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1 **Compound-specific radiocarbon, stable carbon isotope and biomarker analysis of mixed**  
2 **marine/terrestrial lipids preserved in archaeological pottery vessels**

3  
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15  
16 **ABSTRACT**

17 At archaeological sites located on islands or near the coast, the potential exists for lipid  
18 extracts of potsherds to contain fatty acids (FA) from both aquatic and terrestrial organisms,  
19 meaning that consideration must be given to marine reservoir effects (MRE) in radiocarbon  
20 analyses. Here we studied the site of Bornais (Outer Hebrides, UK) where a local MRE,  $\Delta R$   
21 of  $-65 \pm 45$  y was determined through the paired <sup>14</sup>C determinations of terrestrial and marine  
22 faunal bones. Lipid analysis of 49 potsherds, revealed aquatic biomarkers in 45% of the  
23 vessels, and  $\delta^{13}\text{C}$  values of C<sub>16:0</sub> and C<sub>18:0</sub> FAs revealed ruminant and marine product mixing  
24 for 71% of vessels. Compound-specific radiocarbon analysis (CSRA) of FAs yielded  
25 intermediate radiocarbon ages between those of terrestrial and marine bones from the same  
26 contexts, confirming an MRE existed. A database containing  $\delta^{13}\text{C}$  values for FAs from  
27 reference terrestrial and marine organisms provided endmembers for calculating the  
28 percentage marine-derived C ( $\%_{\text{marine}}$ ) in FAs. We show that lipid <sup>14</sup>C dates can be corrected  
29 using determined  $\%_{\text{marine}}$  and  $\Delta R$  values, such that pottery vessels from coastal locations can  
30 be radiocarbon dated by CSRA of FAs.

31  
32 **KEYWORDS:** Pottery vessels, Lipid residues, Compound-specific radiocarbon analysis,  
33 Marine reservoir effect, Mixed marine/terrestrial corrections.

35

## 36 INTRODUCTION

37 Due to their central importance to, and survival in the archaeological record, accurate direct  
38 radiocarbon dating of pottery vessels has been one of the “Holy Grails” of archaeology.  
39 Compound-specific radiocarbon dating of lipids preserved within the clay matrix of  
40 archaeological potsherds is technically extremely challenging, with previous attempts failing  
41 to achieve the accuracy and precision required (e.g. Hedges *et al.* 1992, Stott *et al.* 2001,  
42 2003b; Berstan *et al.* 2008). Recently however, we have reported the first accurate dates  
43 achieved for such residues based on compound-specific <sup>14</sup>C analyses of C<sub>16:0</sub> and C<sub>18:0</sub> fatty  
44 acids (FAs) isolated from the clay walls of Neolithic pottery vessels (Casanova *et al.*, in  
45 press). The preparative-capillary gas chromatography (pcGC) isolation technique required  
46 two major advances, namely a new trap design allowing the solvent-less recovery of the  
47 trapped analytes and a heat-based cleaning method to prevent cross-contamination (Casanova  
48 *et al.* 2018). These methodological improvements have enabled reliable and accurate dating  
49 of the two FAs characteristic of degraded animal fats. Furthermore, the two independent  
50 radiocarbon dates obtained provide an important internal quality control; the radiocarbon age  
51 of the FAs should agree at the 2σ error level (Casanova *et al.* 2018). The samples used thus  
52 far in the validation of the compound-specific pot lipid dating method, outlined in Casanova  
53 *et al.* (in press) have originated from archaeological sites located inland where human dietary  
54 subsistence was dominated by domesticated terrestrial animals, such that the target FAs  
55 derived from dairy or carcass fats of ruminant and non-ruminant animals.

56 None of the pottery dated thus far has originated from coastal areas where the exploitation of  
57 marine products may have occurred. At such locations, FAs preserved in pottery vessels  
58 would likely be affected by a reservoir effect (Heron and Craig 2015), requiring marine  
59 reservoir correction in order to obtain reliable calibrated dates (Cook *et al.* 2015). Particularly  
60 problematic would be potsherds containing mixed marine- and terrestrial-derived FAs  
61 (Cramp and Evershed 2014; Cramp *et al.* 2014a), as this would increase the complexity of  
62 marine reservoir corrections.

63 Marine product processing in pots can be identified by the presence of specific aquatic  
64 biomarkers alongside the C<sub>16:0</sub> and C<sub>18:0</sub> FA, namely: (i) long-chain dihydroxy fatty acids  
65 (DHYAs), (ii) isoprenoid fatty acids (IFAs) and (iii) long-chain ω-(o-alkylphenyl)alkanoic

66 acids (APAAs); (Hansel *et al.* 2004; Evershed *et al.* 2008; Hansel and Evershed 2009;  
67 Cramp and Evershed 2014). Furthermore,  $\delta^{13}\text{C}$  values determined for the  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$  FAs  
68 can reveal the mixing of both terrestrial and marine commodities in the same vessel (Copley  
69 *et al.* 2004, Cramp *et al.* 2014a, 2014b). It is known, however, that the relative abundances of  
70  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$  FAs differ between terrestrial and marine organisms and the relationship  
71 between their mixing proportions and the resulting  $\delta^{13}\text{C}$  values is not necessarily linear  
72 (Mukherjee *et al.* 2005). It is unclear whether this effect will adversely affect the validity of  
73 the internal quality control criteria, such that the  $^{14}\text{C}$  dates obtained for  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$  FAs in  
74 a potsherd are no longer consistent within  $2\sigma$  (Casanova *et al.* 2018, in press). It is certainly  
75 possible that  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$  FAs in sherds arising from mixtures of terrestrial- and marine-  
76 derived food residues may yield different apparent radiocarbon ages.

77 Generally, MRE corrections require generation of terrestrial/marine mixing curves using  
78 dedicated software (e.g. OxCal, CALIB). This requires an understanding of the local  
79 deviation ( $\Delta R$ ) from the global marine calibration curve for a specific time period as well as  
80 the percentage of marine-derived C ( $\%_{\text{marine}}$ ) present (Cook *et al.* 2015). The  $\Delta R$  values can  
81 be obtained by radiocarbon dating historical marine specimens (of known date of collection)  
82 from museum collections, pairing  $^{14}\text{C}$  measurements on terrestrial and marine organisms  
83 from secure contexts at the site of interest or by dating tephra layers deposited at sea and on  
84 land (Ascough *et al.* 2005). The evaluation of the  $\%_{\text{marine}}$ , however, is more challenging. Such  
85 considerations are often applied to bone collagen from omnivores which can feed on both  
86 terrestrial and marine resources (Cook *et al.* 2015). Typically,  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values are  
87 recorded on bulk collagen to understand the local diet and the percentage of marine resources  
88 consumed. Preferably, endmembers for pure terrestrial and pure marine organisms are  
89 recorded for samples local to the site, but in the majority of cases, more general (non-local)  
90 reference values for endmembers are used (Cook *et al.* 2015).

91 Herein, we evaluate whether the approach commonly applied to bone collagen to estimate the  
92 contribution of aquatic resources could be applied to FAs extracted from pottery vessels for  
93 MRE correction of pot lipids  $^{14}\text{C}$  dates. The approach was to undertake radiocarbon dating in  
94 order to determine the influence of aquatic resources on CSRA of lipids from potsherds and  
95 establish appropriate methods to correct for the MRE. We focussed on lipids preserved in  
96 pottery vessels with a clearly mixed marine/terrestrial signal from the site of Bornais (South  
97 Uist, UK). Our approach involved: (i) lipid residue analyses on pottery vessels including  
98 compound-specific  $\delta^{13}\text{C}$  determinations on FAs, (ii) calculation of the local deviation from

99 the global marine calibration curve at the site using paired marine and terrestrial animal  
100 remains, (iii) radiocarbon dating of FAs from a range of pottery vessels, (iv) a multiproxy  
101 investigation (i.e. biomarkers, stable isotopes and  $^{14}\text{C}$  analyses) to evaluate the proportion of  
102 mixing of marine and terrestrial lipids and (v) application of relevant marine reservoir  
103 corrections to the radiocarbon dates obtained from pot lipids.

104

## 105 **METHODS**

### 106 **Site description**

107 The site of Bornais is located on the island of South Uist, in the Outer Hebrides, UK  
108 (Supplementary material S1). The site comprises 4 mounds with a long duration of  
109 occupation defined by 109 radiocarbon dates on seeds and bone collagen, from the late Iron  
110 Age (LIA 1 and LIA 2; 5<sup>th</sup>-6<sup>th</sup> century AD) to the Early, Middle and Late Norse (EN, MN,  
111 LN, respectively; mid-9<sup>th</sup>-14<sup>th</sup> century AD) period (Marshall 2005, 2016, forthcoming;  
112 Sharples forthcoming). The recovery of plant macrofossils indicates the cultivation of rye and  
113 barley, while the faunal assemblage displays a particularly rich diversity of terrestrial animals  
114 (*c.* 18,000 bones), small vertebrates, birds, fish and molluscs; Sharples and Davis  
115 forthcoming, Sharples, *et al.* 2016). Domesticated animals dominate ( $\sim 95\%$ ) the terrestrial  
116 faunal assemblage, which comprised cattle (*ca.* 40%), sheep (*ca.* 45%) and pigs (*ca.* 10%).  
117 The mortality profiles derived from the cattle suggest they were exploited for their milk  
118 (Sharples, *et al.* 2016). Fish bones (eel, saithe, cod, haddock, ray, turbot, mackerel etc.) and  
119 mollusc shells (limpets and winkles) were extremely abundant at the site (*c.* 17,000 identified  
120 specimens), while marine mammal bones, e.g. seal, were rare (Sharples, *et al.* 2016).

### 121 **Lipid residue analysis**

122 Lipid residue analyses of samples of pottery were performed using a methanolic sulphuric  
123 acid extraction procedure (Correa-Ascencio and Evershed 2014). The total lipid extracts  
124 (TLEs) were analysed by gas chromatography (GC) and GC-Mass spectrometry (GC-MS) for  
125 the identification and quantification of biomarkers, including aquatic biomarkers following  
126 established procedures (Evershed *et al.* 1990, Cramp and Evershed 2014). Compound-  
127 specific  $\delta^{13}\text{C}$  values of fatty acids were determined by GC-Combusted-Isotope ratio MS (GC-  
128 C-IRMS; supplementary material S2).

### 129 **Pretreatment methods for radiocarbon analyses**

130 Approximately 300 mg of coarse bone powder were weighed into a culture tube and pre-  
131 treated using a modified Login procedure (Longin 1971) as described in Knowles, *et al.*  
132 (2019). Briefly, bone powder was demineralized in HCl (0.5 M, 10 mL, ~ 18h, room  
133 temperature (RT)) followed by a wash with NaOH (0.1 M, 10 mL, 30 min, RT) and a second  
134 acid wash with HCl (0.5 M, 10 mL, 30 min, RT). The extracted collagen was rinsed with  
135 ultrapure MilliQ-water (MQ-water; 3 x 10 mL) in between each acid and base wash and  
136 centrifuged (3000 rpm, 5 min). The collagen was then gelatinised at pH 3 with HCl (0.001 M,  
137 10 mL, 75 °C, 20 h) and filtered through pre-combusted glass fibre before freeze drying  
138 (Knowles, *et al.* 2019).

139 Surface cleaned shells were ultrasonically agitated in MQ-water (5 mL, 5 min) before drying  
140 at 60°C. When dried, the shells (~ 30 mg) were crushed roughly before the surface was acid  
141 etched (~20 %) with HCl (0.2 M, 10 mL). Samples were rinsed with MQ-water (3 x 10 mL)  
142 and dried at 60°C in a drying cabinet (Knowles, *et al.* 2019).

143 Sherds containing lipid concentrations, typically above 500 µg.g<sup>-1</sup>, were selected for  
144 radiocarbon determinations. Pieces of 2 to 10 g of the potsherd was sampled, depending on  
145 the lipid concentrations and size of the potsherds. The lipids were extracted in culture tubes  
146 using H<sub>2</sub>SO<sub>4</sub>/MeOH (4 % v/v, 3 x 8 mL, 70°C, 1 h). Samples were centrifuged after each  
147 extraction (2500 rpm, 10 min) and the 3 supernatants (methanolic fractions) combined into a  
148 second culture tube containing double-distilled water (5 mL). The lipids, including fatty acid  
149 methyl esters (FAMES) created from the reaction of methanol with the FAs during the first  
150 step, were extracted from the methanolic solution with *n*-hexane (4 x 5 mL) and blown down  
151 to dryness at room temperature under a gentle nitrogen stream. The TLEs were derivatized  
152 with BSTFA (20 µL, 70 °C, 1 h). Excess BSTFA (*N,O*-bis(trimethylsilyl)trifluoroacetamide)  
153 was removed under a nitrogen stream, then ~180 µL of *n*-hexane was added to obtain a  
154 solution containing C<sub>16:0</sub> and C<sub>18:0</sub> FAMES at a concentration at *c.* 5 µg.µL<sup>-1</sup> of carbon. The  
155 solution was transferred to an autosampler vial for isolation of C<sub>16:0</sub> and C<sub>18:0</sub> into individual  
156 traps using a preparative capillary gas chromatography (pcGC) instrument following the  
157 methods described in Casanova *et al.* (2017, 2018, in press).

## 158 **Radiocarbon determinations**

159 Organic materials (FAMES and collagen) were combusted to CO<sub>2</sub> using a Vario Microcube  
160 Elemental Analyser (EA, Elementar). The shells (carbonate-based) were digested in H<sub>3</sub>PO<sub>4</sub> (1

161 mL, 85 % v/v, 70 °C) under a He headspace using a Carbonate Handling System (CHS,  
162 Ionplus; Wacker, *et al.* 2013; Knowles, *et al.* 2019) to generate CO<sub>2</sub>. Resulting CO<sub>2</sub> was  
163 transferred to the Automated Graphitisation Equipment (AGE 3, Ionplus; Wacker *et al.* 2010;  
164 Knowles, *et al.* 2019) under a He stream and adsorbed on Zeolite traps before being released  
165 into reaction tubes. The CO<sub>2</sub> was reduced to graphite under H<sub>2</sub> (580 °C, 2 h, 420 mbar) on a  
166 pre-conditioned iron catalyst. A Pneumatic Sample Press (PSP, Ionplus) was used to press the  
167 graphitised samples into Al targets.

168 All <sup>14</sup>C determinations were performed at the BRAMS (Bristol Accelerator Mass  
169 Spectrometer) facility which is equipped with a mini radiocarbon dating system (BRIS-  
170 MICADAS) instrument (ETH Zurich, Zurich, Switzerland; Synal *et al.* 2007). Samples were  
171 analysed alongside size-matched processing standards and blanks (Casanova *et al.* 2018;  
172 Knowles *et al.* 2019).

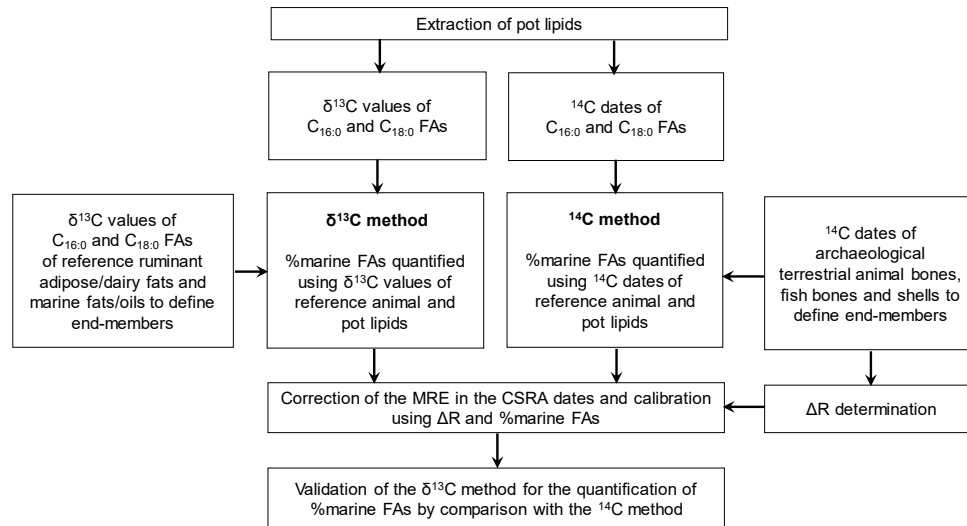
### 173 **Corrections and calibration of <sup>14</sup>C measurements**

174 Radiocarbon measurement on FAs from single pottery vessels were corrected for the  
175 presence of the methyl derivative C (Casanova *et al.* 2017, 2018) and subjected to a 2-sigma  
176 equivalency test and, if successful, combined as described in Casanova *et al.* (in press) before  
177 testing the validity of calibration on mixed marine/terrestrial resources. Reservoir correction  
178 and calibration of the mixed resources was performed in OxCal v 4.3 (Bronk Ramsey 2009)  
179 using the 'Marine/mixed curve' tool using the IntCal13 and Marine13 curves (Reimer *et al.*  
180 2013). This incorporates the percentage of marine derived resources present in the TLEs and  
181 the  $\Delta R$  value for the site (see results section).

182 The local reservoir effect was calculated for every pair combination of terrestrial/marine  
183 organisms in each context using the online  $\Delta R$  calculation tool (Reimer and Reimer, 2016).  
184 These individual  $\Delta R$  values were subjected to a  $\chi^2$  test at the 5 % level (both for each context  
185 and all together) to detect potential outliers before calculation of their weighted average, with  
186 error calculation as recommended by Russell *et al.* (2010).

187 The mixing of marine/terrestrial commodities was quantified in each potsherd using two  
188 independent methods: <sup>14</sup>C dates and  $\delta^{13}\text{C}$  values of C<sub>16:0</sub> and C<sub>18:0</sub> FAs (Figure 1). By  
189 comparing the mixing ratios obtained by the two methods, it is possible to evaluate whether

190 the FA  $\delta^{13}\text{C}$  values (determined by GC-C-IRMS) can be used to estimate the proportion of  
 191 marine-derived C in the FAs for use in MRE corrections of their radiocarbon dates. This  
 192 constitutes an important consideration, especially for sites where terrestrial and marine  
 193 remains are absent from the archaeological record and so cannot be used to provide reference  
 194 radiocarbon ages.



195  
 196 Figure 1

197 The first method of quantifying the  $\%_{marine}$  is based on the weighted average of radiocarbon  
 198 determinations on the short-lived terrestrial and marine organisms, from the same  
 199 context/phase as the potsherds dated, as endmembers using Equation (1).

200 (1) 
$$\%_{marine} = \frac{(Age_{pot} - Age_{terr})}{(Age_{marine} - Age_{terr})} * 100$$

201 Where  $\%_{marine}$  is the percentage of aquatic C in the lipid residue and  $Age_{pot}$ ,  $Age_{terr}$ ,  $Age_{marine}$   
 202 are the combined radiocarbon ages on the individual FAs, for terrestrial animals and marine  
 203 organisms, respectively.

204 The second method uses the  $\delta^{13}\text{C}$  values of the individual FAs of UK reference animals  
 205 (cattle and sheep raised on a pure  $\text{C}_3$  diet; Copley, *et al.* 2003) as the terrestrial endmembers  
 206 (pigs were hypothesised not to have been processed in potsherds; Sharples, *et al.* 2016, and  
 207 fish, winkles and limpets captured from UK waters (corrected for the Suess effect; Cramp  
 208 and Evershed, 2014) to serve as the marine endmembers. The terrestrial endmembers  
 209 correspond to the average values for both  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$  FAs and were found to be



210  $\delta^{13}\text{C}_{16:0} = -30.0 \pm 0.6 \text{ ‰}$  and  $\delta^{13}\text{C}_{18:0} = -32.2 \pm 0.6 \text{ ‰}$  for adipose fats and  $\delta^{13}\text{C}_{16:0} = -29.2 \pm$   
211  $1.0 \text{ ‰}$  and  $\delta^{13}\text{C}_{18:0} = -34.0 \pm 0.9 \text{ ‰}$  for dairy fats. Both ruminant adipose and dairy values  
212 were used as endmembers to evaluate whether one should be used over the another. The  
213 marine endmembers were  $\delta^{13}\text{C}_{16:0} = -22.7 \pm 2.2 \text{ ‰}$  and  $\delta^{13}\text{C}_{18:0} = -21.7 \pm 2.5 \text{ ‰}$ . The  
214 relationship between the relative proportions of marine and terrestrial fats and the  $\delta^{13}\text{C}$  values  
215 is theoretically non-linear, due to the differing relative abundances of the of FAs in the  
216 different foodstuffs (Mukherjee, *et al.* 2005), however, the success of the internal quality  
217 control on the CSRA dates (see results section) suggests a linear relationship within  
218 analytical uncertainty. A linear mixing curve could therefore be employed to estimate the  
219  $\%_{\text{marine}}$  contribution and associated uncertainty using the propagation of analytical errors. The  
220  $\%_{\text{marine}}$  values obtained on both FAs were then combined as a weighted average with  
221 uncertainties calculated according to Russell *et al.* (2010). Such a model is a conservative  
222 approach and probably overestimates the uncertainties. Furthermore, it cannot take into  
223 account the fact that the true  $\%_{\text{marine}}$  values must be constrain between 0 and 100 %.

224 As a comparison, the  $\%_{\text{marine}}$  values of the TLEs were also estimated (using the same  
225 endmembers) using the software package FRUITS (v2.1). This software employs a Bayesian  
226 approach to quantify the contribution of different food sources using isotopic data (Fernandes  
227 *et al.* 2014). The output of this software is given both as means and standard deviations  
228 (represented by box-and whiskers plots) or as probability distributions constrained to between  
229 0 and 100 %. The full range of data points for the probability distributions of the  $\%_{\text{marine}}$  after  
230 Bayesian modelling was exported and implemented as a prior information file into the mixing  
231 marine/terrestrial tool in OxCal.

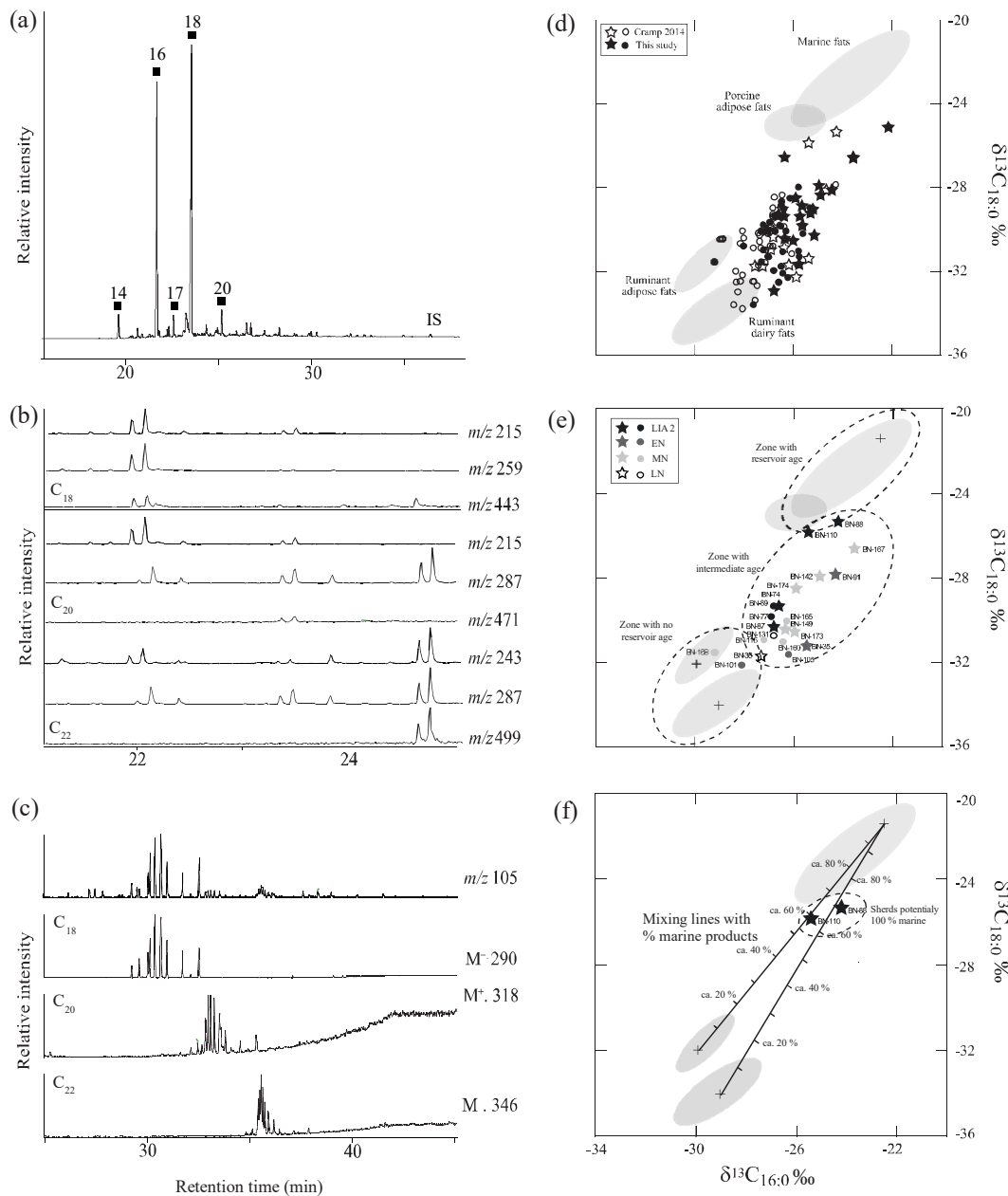
## 232 **RESULTS AND DISCUSSION**

### 233 **Characterisation of lipid residues in pottery vessels**

234 Forty-nine pottery vessels from layer BCC, Mound 2, MN period were subjected to lipid  
235 residue analyses (S2). TLEs with concentrations  $>5 \mu\text{g.g}^{-1}$  were recovered from 96 % ( $n =$   
236 47) of the potsherds, at an average lipid concentration of  $1.2 \text{ mg.g}^{-1}$  (supplementary material  
237 S3). A total of 80 % of the TLEs with residues ( $n = 39$ ) were dominated by the  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$   
238 fatty acids characteristic of degraded animal fats (Figure 2a). Many of the TLEs (47 % of the  
239 sherds with residues;  $n = 22$ ) exhibited marine biomarkers. The long-chain DHYAs ( $\text{C}_{18}$ ,  $\text{C}_{20}$   
240 and  $\text{C}_{22}$ ) were detectable in 30 % ( $n = 14$ ; Figure 2b), long-chain APAAs ( $\text{C}_{18}$ ,  $\text{C}_{20}$  and  $\text{C}_{22}$ ) in

241 23 % ( $n = 11$ ; Figure 2c) and the IFAs (phytanic acid and 4,8,12-trimethyltridecanoic acid  
242 (TMTD)) in 30 % ( $n = 14$ ) of the sherds with residues. In total, only 4 % ( $n = 2$ ; BN-140,  
243 BN-173) of the potsherds with lipid residues contained all 3 classes of aquatic biomarkers, 21  
244 % ( $n = 10$ ) contained 2 aquatic biomarkers and 26 % ( $n = 12$ ) showed one aquatic biomarker.  
245 No aquatic biomarkers were detected in the remainder of the TLEs ( $n = 25$ , 53 % of the  
246 sherds with organic residues). The  $\delta^{13}\text{C}$  values of the palmitic and stearic acids were  
247 determined by GC-C-IRMS (Figure 2d). Significantly, the  $\text{C}_{16:0}$  and  $\text{C}_{18:0}$  fatty acids displayed  
248  $\delta^{13}\text{C}$  values characteristic of mixtures between ruminant and marine or porcine products  
249 (Cramp *et al.* 2014a, 2014b). The extracts yielding the most enriched stable carbon isotope  
250 values also contained aquatic biomarkers, strongly suggesting the processing of marine  
251 products rather than porcine. Several TLEs were relatively more enriched in  $^{13}\text{C}$ , but show no  
252 detectable aquatic biomarkers suggesting they did not survive or could denote the processing  
253 of marine commodities under conditions not conducive to the formation of thermally-  
254 produced aquatic biomarkers (APAAs). The use of  $\Delta^{13}\text{C}$  ( $= \delta^{13}\text{C}_{18:0} - \delta^{13}\text{C}_{16:0}$ ) values allows  
255 the identification of sherds where FAs are predominantly of dairy product origin ( $< -3.1$  ‰;  
256 Copley *et al.* 2003). A total of 22 sherds yielded  $\Delta^{13}\text{C}$  values below  $-3.1$  ‰, with aquatic  
257 biomarker identification for half of them supports the hypothesis of some mixing of dairy  
258 products with marine products. The other sherds with higher  $\Delta^{13}\text{C}$  values of  $> -3.1$  ‰ could  
259 result from the mixing of ruminant carcass and marine products.

260 Additionally, 131 potsherds from all the phases and mounds at the site were previously  
261 analysed by lipid residue analysis (Cramp *et al.* 2014b, forthcoming). The results suggested a  
262 dominance of dairy and ruminant carcass product processing, as well as some mixing of non-  
263 ruminant and marine fat/oil (Figure 1d). Only the pottery from the LIA 1 phase lacked  
264 aquatic biomarkers.



265

266 **Figure 2**

267 A total of 21 potsherds (from all phases) with sufficient lipid concentrations and containing  
 268 either none or at least one aquatic biomarker, were subjected to CSRA (Figure 1e).

269 **ΔR value determination**

270 In order to determine the age of the structures associated with the pots, and the local reservoir  
 271 effect at the site of Bornais a range of fish bones ( $n = 13$ ), marine mollusc shells ( $n = 14$ ) and  
 272 terrestrial animal bones ( $n = 8$ ) were radiocarbon dated (Table 1). These were assessed  
 273 together with other available radiocarbon measurements on terrestrial animal bones and

274 grains (layer BCC,  $n = 7$ ; Marshall *et al.* forthcoming-b). All these materials derive from the  
 275 LIA2, EN, MN and LN settlement structures. On a context-by-context basis, marine and  
 276 terrestrial organisms were subjected to  $\chi^2$  statistical testing to detect outliers for exclusion  
 277 (Table 1). Two marine samples from context BAF and two terrestrial animal bones from  
 278 context BCC were, therefore, excluded from  $\Delta R$  determination. The  $\Delta R$  values calculated  
 279 using all the pairs of terrestrial/marine organism (80 in total) per context are reported in Table  
 280 1. No  $\Delta R$  was calculated for contexts BBA and BBD as they were dated based on only one  
 281 material type, and for AG, the two marine organisms from this context failed the  $\chi^2$  test.

282 With the exception of context BCC, all the contexts demonstrated a negative  $\Delta R$ , varying  
 283 from  $-214 \pm 26$  to  $-45 \pm 21$ . Interestingly, layer BCC (MN phase) shows a  $\Delta R$  of  $28 \pm 150$ ;  
 284 the large uncertainty associated with this value results from high variability in the  
 285 radiocarbon ages of the marine organisms, which could be classified into three distinct  
 286 groups. Group (a) gave a  $\Delta R$  of  $-107 \pm 54$ , Group (b)  $242 \pm 55$  and Group (c)  $-31 \pm 56$ . The  
 287 MRE of Group (c), comprised only fish bones and likely reflects the mobility of the fish  
 288 species (Russell, *et al.* 2011). Groups (a) and (b) comprise both winkles and limpets and their  
 289 MREs do not appear to be species dependent. The grouping could, therefore, either  
 290 correspond to two different collection points of the mollusc shells (likely collection points  
 291 nearby are either completely coastal, or sea lochs with the potential for substantial terrestrial  
 292 runoff) or simply the introduction of older material into a later context (although, this offset  
 293 was only observed for some limpet and winkle shells, but not fish bones).

294

295 Table 1: Radiocarbon determinations on terrestrial and marine organisms at Bornais and  $\Delta R$   
 296 calculated for the diverse contexts based on the multiple paired terrestrial/marine organisms. \*refer to  
 297 statistical outliers that have been excluded from  $\Delta R$  calculation.

Phase	Layer	Terrestrial organisms			Marine organisms			$\Delta R$
		Material	Laboratory#	Conventional $^{14}\text{C}$ age	Material	Laboratory#	Conventional $^{14}\text{C}$ age	
LIA 2	BAC	Cattle	BRAMS-1710	$1,272 \pm 25$	Limpet-1	BRAMS-1727	$1,624 \pm 26$	$-47 \pm 23$
		Caprine	BRAMS-1711	$1,298 \pm 25$	Limpet-2	BRAMS-1728	$1,627 \pm 26$	
		Unidentified	BRAMS-1713.1	$1,258 \pm 25$	Limpet-5	BRAMS-1731	$1,642 \pm 26$	
			BRAMS-1713.2	$1,264 \pm 25$	Limpet-6	BRAMS-1732	$1,616 \pm 26$	
	BAF	Cattle	BRAMS-1712	$1,348 \pm 25$	Fish-9	BRAMS-1725*	$1,651 \pm 25$	$-165 \pm 26$
					Fish-10	BRAMS-1726*	$1,294 \pm 25$	
				Limpet-3	BRAMS-1729	$1,565 \pm 26$		
				Limpet-4	BRAMS-1730	$1,584 \pm 26$		
BAG	Cattle	BRAMS-1708	$1,320 \pm 25$	Fish-1	BRAMS-1717	$1,509 \pm 25$	$-214 \pm 26$	
	Cattle	BRAMS-1709	$1,320 \pm 25$					
EN	BBD	Cattle	BRAMS-1715*	$1,082 \pm 25$	-		-	
		Cattle	BRAMS-1719*	$945 \pm 25$				
	BBA	-			Limpet-7	BRAMS-1733	$1,622 \pm 26$	-

MN	BCC	Cattle	SUERC-2684	925 ± 35	Fish-11	BRAMS-2049 (c)	1,306 ± 25	-102 ± 35 (a) 248 ± 37 (b) -26 ± 40 (c) 35 ± 150 (all)
		Red deer	SUERC-22894*	875 ± 30	Fish-12	BRAMS-2050 (c)	1,318 ± 25	
		Pig	SUERC-22890*	1,035 ± 30	Fish-13	BRAMS-205.1 (c)	1,323 ± 25	
		Seed	GU-18290	-	Fish-14	BRAMS-2052 (c)	1,365 ± 25	
		Cattle	SUERC-22896	970 ± 30	Fish-15	BRAMS-2053 (c)	1,308 ± 25	
		Cattle	SUERC-22897	975 ± 25	Limpet-8	BRAMS-2041*	1,380 ± 24	
		Sheep	OxA-15420	903 ± 27	Limpet-9	BRAMS-2042.1 (a)	1,263 ± 25	
		Cattle	OxA-15522	985 ± 26		BRAMS-2042.2 (a)	1,236 ± 24	
					Limpet-10	BRAMS-2043(b)*	1,575 ± 24	
					Limpet-11	BRAMS-2044 (b)*	1,593 ± 25	
					Winkle-1	BRAMS-2045 (a)	1,257 ± 24	
					Winkle-2	BRAMS-2046 (b)*	1,614 ± 25	
					Winkle-3	BRAMS-2047 (a)	1,243 ± 24	
					Winkle-4	BRAMS-2048 (b)*	1,613 ± 25	
	AD	Cattle	BRAMS-1716	956 ± 25	Fish-4	BRAMS-1720	1,167 ± 25	-84 ± 34
					Fish-5	BRAMS-1721	1,261 ± 25	
					Fish-6	BRAMS-1722	1,257 ± 25	
LN	AG	Sheep	BRAMS-1714	930 ± 25	Fish-7	BRAMS-1723.1*	1,268 ± 25	-
						BRAMS-1723.2*	1,237 ± 25	
Overall site					Fish-8	BRAMS-1724*	1,183 ± 25	-65 ± 46

298 The MREs calculated from Groups (a) and (c) gave statistically indistinguishable <sup>14</sup>C  
299 determinations and are in good agreement with ΔR values calculated for the other contexts.  
300 Only shells from Group (b) were excluded from the overall ΔR determination due to  
301 uncertainty in the security of the context in light of its high ΔR value. The remaining 56 ΔRs  
302 failed the statistical identity test, therefore layer BAG (ΔR = -214 ± 26), showing the  
303 lowest ΔR and the pairs BN-F-14/SUERC-2684, BN-F-14/OxA15420 which showed the  
304 highest ΔR values were excluded from the calculation. With the removal of the organisms  
305 and terrestrial/marine pairs identified as outliers the remaining 53 ΔR values are statistically  
306 identical (T' = 69.3, T'(5 %) = 71.0, v = 53) and average to -65 ± 46. These data suggest  
307 there is no significant difference in the reservoir effect from the LIA2 to LN period at the site.  
308 This ΔR value of -65 ± 46 is also consistent with the previously reported ΔR values for the  
309 North Atlantic, including the (Inner, and Outer) Hebrides Island of -47 ± 52 for the period  
310 3500 BC-1450 cal AD (Reimer, *et al.* 2002; Ascough, *et al.* 2004, 2005, 2006, 2007, 2009,  
311 2017; Russel, *et al.* 2010, 2015; see supplementary material S4).

312

### 313 Radiocarbon dating of pottery vessels

314 C<sub>16:0</sub> and C<sub>18:0</sub> FAs from 21 sherds were dated, of which six were dated in duplicate. This  
315 includes potsherds from all phases present at Bornais, both with and without aquatic  
316 biomarkers present. Of these, 17 sherds successfully passed the internal quality control  
317 criterion, whereby the radiocarbon dates of the C<sub>16:0</sub> and C<sub>18:0</sub> FAs must agree within 95 %  
318 confidence. Three failed, and seven did not yield sufficient C for both FAs to be dated

319 independently. These last ten sherds were therefore not further considered as no internal  
 320 control on the C<sub>16:0</sub> and C<sub>18:0</sub> FAs was present to ensure the security of the dates  
 321 (Supplementary materials S5).

322 Two of the pottery vessels dated in duplicate failed the internal quality control the first time,  
 323 but either passed it the second time (BN-35) or yielded insufficient C for two targets (BN-  
 324 101). Three pot dates that were duplicated gave indistinguishable dates for both extracts. The  
 325 duplicate analysis of potsherd BN-74 produced statistically non-identical results between the  
 326 two extractions. The CSRA dates successfully passed the internal criterion for both  
 327 extractions and as the C<sub>16:0</sub> and C<sub>18:0</sub> dates are essentially independent, it is unlikely that both  
 328 FAs in one extraction could be contaminated to the same degree (giving rise to identical, but  
 329 inaccurate, dates; Casanova, *et al.* 2018). This difference could, therefore, reflect an  
 330 inhomogeneous partitioning of the marine and terrestrial products in the same potsherd (due  
 331 to different filling levels during cooking for example), and could potentially be monitored  
 332 and corrected for in the future by recording  $\delta^{13}\text{C}$  values on the two different TLEs (not  
 333 performed in this case). Table 2 reports the combined measurements on the potsherds which  
 334 passed the internal control.

335 These results suggest that the internal quality control is valid in this case of mixed  
 336 marine/terrestrial resources and can be used as evidence for the reliability of the CSRA  
 337 measurements. The error introduced by mixing the FAs of different abundances is likely  
 338 below the AMS error, justifying the hold of the internal criteria.

339

340 Table 2: Summary of radiocarbon dated potsherds from Bornais, including the presence of aquatic  
 341 biomarkers,  $\delta^{13}\text{C}$  values of individual FAs, combined radiocarbon determinations of C<sub>16:0</sub> and C<sub>18:0</sub>  
 342 FAs (which passed the internal criterion) and the percentage of marine fat/oil within the TLEs. The  
 343 %<sub>marine</sub> were calculated using reference <sup>14</sup>C measurements on marine-terrestrial samples (%<sub>marine</sub> <sup>14</sup>C),  
 344 using a linear mixing with  $\delta^{13}\text{C}$  values on ruminant adipose products (%<sub>marine</sub>  $\delta^{13}\text{C}_{\text{adipose}}$ ) and  $\delta^{13}\text{C}$   
 345 values on ruminant dairy products (%<sub>marine</sub>  $\delta^{13}\text{C}_{\text{milk}}$ ) as endmembers and finally using  $\delta^{13}\text{C}$  values  
 346 implemented in FRUITS (%<sub>marine</sub>  $\delta^{13}\text{C}_{\text{FRUITS}}$ ). \*refers to the preferred endmembers for the  
 347 terrestrial fats based on the  $\Delta^{13}\text{C}$  value (i.e. milk if  $\Delta^{13}\text{C} < -3.1\text{‰}$ , ruminant adipose otherwise) and  
 348 used for the %<sub>marine</sub> calculation using FRUITS (v2.1; here the mean and standard deviation are  
 349 presented and the full probability distribution are in supplementary material S6).

Phase	Layer	Pot#	Aquatic biomarkers	$\delta^{13}\text{C}_{16:0}$ (‰)	$\delta^{13}\text{C}_{18:0}$ (‰)	Laboratory#	Age $\pm 1 \sigma$ (BP)	% <sub>marine</sub> <sup>14</sup> C	% <sub>marine</sub> $\delta^{13}\text{C}_{\text{adipose}}$	% <sub>marine</sub> $\delta^{13}\text{C}_{\text{milk}}$	% <sub>marine</sub> $\delta^{13}\text{C}_{\text{FRUITS}}$
LIA 2	BAC	BN89 (1)	-	-26.8	-29.4	BRAMS-1549.1	1,368 $\pm$ 25	27 $\pm$ 12	30 $\pm$ 22*	37 $\pm$ 22	35 $\pm$ 10
		BN89 (2)	-	-	-	BRAMS-1549.2	1,365 $\pm$ 25	26 $\pm$ 12	-	-	-
		BN74 (1)	APAAs	-26.6	-29.4	BRAMS-1551.1	1,383 $\pm$ 30	31 $\pm$ 12	31 $\pm$ 23*	39 $\pm$ 16	36 $\pm$ 10
		BN74 (2)	-	-	-	BRAMS-1551.2	1,286 $\pm$ 25	4 $\pm$ 11	-	-	-
	BAF	BN77	-	-26.9	-29.9	BRAMS-1605	1,370 $\pm$ 24	-	26 $\pm$ 23*	34 $\pm$ 14	32 $\pm$ 10
		BN87	APAAs	-26.8	-30.4	BRAMS-1604	1,304 $\pm$ 24	-	21 $\pm$ 25	33 $\pm$ 15*	32 $\pm$ 12
		BAG	BN88 (1)	APAAs,	-24.2	-25.4	BRAMS-1548	1,757 $\pm$ 25	-	69 $\pm$ 31*	73 $\pm$ 27

		BN88 (2)	DHYAs			BRAMS-1548	1,762 ± 25	-			
EN	BBD	BN35	APAAs, DHYAs	-25.4	-31.4	BRAMS-1552	1,156 ± 27	-	12 ± 12	31 ± 31*	33 ± 10
		BN105 (1)	-	-26.2	-31.7	BRAMS-1547.1	1,268 ± 25	-	8 ± 36	28 ± 24*	27 ± 10
		BN105 (2)	-			BRAMS-1547.2	1,327 ± 27	-			
	BBA	BN110	APAAs	-25.4	-25.9	BRAMS-1608	1,326 ± 25	-	64 ± 28*	63 ± 24	63 ± 13
MN	BCA	BN115	-	-27.2	-31.0	BRAMS-1609	987 ± 24	-	15 ± 24	28 ± 13*	27 ± 9
	BCC	BN160	-	-26.4	-31.0	BRAMS-2066	1,201 ± 25	74 ± 16	16 ± 31	32 ± 20*	29 ± 10
		BN165	-	-26.3	-30.1	BRAMS-2063	1,060 ± 25	32 ± 14	25 ± 28	38 ± 18*	36 ± 10
		BN174	APAAs	-25.9	-28.5	BRAMS-2062	1,115 ± 26	49 ± 15	40 ± 26*	48 ± 20	46 ± 12
LN	AG	BN36	APAAs, DHYAs	-27.3	-31.8	BRAMS-1607	786 ± 25	-	7 ± 26	24 ± 14*	21 ± 8

350

351 The four sherds from the BAC and BAF contexts of phase LIA2, the three from the EN phase  
352 and the three from the BCC context of the MN phase were shown to have radiocarbon ages  
353 between the age of the terrestrial organisms and their contemporaneous marine analogues  
354 (Table 1 and 2). These include the five sherds (BN-89, BN-77, BN-105, BN160 and BN-165)  
355 which did not exhibit aquatic biomarkers. These dates suggest, therefore, mixing of terrestrial  
356 and marine resources in all the sherds.

357 The sherd BN-88 from the BAG context LIA2 phase exhibited not only the most enriched  
358  $\delta^{13}\text{C}$  values but also the oldest age obtained in this investigation. This date is older than the  
359 marine reference fish bone from this context and, indeed the reference fish bones from other  
360 LIA 2 contexts. The second dating of the potsherd confirmed the accuracy of the compound-  
361 specific  $^{14}\text{C}$  measurement, suggesting the FAs likely derived from a pure marine fat/oil  
362 residue and that the MRE (based on only one pair) was underestimated in this case, unless the  
363 potsherd was residual and corresponds to the LIA1 phase, although no aquatic biomarkers  
364 were detected in potsherds extracts from this particular phase.

365 The potsherd BN-115 ( $987 \pm 24$  BP) from context BCA (not dated) of the MN phase, which  
366 lacked aquatic biomarkers exhibited an age consistent with the MN phase, and thus is likely  
367 to be entirely composed of terrestrial animal fats.

368 For the LN phase, the FA date on the pot BN-36 ( $786 \pm 25$  BP) is younger than that of the  
369 terrestrial organisms (BN-MB-7:  $930 \pm 25$  BP). Based on the  $\delta^{13}\text{C}$  values, the sherd plots  
370 close to the reference dairy fat ellipses despite containing aquatic biomarkers. This result is  
371 surprising and suggests that the dating of this phase, based on only one terrestrial organism  
372 could be erroneous. Younger ages from other LN contexts from Bornais were obtained in a  
373 range of 900 to 650 radiocarbon years BP (uncalibrated), which would support this  
374 hypothesis (Marshall *et al.* 2016, forthcoming).

375 These measurements clearly confirm that lipid dates can be affected by the marine reservoir  
376 effect and such dates will therefore require calibration using relevant  $\Delta R$  values and  
377 proportionately mixed terrestrial/marine curves. The mixing of marine and terrestrial  
378 products influences the determined  $\delta^{13}\text{C}$  values and radiocarbon dates of FAs and this does  
379 not appear to have an adverse effect on the internal quality control criteria. Interestingly,  
380 MREs are evident in TLEs from potsherds lacking detectable aquatic biomarkers. The results  
381 therefore suggest that  $^{14}\text{C}$  dates could be used to detect a (low-)level of marine organism  
382 processing in pots where aquatic biomarkers are undetectable. This would be especially  
383 relevant for sites where potential exists for processing of non-ruminant products or where  
384 aridity effects are possible, shifting the  $\delta^{13}\text{C}$  values away from the ruminant products ellipses.

### 385 **Correction of the MRE and calibration**

386 The  $\%_{\text{marine}}$  in the lipid residues was quantified for the sherds which passed the internal  
387 quality control criterion (Table 2; Supplementary material S5, S6). To ensure a fair  
388 evaluation of the use of FA  $\delta^{13}\text{C}$  values for determination of the degree of marine/terrestrial  
389 product mixing, only potsherds from contexts which were securely dated using more than one  
390 marine/terrestrial organism were used for validating the correction and calibration (BAC and  
391 BCC). The validity of using  $\delta^{13}\text{C}$  values of FAs for the quantification of marine-derived C  
392 was evaluated by comparison with reference values obtained by  $^{14}\text{C}$  dates.

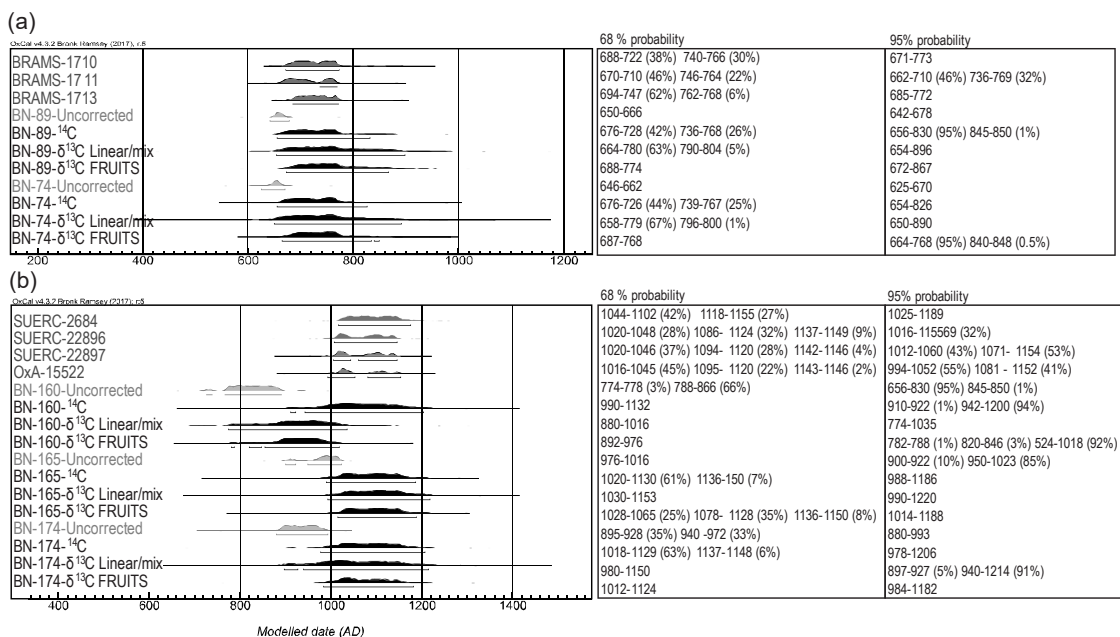
393 Overall, no significant differences in  $\%_{\text{marine}}$  were noted in the use of  $\delta^{13}\text{C}$  values from  
394 ruminant adipose or milk fats as terrestrial end members in a simple linear mixing model  
395 (Table 2, Figure 2f). Therefore, only the one most representative of the terrestrial  
396 endmembers was used for MRE corrections (i.e. milk if  $\Delta^{13}\text{C} < -3.1\text{‰}$ , ruminant adipose if  
397  $\Delta^{13}\text{C} > -3.1\text{‰}$ ).

398 The range of calibrated terrestrial dates on mammals for BRAMS-1710, BRAMS-1711,  
399 BRAMS-1713, in the LIA 2 phase, BAC context, were 672 - 773 cal AD, 662 - 769 cal AD  
400 and 685 - 772 cal AD, respectively (95% probability, Figure 3a). The  $\%_{\text{marine}}$  within the FAs  
401 in pot BN-89 was determined to be  $27 \pm 12 \%$  using  $^{14}\text{C}$  dates,  $30 \pm 22 \%$  using  $\delta^{13}\text{C}$  values  
402 of adipose endmembers in the simple linear mixing and  $35 \pm 10 \%$  (mean and standard  
403 deviation) when implemented in FRUITS (Table 2). All these estimates are statistically



404 indistinguishable and the calibrated ages after MRE correction agrees with the reference age  
 405 of the terrestrial organisms (Figure 3a).

406 Turning to potsherd BN-74, the first extract yielded estimates of  $31 \pm 12$  % marine fat/oil  
 407 using  $^{14}\text{C}$  dates, and  $31 \pm 23$  % and  $36 \pm 10$  % using  $\delta^{13}\text{C}$  values of adipose FAs endmembers  
 408 on the mixing lines and FRUITS, respectively. The calibrated age from pot BN-74 (1<sup>st</sup>  
 409 extract) agrees with the age of terrestrial organisms (Figure 3a). Nonetheless, the 2<sup>nd</sup> extract  
 410 of the pot BN-74, which yielded results statistically different to the 1<sup>st</sup> extract, showed a  
 411 %<sub>marine</sub> of  $4 \pm 11$  % using  $^{14}\text{C}$  as end-members, suggesting an underestimation of the  
 412 proportion of marine products in the TLE based on the  $\delta^{13}\text{C}$  values in this case. As mentioned  
 413 previously, this potsherd is likely affected by a differential deposition of the marine fats in  
 414 certain areas of the vessel, implying that determination of  $\delta^{13}\text{C}$  values and  $^{14}\text{C}$  dates on the  
 415 same TLE is required for a satisfactory quantification of the %<sub>marine</sub> using  $\delta^{13}\text{C}$  values.



416

417 **Figure 3**

418 The range of calibrated terrestrial dates (excluding outliers, Table 1) varies from 993 – 1,052  
 419 cal AD (55 % probability) and 1,081 – 1,152 cal AD (OxA-15522; 41% probability) to 1,039  
 420 – 1,206 cal AD (OxA-1540, 95 % probability) for the MN phase, BCC context (Figure 3b).

421 The results for potsherds BN-165 using  $\delta^{13}\text{C}$  values of milk fatty acids and BN174 using  $\delta^{13}\text{C}$   
 422 values of adipose fatty acids showed, similarly to BN-89 and BN-74 (1<sup>st</sup> extraction), a good

423 agreement with the age of reference terrestrial animals using the different methods (Figure  
424 3b).

425 The  $\%_{marine}$  in the pot BN-160 is, however, estimated to be  $74 \pm 16 \%$  with the use of  $^{14}\text{C}$   
426 dates,  $26 \pm 34 \%$  and  $29 \pm 10 \%$  with the use of  $\delta^{13}\text{C}$  values milk endmembers in the linear  
427 mixing curve and FRUITS, respectively (Table 2). These results are not identical within a  $1 \sigma$   
428 error but are within  $2 \sigma$ . The potsherd BN-160 was calibrated to  $908 - 1,212$  cal AD by  
429 radiocarbon estimates,  $730 - 1,044$  cal AD and to  $780 - 1,016$  cal AD (95 % probability for  
430 all) by dairy endmember in linear mixing and FRUITS, respectively. The end of the last two  
431 distributions overlap only at the start of the calibration of the reference terrestrial organisms  
432 (Figure 3b). It should be noted that for potsherd BN-160, the  $^{14}\text{C}$  dates suggest that marine  
433 fat/oil are dominate in the TLE whereas the  $\delta^{13}\text{C}$  values suggest a dominance of dairy  
434 products. Unless the CSRA date is inaccurate, this implies that this potsherd, like BN-74,  
435 could be affected by a differential partitioning of the marine products and that  $\delta^{13}\text{C}$  values  
436 recorded on another TLE are not representative of the TLE used for  $^{14}\text{C}$  dating.

437 Overall, MRE corrections of lipid residues, using  $\delta^{13}\text{C}$  calculations using the simple linear  
438 mixing model showed a wider probability distribution than those obtained using radiocarbon  
439 dates and  $\delta^{13}\text{C}$  values used in the software FRUITS. However, the calibrated range of the  
440 corrected CSRA determinations on pot lipids using both methods clearly overlaps the  
441 calibrated range of the reference terrestrial organisms. The precision of the calibrated ages  
442 depends almost entirely on the uncertainties associated with the calculated  $\%_{marine}$ , as  
443 illustrated with reduced errors obtained using FRUITS software instead of the simple linear  
444 mixing curve. The results demonstrate no significant difference in the use of ruminant  
445 adipose or dairy  $\delta^{13}\text{C}$  values as end members for the quantification of the  $\%_{marine}$ . In practice,  
446 however, one should be chosen over the other based on the  $\Delta^{13}\text{C}$  values to ensure that the  
447 terrestrial endmember is representative of the animal products processed in the vessels at the  
448 time (i.e. dairy if  $\Delta^{13}\text{C} < -3.1\%$  or adipose if  $\Delta^{13}\text{C} > -3.1\%$ ). On the other hand, the  $^{14}\text{C}$  dates  
449 provide an accurate estimate of the  $\%_{marine}$  present in the FAs and could be used for  
450 quantification of marine products in TLEs instead of a calendar age. The  $\%_{marine}$  in potsherds  
451 BN-74 (2<sup>nd</sup> extraction) and BN-160 were underestimated, leading to inappropriate  
452 corrections. However, this could be accounted for in the future if the  $^{14}\text{C}$  measurements and  
453  $\delta^{13}\text{C}$  values are recorded on the same lipid extract to avoid potential inconsistencies

454 associated with inhomogeneous deposition of the lipids within vessels and use more reliable  
455  $\delta^{13}\text{C}$  values for the quantification of marine products.

456 One limitation of the  $\delta^{13}\text{C}$  approach is the estimation of  $\%_{\text{marine}}$  due to the wide range of  
457 reference values (from ca. -26 ‰ to -20 ‰) observed in modern marine organisms. The  
458 reference ellipses commonly plotted comprise only 68 % of the reference values (i.e. 1- $\sigma$ ).  
459 The average values used to generate an endmember here are not centred in the ellipses  
460 (Figure 2f). Therefore, potsherds with individual FAs  $\delta^{13}\text{C}$  values plotting at the edge of the  
461 reference marine ellipse can be purely marine but, the  $\%_{\text{marine}}$  deposited in the sherd can be  
462 underestimated using the linear mixing curves (Table 2, Figure 2f). This phenomenon is  
463 illustrated in the case of potsherd BN-88 which is likely to contain predominantly marine fats  
464 based on the CSRA dates. The  $\delta^{13}\text{C}$  values plotted just outside the marine reference ellipse  
465 and marine-derived C was quantified to be  $69 \pm 31$  % with adipose endmember, and the  
466  $\%_{\text{marine}}$  appeared to be underestimated in this case. Potsherd BN-110 could also contain a  
467 dominance of marine products based on fatty acid  $\delta^{13}\text{C}$  values (i.e.  $63 \pm 28$  % with adipose  
468 used as endmembers; Table 2, Figure 2f). This suggests that the linear mixing does not  
469 account particularly well for the dominance of marine products at the boundaries. Therefore,  
470 the use of the mean and standard deviation for the reported  $\%_{\text{marine}}$  products would lead to  
471 some underestimation when applying MRE corrections in the case of dominance of marine  
472 products. This would be overcome using the full probability distribution calculated in  
473 FRUITS as prior information on the percentage marine.

474 We suggest that a linear mixing curve can give valid corrections if marine products are not  
475 dominant in the TLE, however, the FRUITS software would deal more adequately with the  
476 boundaries (if probability distribution are used) and should be used preferably to access the  
477  $\%_{\text{marine}}$  in the TLEs.

478

## 479 **CONCLUSION**

480 The processing of mixed terrestrial/marine fats in pottery vessels at the site of Bornais was  
481 revealed through lipid biomarker and CSRA analyses. CSRA and comparison with the  
482 radiocarbon dates of associated marine and terrestrial samples also enabled the detection of  
483 marine product processing in cases where no aquatic biomarkers were detected. We therefore  
484 suggest that in such circumstances,  $^{14}\text{C}$  measurements could be used as a tracer for the

485 detection and quantification of marine products processing in pots. Compound-specific dates  
486 from potsherds from Bornais were successfully subjected to MRE correction, and assessed  
487 against independent ages determined for contemporaneous terrestrial organisms using:

- 488 (i) An appropriate  $\Delta R$  ( $-65 \pm 45$ ) for the site and time period.
- 489 (ii) An estimate of the proportion of marine resource processed in the pots calculated  
490 using  $\delta^{13}\text{C}$  values on individual pot FAs and from a modern reference database  
491 (linear mixing or implementation in FRUITS).
- 492 (iii) A mixed calibration approach in OxCal software.

493 These corrected ages agreed well with the calibrated age of terrestrial samples which  
494 confirmed the efficacy of using FA  $\delta^{13}\text{C}$  values to estimate the  $\%_{\text{marine}}$ , meaning that an  
495 approach similar to that commonly adopted for bone collagen can be used to correct for MRE  
496 present in lipids. For future MRE corrections and calibration on lipids dates, we recommend:

- 497 (i) Calculating a  $\Delta R$  for the site using a paired terrestrial/marine sample approach or  
498 using a previously published  $\Delta R$  relevant for the spatiotemporal area.
- 499 (ii) Recording  $\delta^{13}\text{C}_{16:0}$  and  $\delta^{13}\text{C}_{18:0}$  values from the same TLE as that used for  $^{14}\text{C}$   
500 dating to determine the  $\%_{\text{marine}}$ , avoiding the negative impact of potential  
501 inhomogeneity of lipid distribution in vessels.
- 502  
503 (iii) Using mixing model endmembers calculated from modern reference values valid  
504 for the location of interest, by using either those of the database for UK animals  
505 (excluding the species not present at the site; Copley, *et al.* 2003; Cramp and  
506 Evershed 2014) or  $\delta^{13}\text{C}$  values recorded from reference animals, representative of  
507 other locations and environmental conditions (e.g. arid environments, Dunne *et al.*  
508 2012).
- 509 (iv) Using endmembers from dairy reference fats in the case of potsherds with  $\Delta^{13}\text{C} < -$   
510  $3.1$  ‰ or using endmembers from the reference ruminant adipose fats values for  
511 potsherds with  $\Delta^{13}\text{C} > - 3.1$  ‰, to determine  $\%_{\text{marine}}$  in the TLEs.

- 512 (v) Employ FRUITS or other Bayesian approaches (if available) to quantify %*marine* in  
513 the TLEs using a probability density function.
- 514 (vi) Correct CSRA dates for the MRE using mixed atmospheric and marine calibration  
515 curves (e.g. in OxCal).

516

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524

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663

664 **Figure captions:**

665 Figure 1: Flowchart showing the method used to assess the validity of the  $\delta^{13}\text{C}$  values method for the  
666 estimation of the  $\%_{\text{marine}}$  products and correction of the CSRA dates on the FAs.

667

668 Figure 2: Partial gas chromatogram of the TLE (a), and GC/MS SIM mass chromatograms showing  
669 detection of DHYAs (b) and APAAs (c), for potsherd BN-173. Scatter plots of  $\delta^{13}\text{C}_{16:0}$  plotted against  
670  $\delta^{13}\text{C}_{18:0}$  from lipid residues characteristic of animal fats at Bornais for all the TLEs (Cramp *et al.*,  
671 2014b, forthcoming and this study), (d) for the 22 potsherd extracts selected for  $^{14}\text{C}$  dating by CSRA  
672 and position of the average reference values (crosses) (e), and the theoretical mixing lines of  
673 terrestrial and marine end-members with the approximate percentage of marine fat/oil marked on the  
674 lines (f). Stars denote the detection of aquatic biomarkers. Shaded areas indicate the reference ellipses  
675 for  $\delta^{13}\text{C}$  values on modern animals and the crosses are the values used as endmembers. The dashed  
676 lines correspond to the areas where lipid residues are hypothesised to be affected to varying degrees  
677 by the MRE.

678

679 Figure 3: Corrections and calibration for potsherds of the (a) LIA phase, (b) MN phase in OxCal v4.3  
680 against the IntCal13 calibration curve (Bronk Ramsey 2009, Reimer *et al.* 2013). The distributions  
681 plotted in dark grey correspond to the reference age of terrestrial animal bones, in light grey the  
682 uncorrected determinations on pot lipids and in black the corrected determinations on pot lipids using  
683 either the  $^{14}\text{C}$  or  $\delta^{13}\text{C}$  methods using adipose or milk as endmembers.

684

## Supporting Materials

### **Compound-specific radiocarbon, stable carbon isotope and biomarker analysis of mixed marine/terrestrial lipids preserved in archaeological pottery vessels**

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### S1- Site location and stratigraphic information of Mound 2

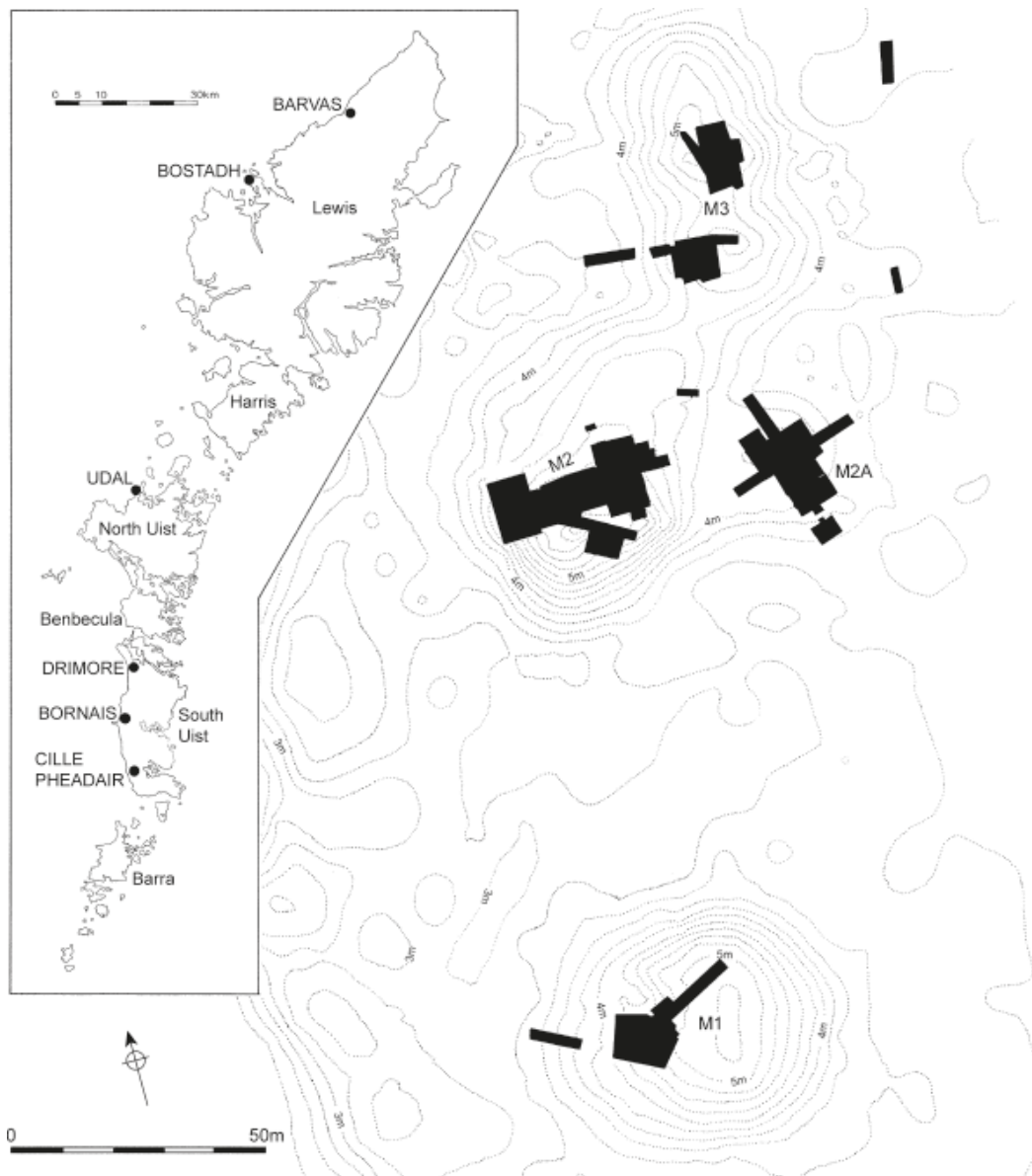


Figure S1: Location of the site of Bornais and the 4 mounds excavated. From Sharples *et al.* (2016), Fig.17.1.

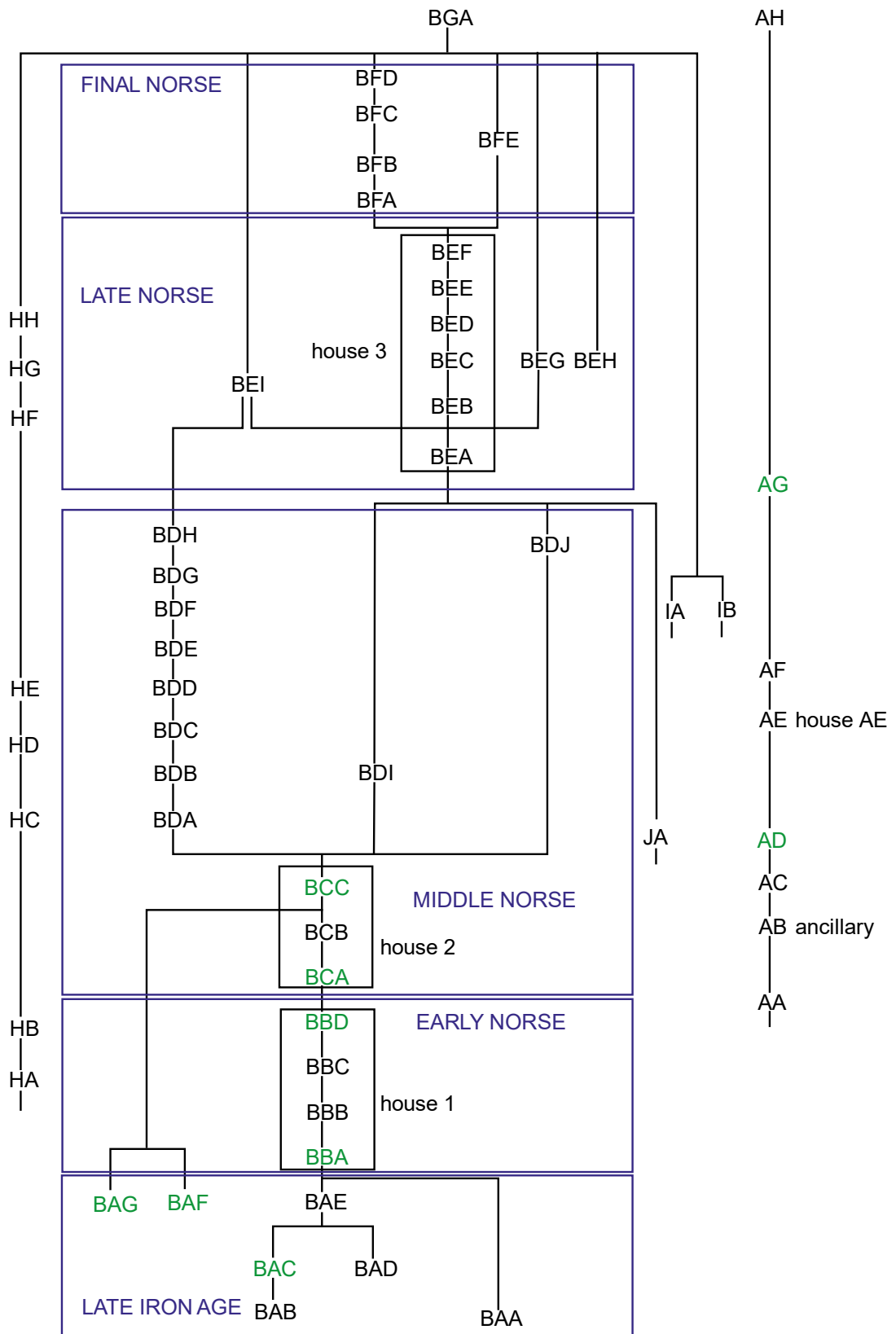


Figure S2: Schematic diagram showing the stratigraphic information for the Mound 2. Green contexts correspond to the ones studied in this paper.

## S2- Detailed method for lipid residue analyses

To remove surface contaminants, a small part of the potsherd (~ 2 - 7 g) was cleaned with a modelling drill then sampled using hammer and chisel and ground to fine powder using mortar and pestle. Approximately 1 to 2 g of ground potsherd were weighed into a clean culture tube (I) with stopper and 20  $\mu\text{L}$  of internal standard (IS; *n*-tetratriacontane) at 1  $\text{mg}\cdot\text{mL}^{-1}$  was added. The lipids were extracted using a solution of  $\text{H}_2\text{SO}_4/\text{MeOH}$  (4% v/v, 5 mL, 70 °C, 1 h.). The supernatant of culture tube I was then centrifuged (2500 rpm, 10 min) and transferred to a clean culture tube (II) before adding double distilled water (2 mL). *n*-hexane was added (2 x 3 mL) to culture tube I and the supernatant transferred to culture tube II. Following this, 2 x 2 mL *n*-hexane was added directly to the  $\text{H}_2\text{SO}_4/\text{MeOH}$  solution in culture tube II and whirlimixed to extract the remaining residues, then transferred to the 3.5 mL vials and blown down until a full vial of *n*-hexane remained. A procedural blank was prepared and analysed alongside every batch of archaeological materials to assess whether contamination was introduced during the protocol. Before analysis, an aliquot of the total lipid extract (TLE; 1/4) was derivatised by the addition of BSTFA (*N,O*-bis(trimethylsilyl)trifluoroacetamide; 20  $\mu\text{L}$ , 70 °C, 1 h). Excess BSTFA was blown down at 40 °C under a gentle stream of nitrogen, and an appropriate amount of *n*-hexane was added, prior to analysis with GC, GC-MS and GC-C-IRMS (Correa-Ascencio and Evershed 2014).

GC analysis of TLEs (Section 2.3.2.1) for quantification was performed on a Hewlett Packard 5890 series II gas chromatograph or an Agilent Technologies 7890A GC. Helium was used as carrier gas at constant flow (2  $\text{mL}\cdot\text{min}^{-1}$ ), and a flame ionisation detector (FID) used to monitor column effluent. Lipids extracts (1  $\mu\text{L}$ ) were injected into a non-polar fused silica capillary column (50 m x 0.32 mm i.d., DB1 stationary phase (100 % dimethylpolysiloxane), 0.17  $\mu\text{m}$  film thickness, Agilent technologies). The oven temperature program started with an isothermal hold at 50 °C for 2 min, then the temperature was increased at 10  $^\circ\text{C}\cdot\text{min}^{-1}$  to 300 °C and held for 10 min (Evershed *et al.* 1990).

GC-MS analysis of TLEs for molecular identification was performed on a Finnigan Trace MS quadrupole instrument coupled to a Trace GC, or on a Thermo Scientific ISQ LT single quadrupole GC-MS coupled to a Trace 1300, with manual or auto-sampling injections. The lipid extracts (1  $\mu\text{L}$ ) were introduced into a non-polar fused silica capillary column (50 m x 0.32 mm i.d., DB1 stationary phase, 0.17  $\mu\text{m}$  film thickness, Agilent Technologies). For TLEs analysis the oven temperature program started with an isothermal hold at 50 °C during 2 min, then the temperature increased at 10  $^\circ\text{C}\cdot\text{min}^{-1}$  to 300 °C and held for 10 min. The MS used electron ionization (EI) mode operating at 70 eV with a GC interface temperature of 300 °C and a source temperature of 200 °C. Acquisition used the total ion current (TIC) mode over the range *m/z* 50-650 Daltons at 8.3 scans. $\cdot\text{s}^{-1}$  (Evershed *et al.* 1990). Screening for di-hydroxy fatty acid methyl esters (DHYAs, aquatic biomarkers) used selected ion monitoring

(SIM) mode, monitoring  $m/z$  159, 187, 215, 243, 259, 287, 315, 443, 459, 471, 487, 499 and 515 (for -COOMe derivatives instead of -COOTMS as published; Cramp and Evershed 2014).

In order to determine the presence of other aquatic biomarkers (APAAs and isoprenoid acids), TLEs were run on a polar column (60 m x 0.32 mm i.d., VF-23ms stationary phase (polydimethylsiloxane highly substituted with cyanopropyl groups), 0.15  $\mu\text{m}$  film thickness, Agilent Technologies). The temperature program started with an isothermal hold at 70 °C for 2 min, followed by a ramp at 10 °C.min<sup>-1</sup> to 220 °C, then a ramp at 4 °C.min<sup>-1</sup> to 300 °C and finally an isothermal hold for 10 min. Full scan mode  $m/z$  50-650 and SIM mode, screening for the masses  $m/z$  105, 262, 290, 318 and 346, were performed for the detection of APAAs (Cramp and Evershed 2014).

The GC-C-IRMS analyses on C<sub>16:0</sub> and C<sub>18:0</sub> FAs (for identification of the source of animal fats) was performed on an Agilent Technologies 7890A, coupled via an IsoPrime GC5 combustion interface (CuO and silver reactor, 850 °C) to an IsoPrime 100 mass spectrometer. The FAME extracts (1  $\mu\text{L}$ ) were injected into a non-polar column (50 m x 0.32 mm i.d., DB1 stationary phase, 0.17  $\mu\text{m}$  film thickness, Agilent technologies). The GC oven temperature was held for 2 min at 40 °C and increased to 300 °C at 10 °C.min<sup>-1</sup> and held for 10 min. The MS used EI at 70 eV and had three Faraday cups collecting for the masses  $m/z$  44, 45 and 46. Data were acquired and processed by the IonVantage software (Copley *et al.* 2003).

### S3- Results of lipid residue analysis

Table S1: Results of lipid residue analysis of potsherds from the site of Bornais Mound 2, House 2. P corresponds to the Phytanic acid and TMTD to the 4,8,12-trimethyltridecanoic acid. Compounds in brackets corresponds to trace amounts (not clearly identified).

Sherd #	Block	Code	C ( $\mu\text{g}\cdot\text{g}^{-1}$ )	FAs	DHYAs	APAAs	IFAs	$\delta^{13}\text{C}_{16:0}$ (‰)	$\delta^{13}\text{C}_{18:0}$ (‰)	$\Delta^{13}\text{C}$ (‰)	Assignment (before CSRA dating)
BN-132	BCA	1259/3790	849	C <sub>14</sub> -C <sub>18</sub>	-	-	(TMTD), P	-25.8	-31.0	-5.2	Mixture dairy, non-ruminant fats
BN-133	BCA	1280/9448/3/3887	137	C <sub>14</sub> -C <sub>18</sub>	C <sub>18</sub>	-	-	-	-	-	<i>nd</i>
BN-134	BCB	6/8656	190	C <sub>14</sub> -C <sub>22</sub>	-	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	-	-	-	<i>nd</i>
BN-135	BCB	1089/8654	1444	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	-	(TMTD), P	-26.3	-26.6	-0.2	Mixture ruminant adipose, marine fats
BN-136	BCB/C	1074/2/3529	207	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub>	-	-26.7	-32.9	-6.2	Mixture dairy, marine fats
BN-137	BCC	182/1/8659	676	C <sub>14</sub> -C <sub>22</sub>	-	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	-24.9	-28.4	-3.5	Mixture dairy, marine fats
BN-138	BCC	528/2/199	19	C <sub>14</sub> -C <sub>22</sub>	-	-	-	-	-	-	<i>nd</i>
BN-139	BCC	549/2/264	3135	C <sub>14</sub> -C <sub>20</sub>	C <sub>18</sub> , C <sub>22</sub>	C <sub>18</sub>	-	-26.6	-29.3	-2.7	Ruminant adipose fats
BN-140	BCC	550/5/2458	216	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	-22.1	-25.2	-3.1	Marine fats
BN-141	BCC	550/5/2458	28	C <sub>16</sub> -C <sub>22</sub>	(C <sub>18</sub> )	-	-	-26.6	-32.5	-5.9	Mixture dairy, non-ruminant fats
BN-142	BCC	557/5/2341	2220	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> )	(TMTD), P	-24.9	-27.9	-3.0	-
BN-143	BCC	557/5/8660	5630	C <sub>14</sub> -C <sub>20</sub>	C <sub>18</sub>	C <sub>18</sub>	P	-25.8	-27.9	-2.2	Mixture ruminant adipose, marine fats
BN-144	BCC	558/5/8670	2483	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub>	(TMTD), P	-25.6	-28.9	-3.3	Mixture dairy, marine fats
BN-145	BCC	565/8655	1425	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub>	-	P	-27.0	-31.3	-4.3	Mixture dairy, non-ruminant fats
BN-146	BCC	921/2/992	1349	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub>	C <sub>18</sub>	P	-25.6	-30.2	-4.6	Mixture dairy, non-ruminant fats
BN-147	BCC	1008/3063	56	C <sub>16</sub> -C <sub>22</sub>	(C <sub>18</sub> )	-	-	-	-	-	<i>nd</i>
BN-148	BCC	1008/9452/8657	60	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	TMTD, P	-25.3	-29.2	-4.0	Mixture dairy, marine fats
BN-149	BCC	1010/9685/2/3279	3984	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub>	(TMTD), P	-26.3	-30.4	-4.1	Mixture dairy, marine fats
BN-150	BCC	1010/2/3266	1460	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	-25.2	-29.0	-3.8	Mixture dairy, marine fats
BN-151	BCC	1049/9809/8661	2141	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	C <sub>18</sub>	P	-26.9	-29.6	-2.7	Mixture ruminant, non-ruminant adipose fats
BN-152	BCC	1057/9895/8656	0	-	-	-	-	-	-	-	-
BN-153	BCC	1057/9894/3522	1409	C <sub>14</sub> -C <sub>22</sub>	(C <sub>18</sub> )	C <sub>18</sub>	-	-26.5	-29.8	-3.3	Mixture dairy, non-ruminant fats
BN-154	BCC	1057/9893/3461	1153	C <sub>14</sub> -C <sub>20</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	TMTD, P	-27.0	-30.1	-3.0	Mixture ruminant adipose, marine fats
BN-155	BCC	1057/9894/3535	4198	C <sub>14</sub> -C <sub>22</sub>	-	-	-	-26.5	-28.6	-2.2	Mixture ruminant, non-ruminant adipose fats
BN-156	BCC	1057/9894/8666	276	C <sub>14</sub> -C <sub>18</sub>	-	-	-	-25.3	-28.9	-3.6	Mixture dairy, non-ruminant fats
BN-157	BCC	1079/2/8671	11	C <sub>16</sub> -C <sub>22</sub>	C <sub>18</sub>	-	TMTD, P	-	-	-	<i>nd</i>
BN-158	BCC	1220/9991/3662	12	-	(C <sub>18</sub> )	-	-	-	-	-	<i>nd</i>
BN-159	BCC	1220/9991/8664	1178	C <sub>14</sub> -C <sub>20</sub>	C <sub>18</sub>	C <sub>18</sub>	P	-26.4	-32.0	-5.6	Mixture dairy, non-ruminant fats
BN-160	BCC	1234/9492/8665	4559	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub>	C <sub>18</sub>	(TMTD), P	-26.4	-31.0	-4.6	Mixture dairy, non-ruminant fats
BN-161	BCC	1260/9467/3779	546	C <sub>14</sub> -C <sub>18</sub>	(C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub> )	-	-	-25.7	-31.2	-5.5	Mixture dairy, non-ruminant fats
BN-162	BCC	1260/9465/8672	1574	C <sub>16</sub> , C <sub>18</sub>	(C <sub>18</sub> )	-	-	-26.7	-29.3	-2.6	Mixture ruminant, non-ruminant adipose fats
BN-163	BCC	2192/11902/10/2192	452	C <sub>14</sub> -C <sub>18</sub>	C <sub>18</sub>	C <sub>18</sub>	-	-27.3	-31.5	-4.2	Mixture dairy, non-ruminant fats
BN-164	BCC	2225/11943/6230	129	C <sub>16</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub>	TMTD, P	-25.8	-31.6	-5.8	Mixture dairy, marine fats



Sherd #	Block	Code	C ( $\mu\text{g}\cdot\text{g}^{-1}$ )	FAs	DHYAs	APAAs	IFAs	$\delta^{13}\text{C}_{16:0}$ (‰)	$\delta^{13}\text{C}_{18:0}$ (‰)	$\Delta^{13}\text{C}$ (‰)	Assignment (before CSRA dating)
BN-165	BCC	2231/11968/14/6214	1823	C <sub>14</sub> -C <sub>18</sub>	(C <sub>18</sub> )	-	-	-26.3	-30.1	-3.7	Mixture dairy, non-ruminant fats
BN-166	BCC	2258/11287/15/8663	246	C <sub>16</sub> , C <sub>18</sub>	C <sub>18</sub>	C <sub>18</sub> , C <sub>20</sub>	TMTD, P	-25.8	-31.7	-5.9	Mixture dairy, marine fats
BN-167	BCC	2264/6314	946	C <sub>14</sub> -C <sub>22</sub>	(C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub> )	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	-23.5	-26.6	-3.1	Marine fats
BN-168	BCC	2264/14/6312	1105	C <sub>16</sub> , C <sub>18</sub>	C <sub>18</sub>	C <sub>18</sub>	-	-29.2	-31.5	-2.3	Ruminant adipose fats
BN-169	BCC	2264/14/8667	2	C <sub>18</sub> , C <sub>22</sub>	-	-	-	-	-	-	-
BN-170	BCC	2285/11278/19/8662	4231	C <sub>14</sub> -C <sub>22</sub>	-	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	nd	nd	nd	Marine fats
BN-171	BCC	2297/11318/19/8658	872	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub>	C <sub>18</sub>	P	-26.2	-32.2	-6.0	Mixture dairy, non-ruminant fats
BN-172	BCC	2613/9/6398	834	C <sub>16</sub> -C <sub>20</sub>	(C <sub>18</sub> , C <sub>20</sub> )	C <sub>18</sub>	P	-26.1	-28.5	-2.4	Mixture ruminant, non-ruminant adipose fats
BN-173	BCC	2637/11398/6500	2376	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	TMTD, P	-26.0	-30.5	-4.5	Mixture dairy, marine fats
BN-174	BCC	2657/11464/16/6523	1176	C <sub>14</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	(TMTD), P	-25.9	-28.5	-2.6	Mixture ruminant adipose, marine fats
BN-175	BCC	2673/11489/13/8669	41	C <sub>16</sub> -C <sub>22</sub>	C <sub>18</sub> , (C <sub>22</sub> )	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	(TMTD), P	-25.6	-29.8	-4.2	Mixture dairy, marine fats
BN-176	BCC	2692/12041/7300	924	C <sub>16</sub> -C <sub>20</sub>	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	C <sub>18</sub> , (C <sub>20</sub> , C <sub>22</sub> )	P	-26.4	-29.0	-2.6	Mixture ruminant adipose, marine fats
BN-177	BCC	2700/12057/6631	497	C <sub>16</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , (C <sub>20</sub> )	TMTD, P	-25.1	-30.3	-5.2	Mixture dairy, marine fats
BN-178	BCC	2715/12090/20/6637	280	C <sub>16</sub> -C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	C <sub>18</sub> , C <sub>20</sub> , (C <sub>22</sub> )	TMTD, P	-	-	-	-
BN-179	BCC	2731/1204/20/6686	1083	C <sub>14</sub> -C <sub>20</sub>	(C <sub>18</sub> )	C <sub>18</sub>	P	-26.5	-29.8	-3.2	Mixture dairy, non-ruminant fats
BN-180	BCC	545+39/2501	536	C <sub>16</sub> -C <sub>20</sub>	(C <sub>18</sub> )	C <sub>18</sub> , C <sub>20</sub> , C <sub>22</sub>	P	-24.4	-28.1	-3.7	Mixture dairy, marine fats

#### S4- Summary of published $\Delta$ determinations in the Hebrides Islands

Table S2: Summary of  $\Delta\text{R}$  values published for the Hebridian Islands

Location	Site	$\Delta\text{R} \pm 1\sigma$	Time period	Reference
Outer Hebrides Lewis and Harris	Guinnesso	-130 $\pm$ 36	1,460 - 1,630 AD	Ascough <i>et al.</i> 2017
	Bostadh	-56 $\pm$ 14	893 - 984 AD	Ascough <i>et al.</i> 2009
	Garenin	-85 $\pm$ 17	887 - 995 AD	Ascough <i>et al.</i> 2009
	Traigh na Beirigh	-126 $\pm$ 39	4,540 - 4,240 BC	Ascough <i>et al.</i> 2017
	Northton	64 $\pm$ 41 79 $\pm$ 32	6,390 - 6,290 BC 6,390 - 6,230 BC	Ascough <i>et al.</i> 2017 Ascough <i>et al.</i> 2007
Outer Hebrides North Uist	Baleshare	-79 $\pm$ 17	252 BC - 149 AD	Ascough <i>et al.</i> 2004
		68 $\pm$ 95	77 BC - 111 AD	Reimer <i>et al.</i> 2002
Outer Hebrides South Uist	Hornish point	-79 $\pm$ 17	252 BC - 149 AD	Ascough <i>et al.</i> 2004
		-184 $\pm$ 122 -146 $\pm$ 71	394 BC - 24 AD	Reimer <i>et al.</i> 2002
Inner Hebrides (& Mainland)	Carding Mill Bay	150 $\pm$ 28	3,641 - 3,521 BC	Russell <i>et al.</i> 2015
		86 $\pm$ 67	3,942 - 3,653 BC	Reimer <i>et al.</i> 2002
		-44 $\pm$ 91	3,965 - 3,714 BC	Reimer <i>et al.</i> 2002
	Sand	64 $\pm$ 41	6,480 - 6,420 BC	Ascough <i>et al.</i> 2017
64 $\pm$ 19		6,480 - 6,420 BC	Ascough <i>et al.</i> 2007	

## S5- Details on CSRA dates

Table S3: Details of CSRA dates on pottery vessels and aquatic biomarker detection

Sample	Phase	Layer	BRAMS #	mCO <sub>2</sub> (µg)	Age ± 1σ (BP)	σ range	Comments
<b>BN89-C<sub>16:0</sub></b>	LIA 2	BAC	1549.1.1	199	1396 ± 29	••	No aquatic biomarkers
<b>BN89-C<sub>18:0</sub></b>	LIA 2	BAC	1549.1.2	312	1336 ± 27		
<b>BN89-C<sub>16:0</sub></b>	LIA 2	BAC	1549.2.1	230	1404 ± 30		
<b>BN89-C<sub>18:0</sub></b>	LIA 2	BAC	1549.2.2	373	1322 ± 29		
<b>BN74-C<sub>16:0</sub></b>	LIA 2	BAC	1551.1.1	388	1394 ± 27	••	APAAs
<b>BN74-C<sub>18:0</sub></b>	LIA 2	BAC	1551.1.2	534	1369 ± 26		
<b>BN74-C<sub>16:0</sub></b>	LIA 2	BAC	1551.2.1	379	1299 ± 29		
<b>BN74-C<sub>18:0</sub></b>	LIA 2	BAC	1551.2.2	448	1273 ± 29		
<b>BN77-C<sub>16:0</sub></b>	LIA 2	BAF	1605.1.1	234	1364 ± 29	•	No aquatic biomarkers
<b>BN77-C<sub>18:0</sub></b>	LIA 2	BAF	1605.1.2	293	1375 ± 28		
<b>BN87-C<sub>16:0</sub></b>	LIA 2	BAF	1604.1.1	175	1292 ± 29	•	APAAs
<b>BN87-C<sub>18:0</sub></b>	LIA 2	BAF	1604.1.2	246	1315 ± 28		
<b>BN88-C<sub>16:0</sub></b>	LIA 2	BAG	1548.1.1	216	1720 ± 28	•	APAAs, DHYAs
<b>BN88-C<sub>18:0</sub></b>	LIA 2	BAG	1548.1.2	141	1729 ± 30		
<b>BN88-C<sub>16:0</sub></b>	LIA 2	BAG	1548.2.1	228	1726 ± 30		
<b>BN88-C<sub>18:0</sub></b>	LIA 2	BAG	1548.2.2	165	1728 ± 32		
<b>BN35-C<sub>16:0</sub></b>	EN	BBD	1552.1.1	112	1389 ± 33	X	APAAs, DHYAs
<b>BN35-C<sub>18:0</sub></b>	EN	BBD	1552.1.2	142	1255 ± 30		
<b>BN35-C<sub>16:0</sub></b>	EN	BBD	1552.2.1	136	1151 ± 33		
<b>BN35-C<sub>18:0</sub></b>	EN	BBD	1552.2.2	125	1161 ± 34		
<b>BN91-C<sub>16:0</sub></b>	EN	BBD	1603.1.1	207	1372 ± 29	-	APAAs, DHYAs - No internal control
<b>BN91-C<sub>18:0</sub></b>	EN	BBD	1603.1.2	79	-		
<b>BN101-C<sub>16:0</sub></b>	EN	BBD	1550.1.1	110	1033 ± 27	X	No aquatic biomarkers
<b>BN101-C<sub>18:0</sub></b>	EN	BBD	1550.1.2	105	1420 ± 31		
<b>BN101-C<sub>16:0</sub></b>	EN	BBD	1550.2.1	110	1469 ± 33		
<b>BN101-C<sub>18:0</sub></b>	EN	BBD	1550.2.2	70	-		
<b>BN105-C<sub>16:0</sub></b>	EN	BBD	1547.1.1	164	1288 ± 30	•	No aquatic biomarkers
<b>BN105-C<sub>18:0</sub></b>	EN	BBD	1547.1.2	128	1247 ± 31		
<b>BN105-C<sub>16:0</sub></b>	EN	BBD	1547.2.1	120	1370 ± 34		
<b>BN105-C<sub>18:0</sub></b>	EN	BBD	1547.2.2	117	1280 ± 34		
<b>BN110-C<sub>16:0</sub></b>	EN	BBA	1608.1.1	184	1360 ± 29	••	APAAs
<b>BN110-C<sub>18:0</sub></b>	EN	BBA	1608.1.2	153	1288 ± 30		
<b>BN115-C<sub>16:0</sub></b>	MN	BCA	1609.1.1	219	989 ± 28	•	No aquatic biomarkers
<b>BN115-C<sub>18:0</sub></b>	MN	BCA	1609.1.2	214	984 ± 29		
<b>BN142-C<sub>16:0</sub>C<sub>18:0</sub></b>	MN	BCC	2069.1.1	238	1007 ± 29	-	DHYAs - No internal control
<b>BN149-C<sub>16:0</sub>C<sub>18:0</sub></b>	MN	BCC	2064.1.1	145	1768 ± 34	-	DHYAs - No internal control
<b>BN160-C<sub>16:0</sub></b>	MN	BCC	2066.1.1	-	1230 ± 29	•	No aquatic biomarkers
<b>BN160-C<sub>18:0</sub></b>	MN	BCC	2066.1.2	216	1165 ± 31		
<b>BN165-C<sub>16:0</sub></b>	MN	BCC	2063.1.1	311	1080 ± 29	•	No aquatic biomarkers
<b>BN165-C<sub>18:0</sub></b>	MN	BCC	2063.1.2	285	1040 ± 29		
<b>BN167-C<sub>16:0</sub>C<sub>18:0</sub></b>	MN	BCC	2068.1.1	225	1295 ± 31	-	APAAs, TMTD - No internal control
<b>BN168-C<sub>16:0</sub></b>	MN	BCC	2061.1.1	123	1211 ± 34	X	No aquatic biomarkers
<b>BN168-C<sub>18:0</sub></b>	MN	BCC	2061.1.2	253	1062 ± 34		
<b>BN173-C<sub>16:0</sub>C<sub>18:0</sub></b>	MN	BCC	2067.1.1	167	1234 ± 32	-	APAAs, DHYAs, TMTD - No internal control
<b>BN174-C<sub>16:0</sub></b>	MN	BCC	2062.1.1	219	1152 ± 31	••	APAAs
<b>BN174-C<sub>18:0</sub></b>	MN	BCC	2062.1.2	296	1065 ± 33		
<b>BN38-C<sub>16:0</sub></b>	MN	AD	1606.1.1	121	871 ± 31	-	No aquatic biomarkers - No internal control
<b>BN36-C<sub>16:0</sub></b>	LN	AG	1607.1.1	201	767 ± 29	••	APAAs, DHYAs
<b>BN36-C<sub>18:0</sub></b>	LN	AG	1607.1.2	115	816 ± 31		

## S6- Quantification of marine derived-C using FRUITS (v2.1)

Table S4: Determination of the percentage marine products in pottery vessels CSRA dated using FRUITS.

Target/ Consumer	Source/Food	Mean	sd	2.5pc	median	97.5pc
<b>BN89</b>	ruminant adipose	0.646	0.1001	0.4117	0.6601	0.8014
	marine	0.354	0.1001	0.1986	0.3399	0.5884
<b>BN74</b>	ruminant adipose	0.6361	0.09734	0.4126	0.6486	0.7901
	marine	0.3639	0.09734	0.21	0.3514	0.5875
<b>BN174</b>	ruminant adipose	0.537	0.1158	0.2521	0.5545	0.7138
	marine	0.463	0.1158	0.2862	0.4455	0.7491
<b>BN160</b>	milk	0.7067	0.09375	0.5054	0.7157	0.8641
	marine	0.2933	0.09375	0.136	0.2843	0.4951
<b>BN165</b>	milk	0.6372	0.1027	0.389	0.6489	0.7997
	marine	0.3628	0.1027	0.2004	0.3511	0.6111
<b>BN77</b>	ruminant adipose	0.6823	0.09332	0.4724	0.6928	0.8323
	marine	0.3177	0.09332	0.1678	0.3072	0.5276
<b>BN88</b>	ruminant adipose	0.2925	0.1237	0.03621	0.3041	0.505
	marine	0.7075	0.1237	0.4951	0.696	0.9638
<b>BN110</b>	ruminant adipose	0.3656	0.1253	0.08533	0.3807	0.5757
	marine	0.6344	0.1253	0.4243	0.6193	0.9148
<b>BN87</b>	milk	0.6823	0.08895	0.4844	0.6914	0.8288
	marine	0.3177	0.08895	0.1713	0.3086	0.5156
<b>BN35</b>	milk	0.6692	0.1059	0.4316	0.681	0.8465
	marine	0.3308	0.1059	0.1536	0.319	0.5685
<b>BN105</b>	milk	0.7343	0.094	0.5228	0.7439	0.8934
	marine	0.2657	0.094	0.1067	0.2561	0.4774
<b>BN115</b>	milk	0.4731	0.1099	0.2199	0.4864	0.6534
	marine	0.5269	0.1099	0.3468	0.5136	0.7803
<b>BN36</b>	milk	0.4835	0.1136	0.2177	0.4982	0.6682
	marine	0.5165	0.1136	0.3323	0.5018	0.7824

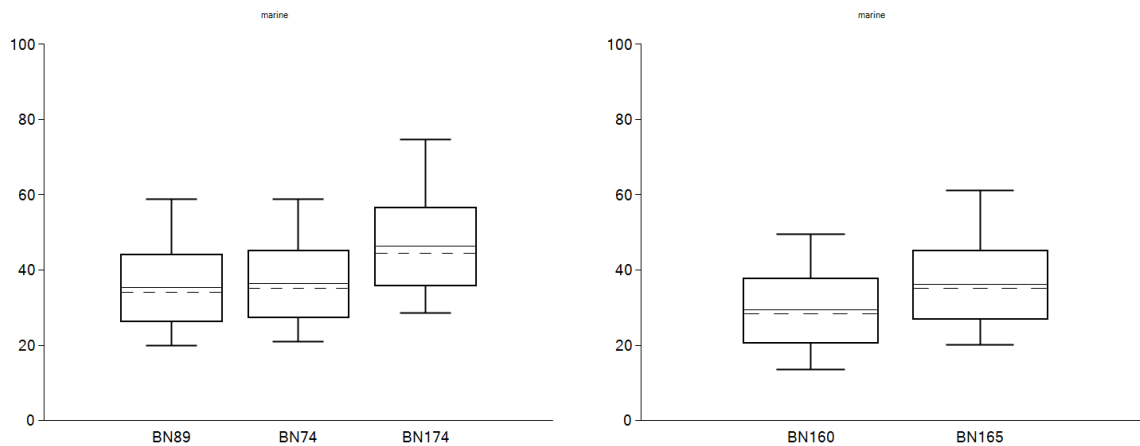


Figure S3: Box and whisker plot for the marine contribution in potsherds dominated by (a) ruminant adipose fats and (b) dairy fats for the potsherds CSRA dated and corrected in the main paper (from FRUITS v2.1).

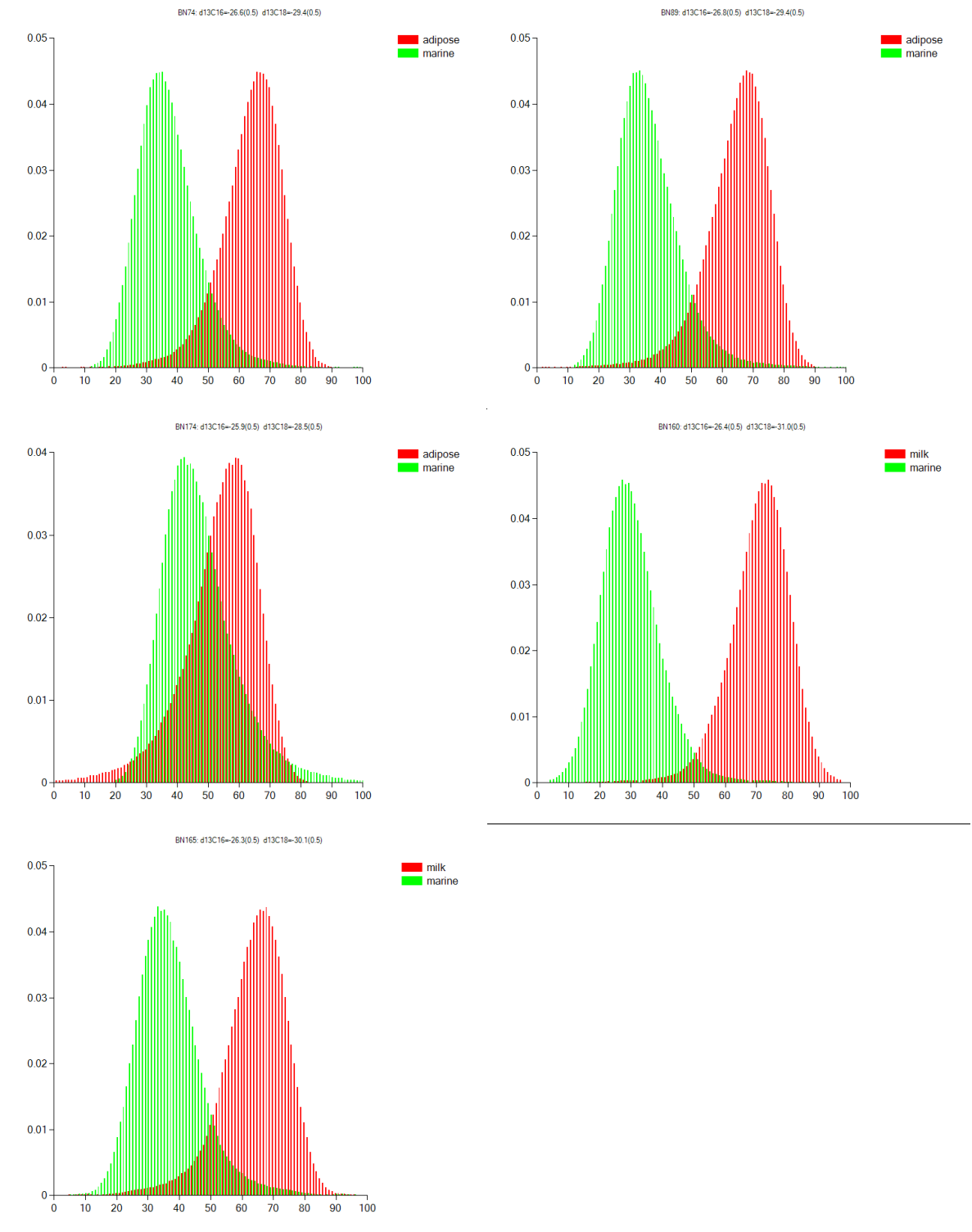


Figure S4: Probability distribution for the proportion of marine (green) and terrestrial (red) resources for the potsherds CSRA dated and corrected in the main paper (from FRUITS v2.1)

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