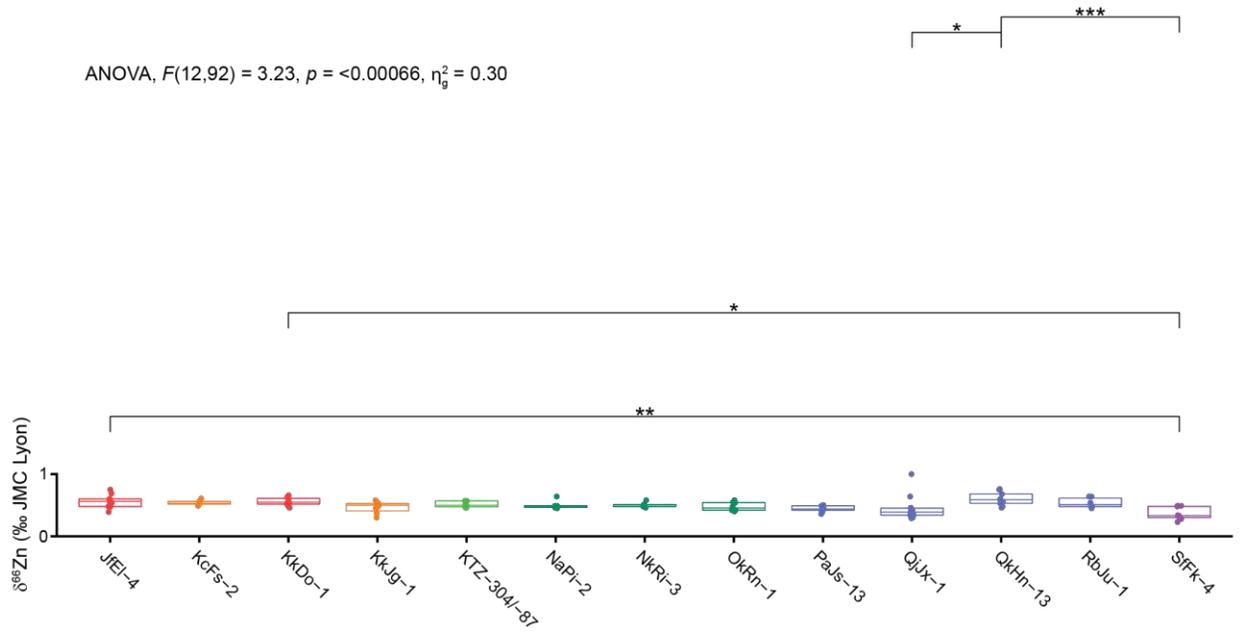
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593
 594 **Extended Data Figure 3:** Results for ANOVA testing and post-hoc Tukey pair-wise comparisons between
 595 sites (indicated by their Borden code) for $\delta^{13}\text{C}$ values of *P. hispidia*. A single *P. hispidia* specimen from Little
 596 Cornwallis Island¹³ was excluded from statistical analysis as it was singled-out as an extreme outlier ($\delta^{66}\text{Zn}$
 597 = 1.00 ‰). The boxes from the box and whisker plots represent the 25th–75th percentiles, with the
 598 median as a bold horizontal line. Significance level is indicated by “*” (p -value < 0.05), “**” (p -value <
 599 0.005), “***” (p -value < 0.0005) and “****” (p -value < 0.00005).

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ANOVA, $F(12,92) = 3.23$, $p = <0.00066$, $\eta_p^2 = 0.30$



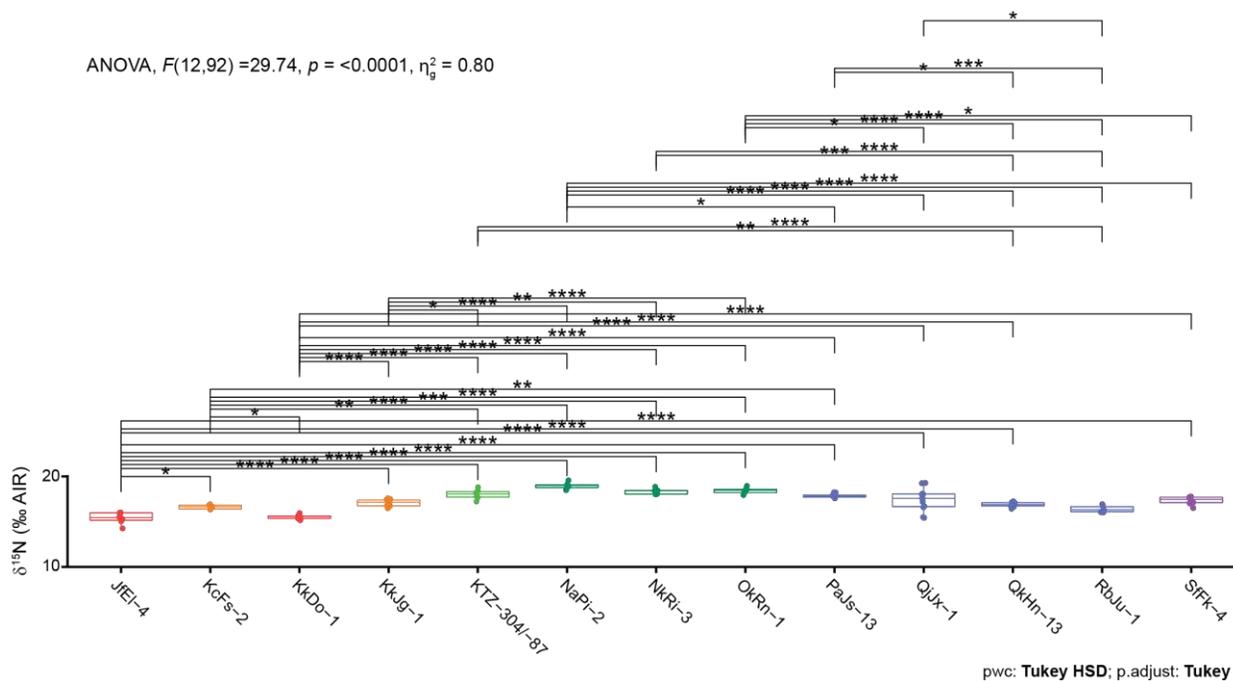
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602
 603 **Extended Data Figure 4:** Results for ANOVA testing and post-hoc Tukey pair-wise comparisons between
 604 sites (indicated by their Borden code) for $\delta^{66}\text{Zn}$ values of *P. hispidus* including an extreme outlier value from
 605 Little Cornwallis Island¹³. The boxes from the box and whisker plots represent the 25th–75th percentiles,
 606 with the median as a bold horizontal line. Significance level is indicated by “*” (p -value < 0.05), “***” (p -
 607 value < 0.005), “****” (p -value < 0.0005) and “*****” (p -value < 0.00005).

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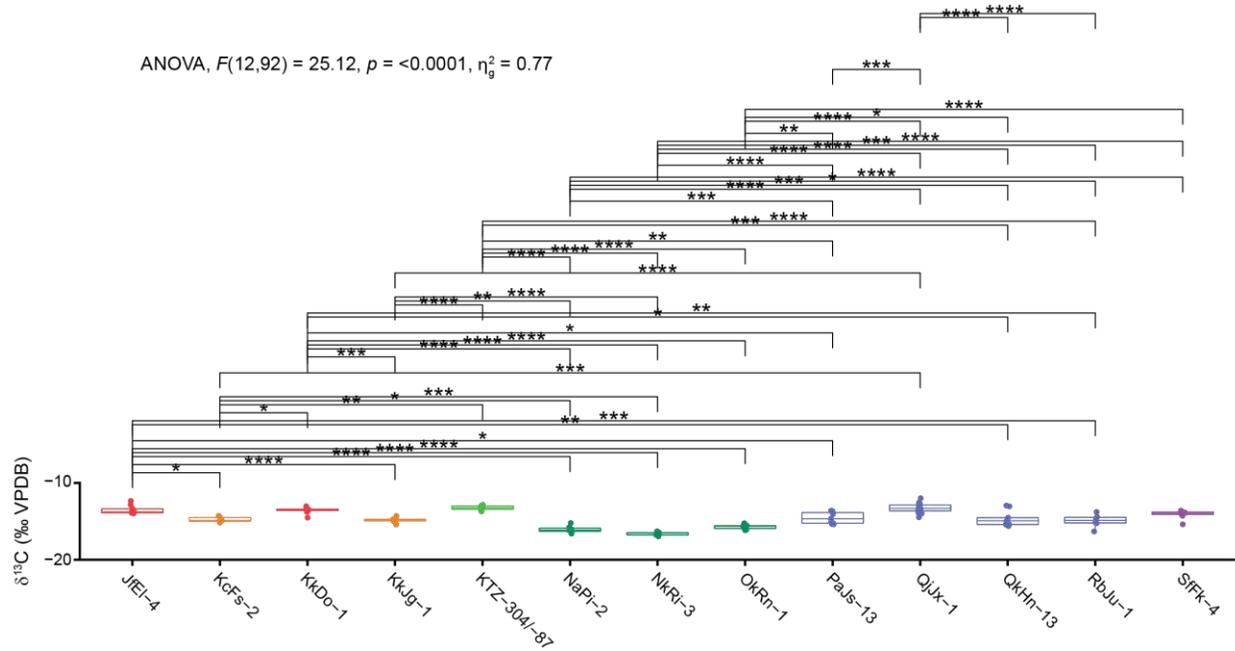
609

ANOVA, $F(12,92) = 29.74$, $p = < 0.0001$, $\eta_p^2 = 0.80$



610
 611 **Extended Data Figure 5:** Results for ANOVA testing and post-hoc Tukey pair-wise comparisons between
 612 sites (indicated by their Borden code) for $\delta^{15}\text{N}$ values of *P. hispida* including an extreme outlier value from
 613 Little Cornwallis Island¹³. The boxes from the box and whisker plots represent the 25th–75th percentiles,
 614 with the median as a bold horizontal line. Significance level is indicated by “*” (p -value < 0.05), “**” (p -
 615 value < 0.005), “***” (p -value < 0.0005) and “****” (p -value < 0.00005).

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 619 **Extended Data Figure 6:** Results for ANOVA testing and post-hoc Tukey pair-wise comparisons between
 620 sites (indicated by their Borden code) for $\delta^{13}\text{C}$ values of *P. hispida* including an extreme outlier value from
 621 Little Cornwallis Island¹³. The boxes from the box and whisker plots represent the 25th–75th percentiles,
 622 with the median as a bold horizontal line. Significance level is indicated by “*” (*p*-value < 0.05), “**” (*p*-
 623 value < 0.005), “***” (*p*-value < 0.0005) and “****” (*p*-value < 0.00005).

Figures

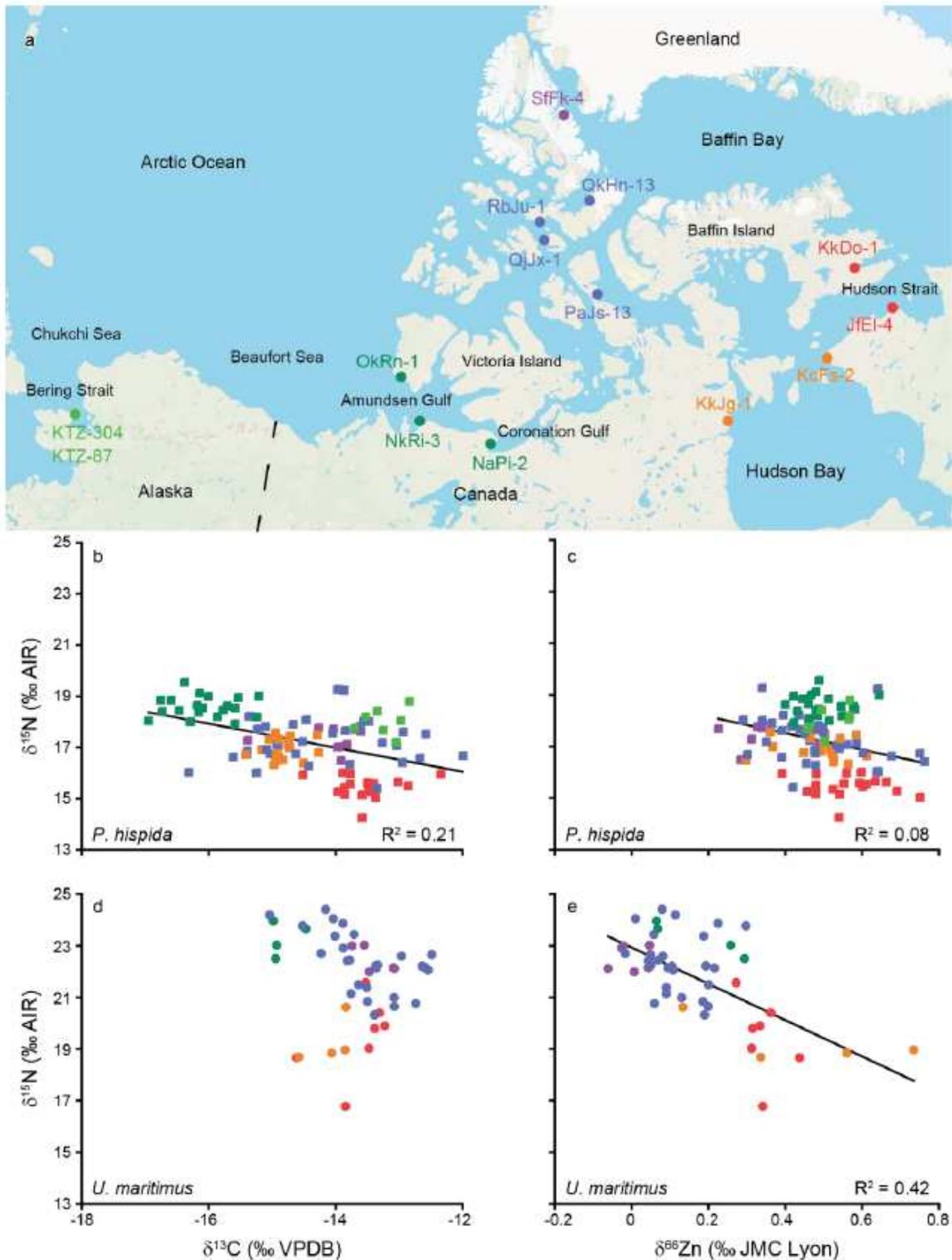


Figure 1

Isotopic composition of *P. hispida* (squares, $n = 104$) and *U. maritimus* (dots, $n = 47$) samples from Arctic archaeological sites colour coded as geographic groups. a) Map indicating the archaeological sites analysed and geographic colour coding: Light green for the Bering Strait; dark green for the Amundsen

and Coronation Gulf; blue for the CAA; orange for the Hudson Bay; purple for the North Water Polynya; and red for sites influenced by the Labrador Sea in the Hudson Strait and Frobisher Bay. b) $\delta^{15}\text{N}$ versus $\delta^{13}\text{C}$ plot for *P. hispida* samples (p-value < 0.05; $R^2 = 0.21$; n = 104). c) $\delta^{15}\text{N}$ versus $\delta^{66}\text{Zn}$ plot for *P. hispida* samples (p-value < 0.05; $R^2 = 0.08$; n = 104). d) $\delta^{15}\text{N}$ versus $\delta^{13}\text{C}$ plot for *U. maritimus* samples (no correlation, p-value > 0.05; n = 47). e) $\delta^{15}\text{N}$ versus $\delta^{66}\text{Zn}$ plot for *U. maritimus* samples (p-value < 0.05; $R^2 = 0.42$; n = 47). We included already published $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values^{4,13,43,44} and already published $\delta^{66}\text{Zn}$ values from QjJx-113. The map is redrawn after www.google.com/maps. Note: The designations employed and the presentation of the material on this map do not imply the expression of any opinion whatsoever on the part of Research Square concerning the legal status of any country, territory, city or area or of its authorities, or concerning the delimitation of its frontiers or boundaries. This map has been provided by the authors.

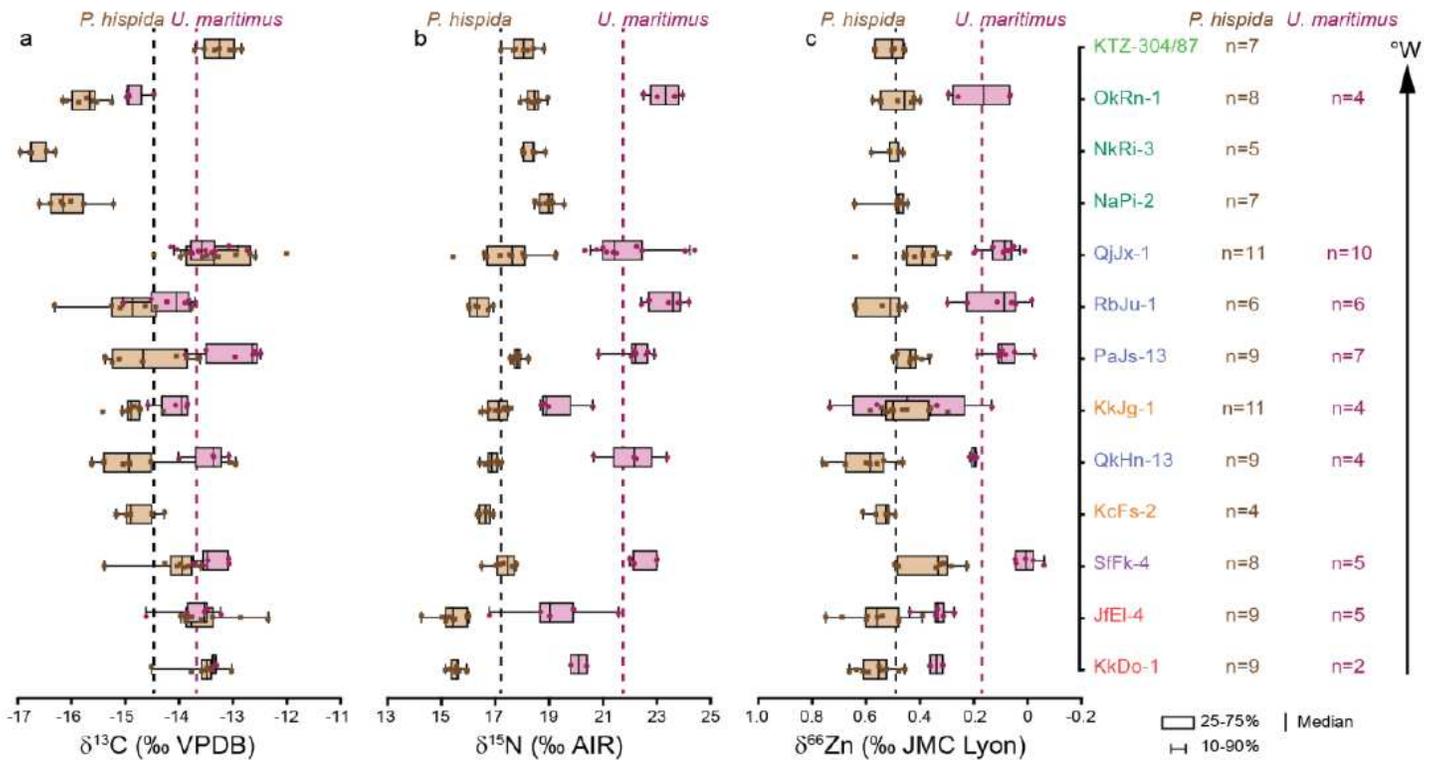


Figure 2

Range of $\delta^{13}\text{C}$ (a), $\delta^{15}\text{N}$ (b) and $\delta^{66}\text{Zn}$ (c) values for *P. hispida* and *U. maritimus* bones for all locations. Site names are colour coded following figure 1. Dashed lines represent mean values when including all sites. We included already published $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values from sites RbJu-1, PaJs-13, QkHn-13, KTZ and QjJx-1 sites^{4,13,43,44} and already published $\delta^{66}\text{Zn}$ values from QjJx-113.

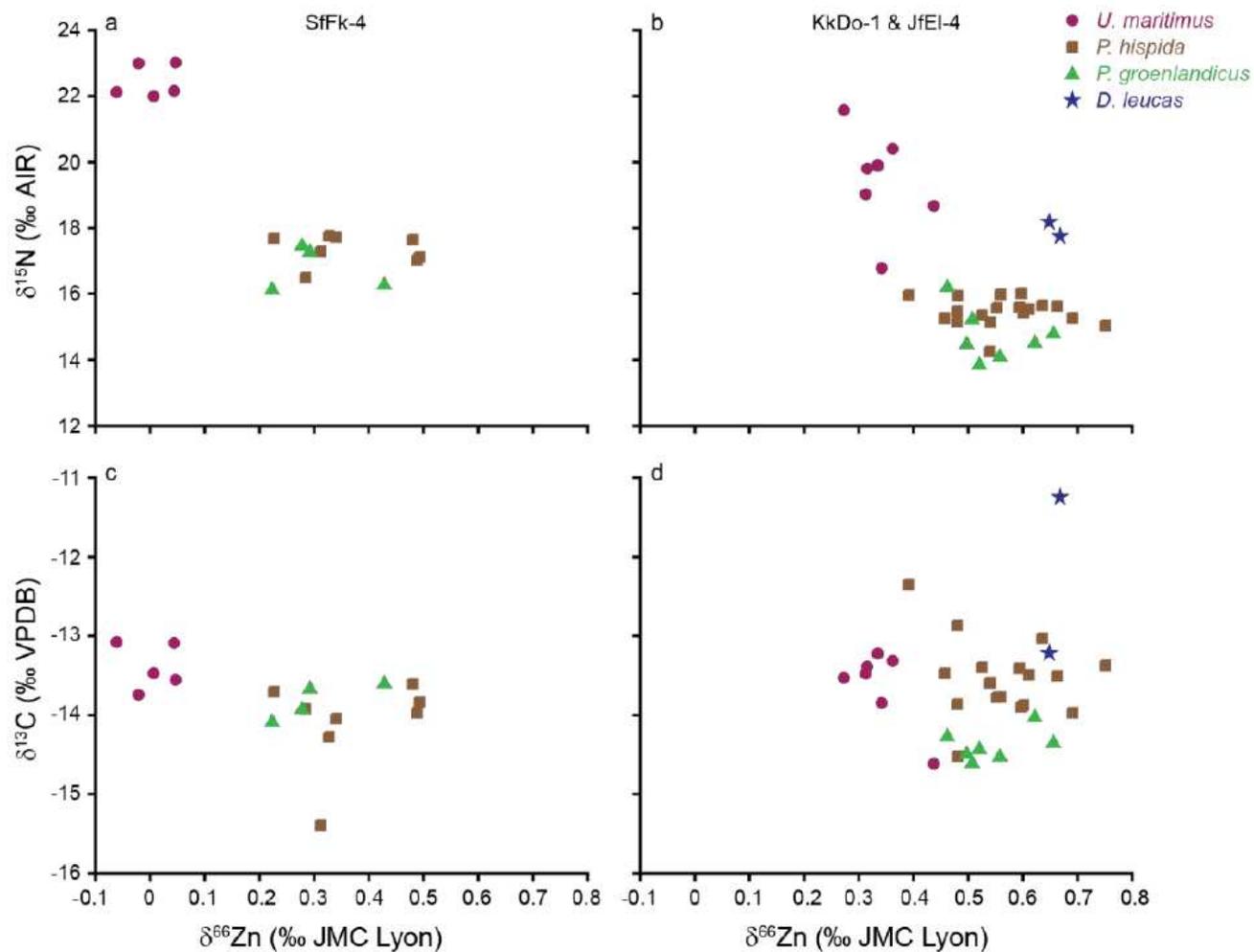


Figure 3

Isotopic composition (δ¹⁵N, δ¹³C versus δ⁶⁶Zn) of *U. maritimus* (dots), *P. hispida* (squares), *P. groenlandicus* (triangle) and *D. leucas* (star) bones for the SIFk-4 (a, c) and combined KkDo-1 and JfEI-4 sites (b, d).

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [SupplementaryInformationMcCormacketal.pdf](#)
- [SupplementaryTable1.xlsx](#)
- [SupplementaryTable2.xlsx](#)