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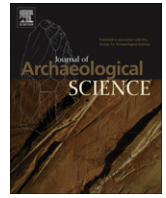
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## Interpreting the expansion of sea fishing in medieval Europe using stable isotope analysis of archaeological cod bones

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### ABSTRACT

Archaeological fish bones reveal increases in marine fish utilisation in Northern and Western Europe beginning in the 10th and 11th centuries AD. We use stable isotope signatures from 300 archaeological cod (*Gadus morhua*) bones to determine whether this sea fishing revolution resulted from increased local fishing or the introduction of preserved fish transported from distant waters such as Arctic Norway, Iceland and/or the Northern Isles of Scotland (Orkney and Shetland). Results from 12 settlements in England and Flanders (Belgium) indicate that catches were initially local. Between the 9th and 12th centuries most bones represented fish from the southern North Sea. Conversely, by the 13th to 14th centuries demand was increasingly met through long distance transport – signalling the onset of the globalisation of commercial fisheries and suggesting that cities such as London quickly outgrew the capacity of local fish supplies.

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### 1. Introduction

The analysis of fish bones from archaeological sites indicates increases in marine fishing in Northern and Western Europe around the turn of the first and second millennia AD (Benecke,

1982; Barrett et al., 1999, 2004a; Enghoff, 1999, 2000; Ervynck et al., 2004; Jones, 1988; Løugas, 2008; Makowiecki, 2008; Müldner, 2009). Past research has identified two alternative scenarios that might explain this change in the relationship between people and the sea – a transformation that started the globalisation of the world's fisheries and that has continued to the present (Barrett et al., 2004a; Hoffmann, 1996, 2002; Perdikaris and McGovern, 2008; Roberts, 2007). One possibility is that the

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medieval rise in sea fish utilisation resulted from a change in supply – the start of long-range trade in staple goods from the north-east Atlantic, Arctic Norway and the Baltic Sea to the urbanised southern North Sea littoral. A development of this kind could have been driven by the Viking Age Scandinavian diaspora (connecting producer and consumer regions in new networks), increased ship capacities (allowing more bulky cargoes) and the Medieval Warm Period (influencing the success of seafaring and the productivity of Arctic cod stocks). Alternatively, the medieval fishing revolution could have resulted from a demand-led intensification of local marine fishing – to meet new food requirements influenced by growing human populations, Christian fasting practises (fish were substituted for the meat of quadrupeds), the over-exploitation of freshwater fish stocks and the effects of human-induced habitat change on freshwater species. It is important to determine which of these alternative hypotheses is most probable. The resulting growth of intensive and/or extensive marine fishing has had profound social, economic and environmental impacts on both short and long timescales (Hoffmann, 2002; Jackson, 2008; Jackson et al., 2001; Rick and Erlandson, 2008; Roberts, 2007).

The written evidence regarding Europe's medieval fisheries is sporadic until the 14th century (Holm et al., 1996; Kowaleski, 2003; Starkey et al., 2009) – postdating the years during which, as demonstrated by archaeological evidence, large-scale sea fishing first developed (Barrett et al., 2004a). Thus to differentiate the alternative scenarios we ideally need a way to discover where the fish whose bones have been recovered from archaeological excavations were originally caught. At the very least it is essential to reconstruct when preserved sea fish were first transported over hundreds or thousands of kilometres, and what the relative magnitude of this long-range transport was in comparison with the produce of local marine fishing.

Having developed such a technique, based on stable carbon and nitrogen isotope analysis of fish bone collagen (Barrett et al., 2008), we apply it here to finds of cod (*Gadus morhua*) bones from London and 11 other urban centres and villages bordering the southern North Sea in England and Flanders (Belgium). Alternative methodologies exist for determining the origin of modern cod catches, but these require otoliths which rarely preserve in archaeological deposits, employ trace elements susceptible to contamination from the burial environment or use genetic markers which may have changed over time due to modern fishing pressure (Hutchinson et al., 2003; Pampoulie et al., 2008; Svedäng et al., 2010).

In addition to cod, the taxa most important in Europe's sea fishing revolution included herring (*Clupea harengus*), flatfish (Pleuronectidae), haddock (*Melanogrammus aeglefinus*) and other gadids (Gadidae). Of these, herring is the most consistently abundant find in the southern North Sea and Baltic regions (Barrett et al., 1999, 2004a; Enghoff, 1999, 2000; Ervynck et al., 2004; Jones, 1988; Lõugas, 2008; Makowiecki, 2008). However, cod is the best archaeological proxy for the study of long-range fish trade. This species is known to have been widely traded in the past (Starkey et al., 2009), has bones of an ideal size to extract sufficient purified collagen for analysis and (having large bones) can be recovered on archaeological sites even when sediment is not sieved. Crucially, it also has well-understood migration patterns, with many populations remaining close to their spawning grounds (Bagge et al., 1994; Deutsch and Berth, 2006; Holmes et al., 2008; Sarvas and Fevolden, 2005; Svedäng et al., 2007, 2010; Wright et al., 2006).

Our study is based on two steps, involving the stable isotope analysis of collagen from archaeological cranial (control) and postcranial (target) cod bones respectively. Cranial bones are likely to derive from relatively local catches because, prior to

refrigeration, cod were typically decapitated and dried (with or without salting) for long-range transport or trade. Conversely, postcranial bones (specifically vertebrae and cleithra, the latter of which support the pectoral fins) were typically left in traditional fish products such as stockfish (air-dried cod) before transport to consumers (Barrett, 1997; Coy and Hamilton-Dyer, 2005). Thus target specimens could be from either local catches or long-range transport – and a comparison of the stable isotope values of control and target specimens can reveal the introduction of non-local catches (Barrett et al., 2008).

## 2. Material and methods

### 2.1. Control and target samples

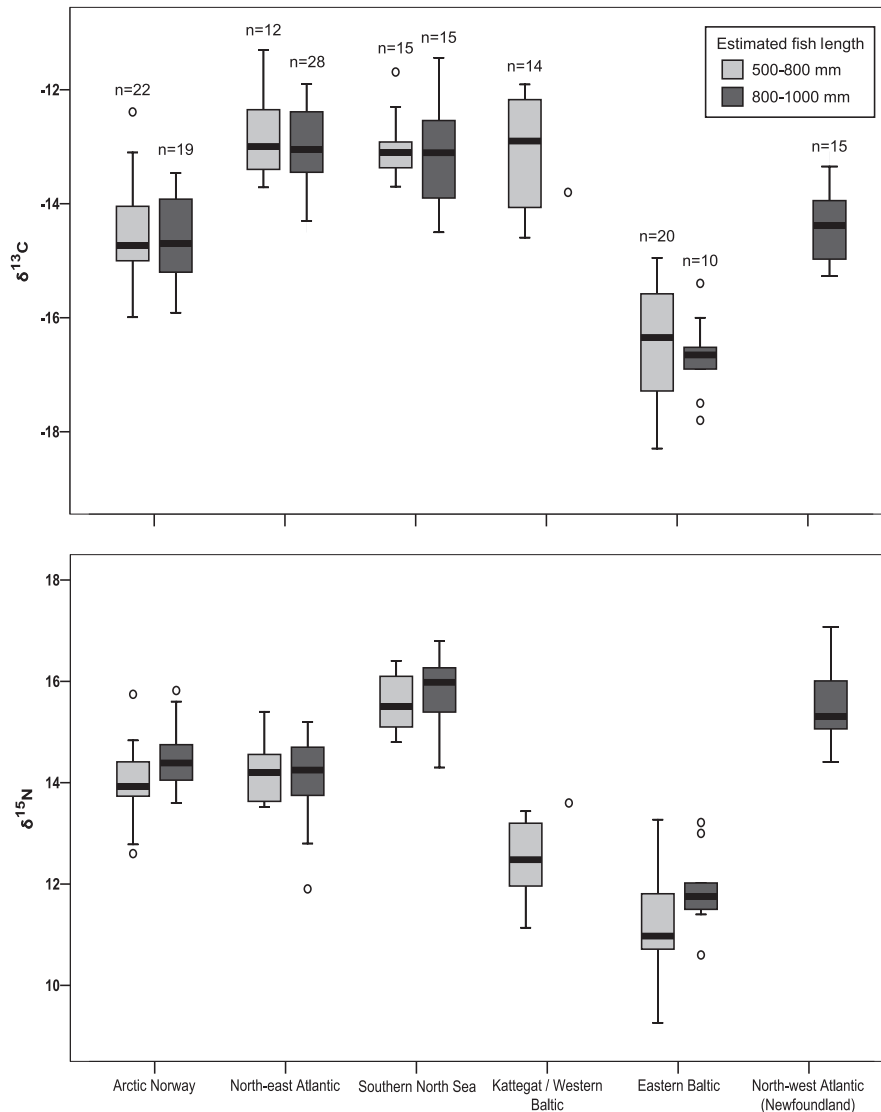
Samples of archaeological cod cranial ( $n = 208$ ), and postcranial ( $n = 188$ ) bones for  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  analysis of collagen were obtained from over 50 archaeological assemblages from six analytical regions: Newfoundland, Arctic Norway, the north-east Atlantic (Iceland and northern Scotland), the southern North Sea (below  $55^\circ\text{N}$ ) and English Channel, the Kattegat/western Baltic Sea and the eastern Baltic Sea. Studies of cod migration patterns indicate limited natural movement between the designated regions (Bagge et al., 1994; Deutsch and Berth, 2006; Holmes et al., 2008; Sarvas and Fevolden, 2005; Svedäng et al., 2007, 2010; Wright et al., 2006). Where practicable, sampling regions were specifically defined based on the known existence of separate cod populations (e.g. the Kattegat/western Baltic versus the eastern Baltic) (Bagge et al., 1994; Deutsch and Berth, 2006). The regions considered include known fishing grounds exploited by Europeans in medieval and/or post-medieval times (Starkey et al., 2009).

Sample selection aimed at good geographical and temporal coverage of each region, but was constrained by standard archaeological contingencies of preservation and accessibility, exacerbated by variable laboratory success rates. 171 of the cranial specimens and 129 of the postcranial specimens were successful, producing high yields of collagen with acceptable atomic C:N ratios (see Section 2.6). All except two of the target (postcranial) specimens, both from Wharram Percy in England (Barrett et al., 2008), are reported here for the first time. Data regarding a subset of 74 of the control (cranial) specimens was summarised previously in a pilot study (Barrett et al., 2008).

Outliers within the control (cranial) samples might reflect rare instances of fish imported without prior decapitation (Jonsson, 1986). Thus we excluded the single clearest outlier from each region, defined by the greatest mean of the z-scores for  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$ . No outliers were excluded from the target data set.

### 2.2. Size-based variability

Since higher trophic levels in older, larger fish could result in elevated  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values (DeNiro and Epstein, 1978; Jennings et al., 2002; Minagawa and Wada, 1984; Schoeninger and DeNiro, 1984), the archaeological samples were restricted to cod with estimated total lengths in the 500–1000 mm range (based on bone measurements and comparison with reference specimens of known size) (Barrett et al., 2008). Size effects were also mitigated by analysing complete bone cross-sections – providing approximate life-time averages given the incremental nature of fish bone growth (Van Neer et al., 1999). Comparison of  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values between smaller (500–800 mm) and larger (800–1000 mm) fish within each region using  $t$ -tests revealed only one significant difference, for  $\delta^{15}\text{N}$  in Arctic Norway ( $t = -2.11$ ,  $df = 39$ ,  $p = 0.041$ ) (see Fig. 1). In future research it may be possible to further control for trophic level effects using amino acid-specific stable isotope



**Fig. 1.** Boxplots showing distributions of  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  for smaller (500–800 mm total length) and larger (800–1000 mm total length) cod from the control groups, by region.

compositions (Hannides et al., 2009). The present demonstration of spatial patterning in collagen stable isotope values is, however, a necessary precursor to work of this kind.

### 2.3. Temporal variability

To address temporal variability in stable isotope signatures we did not include any archaeological samples of 19th–21st-century date that might have altered  $\delta^{15}\text{N}$  values – due to radical changes in food webs and/or pollutants such as agricultural fertilisers and sewage (Christensen and Richardson, 2008; Doney, 2010; Hansson et al., 1997; Struck et al., 2000). The archaeological control samples of 9th–18th century date were also classified into five two-century groups and compared within each region using boxplots (see Fig. 2) and ANOVA. Significant temporal differences in  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values were absent from all regions except the eastern Baltic, where there is a decrease in  $\delta^{13}\text{C}$  between the 13th to 14th and 15th to 16th centuries ( $F(\text{df } 1, 28) = 15.58, p < 0.001$ ). The magnitude of this effect is small relative to differences between regions, and the eastern Baltic remains clearly separated from other regions when  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  are taken together (Fig. 3).

### 2.4. Variability within regions

Since the regions are large, treating them as single units may mask finer-scale geographic variation. However, ANOVA of the different settlements represented in each region produced only one significant result, for  $\delta^{13}\text{C}$  values from the eastern Baltic ( $F(\text{df } 2, 27) = 3.50, p = 0.045$ ). As for temporal variability, the eastern Baltic nevertheless remains clearly separated when  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  are taken together (Fig. 3).

### 2.5. Intra-skeletal variability

To confirm that there is unlikely to be systematic intra-skeletal variability in the stable isotope signatures employed we analysed the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  of collagen extracted from one cranial bone (a praemaxilla) and two postcranial bones (a cleithrum and an anterior caudal vertebra) from each of eight cod caught in the North Sea in 1992. The individuals ranged from 460 to 950 mm in total length. There was no systematic difference in the isotope values between skeletal elements and all fish showed small intra-individual variability. The maximum ranges between skeletal

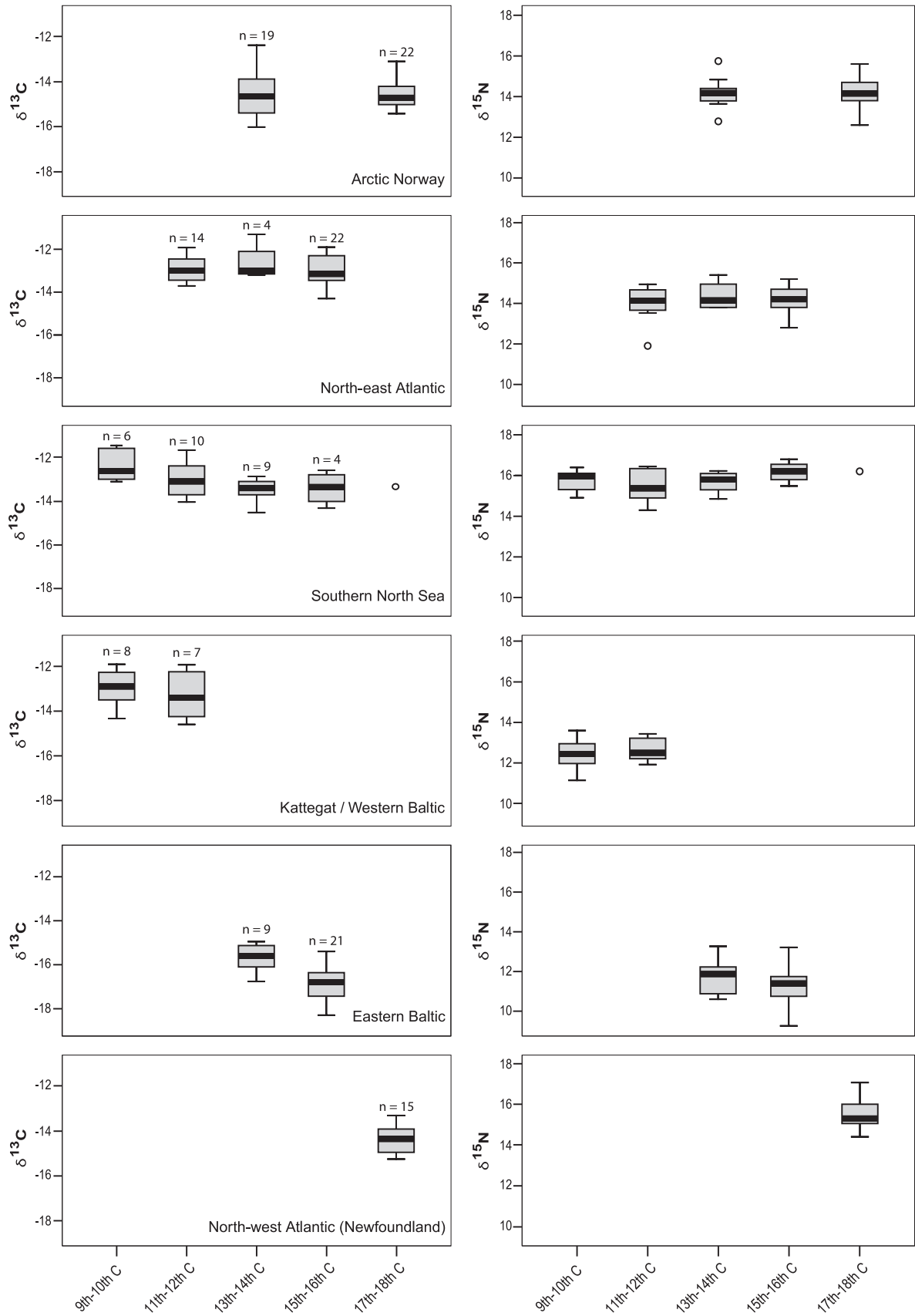
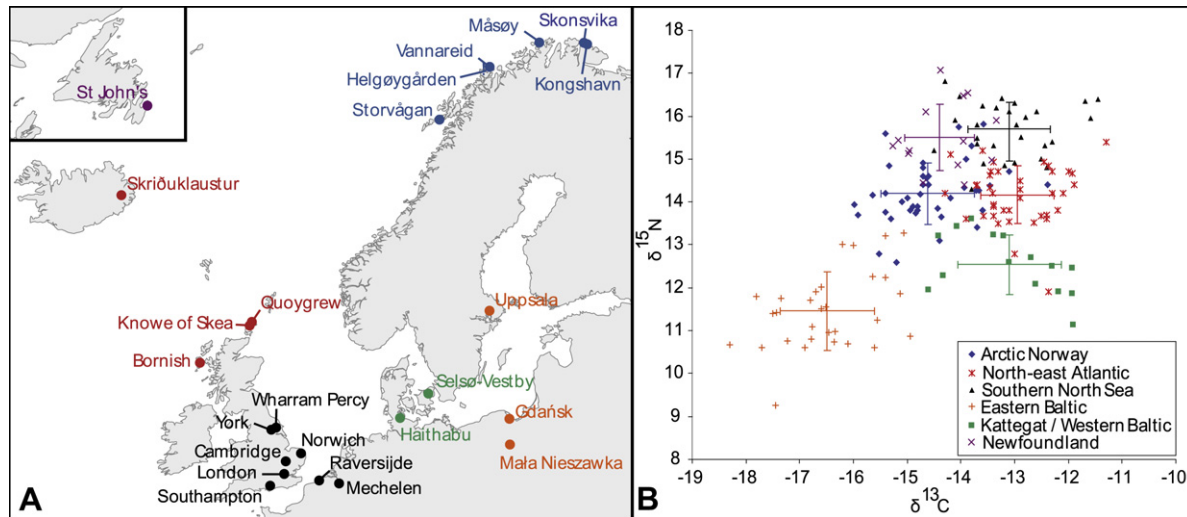


Fig. 2. Boxplots comparing distributions of  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  over time within the control groups, by region.



**Fig. 3.** (A) The archaeological sites from which cod cranial (control) bones were sampled. (B)  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values, including means  $\pm$  one standard deviation, for the 171 control samples grouped into six analytical regions indicated by different coloured labels in 3A.

elements from the same fish for  $\delta^{13}\text{C}$  were 0.1, 0.1, 0.2, 0.2, 0.2, 0.3, 0.4 and 0.6. The equivalent ranges for  $\delta^{15}\text{N}$  were 0.2, 0.2, 0.3, 0.4, 0.4, 0.4, 0.6 and 1.0.

## 2.6. Laboratory methods

Cod bones greater than 1 g in mass were sawn in two, with one subsample archived for further study. For the second subsample (or whole specimen if under 1 g), a complete cross-section (totalling 100–200 mg) of each bone was then processed for stable isotope analysis. Bone collagen was extracted following standard procedures outlined by Richards and Hedges (1999), with the addition of an ultrafiltration step for archaeological samples (Brown et al., 1988). Whole bone samples were demineralised in 0.5M aqueous HCl at 4 °C for 2–5 days until demineralised, and then gelatinised at 70 °C for 48 h, and the resulting solution filtered through a 5–8  $\mu\text{m}$  filter. For the modern samples (used only to study intra-skeletal variability), the gelatinised solution was then lyophilised for 48 h. For the archaeological samples, the gelatinised solution was ultra-filtered through a 30 kDa filter, and then the >30 kDa fraction was lyophilised for 48 h. The resultant ‘collagen’ was analysed in duplicate or triplicate using continuous flow isotope-ratio-monitoring mass spectrometry: a ThermoFinnigan Flash EA coupled to a Delta Plus XP mass spectrometer at the Department of Human Evolution, Max Planck Institute for Evolutionary Anthropology, Leipzig, Germany for the archaeological samples, and a Costech EA coupled to a Delta V mass spectrometer at the Godwin Laboratory, Department of Earth Sciences, University of Cambridge, for the modern samples. The  $\delta^{13}\text{C}$  values are reported relative to the V-PDB scale, and  $\delta^{15}\text{N}$  values relative to the AIR scale. Errors on both the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  measurements are less than 0.2‰. Amounts of carbon and nitrogen in the collagen extract were measured, and we only report isotope values from those samples with acceptable atomic C:N ratios, defined as between 2.9 and 3.6 (DeNiro, 1985).

## 3. Results

The resulting control data – from cranial bones – show that cod from within particular regions of ocean and sea share similar isotopic values, yet there are clear differences between regions (Fig. 3, see Appendix 1 for the data set). It is therefore possible, within meaningful limits of probability, to use the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$

control data to attribute individual archaeological cod bones to the regions considered (having ruled out systematic variability by skeletal element, size and time). We attribute the target bones to probable source using discriminant function analysis (DFA), the utility of which can be evaluated by considering how successful the method is at reclassifying the original data. Treating all six regions as separate sources the reclassification success rates range from 61% to 93% (Table 1). However, by combining the north-east Atlantic and Arctic Norwegian regions they increase to between 73% and 93%. Grouping the data in this way is appropriate because historical sources indicate that dried cod produced in both Norway and the Scandinavian colonies of the north-east Atlantic (particularly Iceland) were traded via the port of Bergen during the Middle Ages (Starkey et al., 2009). For the consumers, these catches thus all represented the same product.

We take the cautious view that cod from the north-west Atlantic are unlikely to have been traded to Europe before the transatlantic expedition of Zuan Caboto (‘John Cabot’) in 1497 (cf. McGhee, 2003). Thus DFA has been performed both excluding and including the control data from Newfoundland – for applicability to

**Table 1**

Discriminant function analysis reclassification success rates for the archaeological control samples of cod bone.

Regions (All separate)	Excluding Newfoundland			Including Newfoundland		
	n	Correct	%	n	Correct	%
Arctic Norway	41	29	71	41	25	61
North-east Atlantic	40	32	80	40	32	80
Southern North Sea	30	24	80	30	21	70
Eastern Baltic	30	28	93	30	28	93
Kattegat/Western Baltic	15	12	80	15	12	80
Newfoundland				15	10	67
Overall	156	125	80	171	128	75

Regions (the ‘North’ Combined)	Excluding Newfoundland			Including Newfoundland		
	n	Correct	%	n	Correct	%
Arctic Norway and NE Atlantic	81	66	81	81	62	77
Southern North Sea	30	27	90	30	23	77
Eastern Baltic	30	28	93	30	28	93
Kattegat/Western Baltic	15	12	80	15	12	80
Newfoundland				15	11	73
Overall	156	133	85	171	136	80

earlier and later archaeological case studies respectively. Without Newfoundland, the DFA reclassification success rates range from 80% to 93% when the north-east Atlantic and Arctic Norway are combined. Overall the success rates are high. This result makes ecological sense given that there are established relationships between the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values of fish tissues and environment, diet and trophic level – which vary from region to region (Fredriksen, 2003; Jennings and Warr, 2003; Sweeting et al., 2007; Weidman and Millner, 2000).

Turning to the second step of the study, because target bones could be from either local catches or long-range transport, differences in the relative abundance of local and imported fish can be evaluated by comparing their probable sources based on DFA of the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  data. The analysed bones include the earliest known finds of cod from medieval settlements such as York, Flixborough, Norwich, London, Southampton, Ghent and Mechelen – plus (within the limits of available samples) subsequent time series ending in the 15th–16th centuries for England and the 17th–18th centuries for Flanders. Following the conservative strategy noted above, the north-west Atlantic (Newfoundland) is only included as a potential source for DFA of samples dating from the 15th to 16th centuries and later.

Details of each target specimen, including probabilities of group membership based on DFA, are provided in Appendix 2. Like all

archaeological characterisation studies, the attributions to source are not absolute. Not all possible fishing regions are represented in the control data, the limited number of control samples (an unavoidable characteristic of archaeology) is unlikely to represent fully the variability within each region and there is some overlap in the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values of the data from each potential source. Nevertheless, the method can serve as a guide to when, where and in what proportions (*vis-à-vis* local catches) cod transported from distant waters first appeared in settlements around the southern North Sea.

Fig. 4 summarises the DFA results for the target data from London and other towns and villages in England and Flanders. It shows that the marine fishing revolution at the turn of the first and second millennia AD (Barrett et al., 2004a) was predominantly the result of local fishing – rather than the introduction of long-range trade goods. If this conclusion is true for cod it is reasonable to believe that the bones of other gadids, herring and flatfish found in archaeological remains dated to this time (Barrett et al., 2004b; Eryvynck et al., 2004) were also from local catches in the southern North Sea. The 10th–11th century increase in sea fishing therefore seems to derive predominantly from the intensification of local (demand-led) production rather than changes in distant supply. Based on our DFA analyses, cod of northern origin may occasionally have been imported to towns such as York and London in the

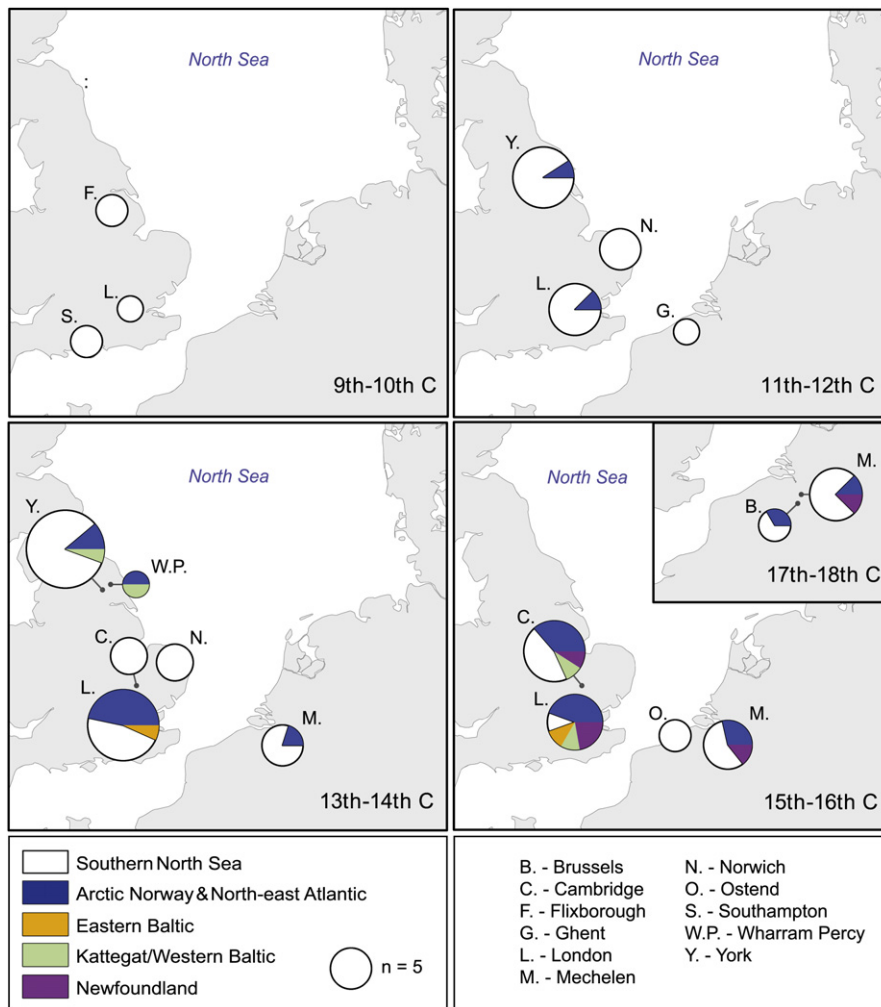
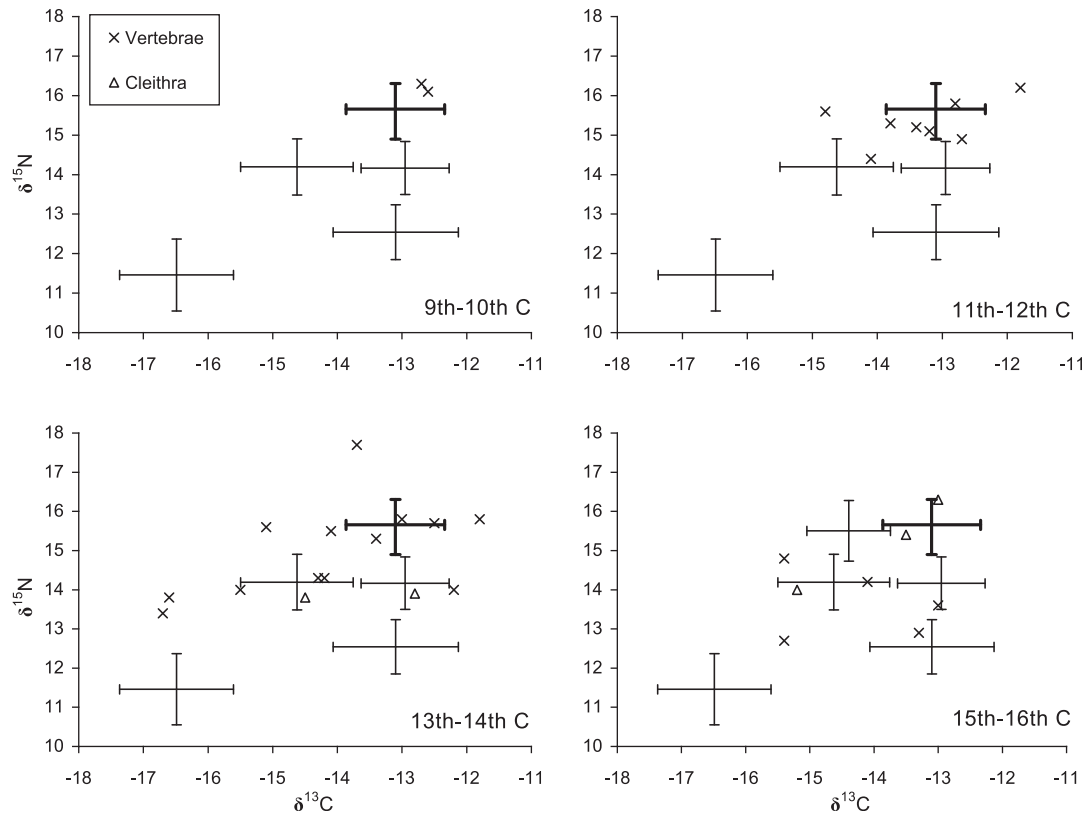


Fig. 4. Local (southern North Sea) and imported (other regions) cod bones in England and Flanders (Belgium) based on discriminant function analysis of  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  measurements on 129 archaeological vertebrae and cleithra dating between the 9th and 18th centuries. The size of each pie chart reflects the number of bones analysed. Sites with only one successful analysis are omitted for clarity.



**Fig. 5.**  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  values for cod target bones (vertebrae and cleithra) from London dating between the 9th and 16th centuries, superimposed on the means  $\pm$  one standard deviation for the control data from Fig. 3B. The local (southern North Sea) standard deviation values are emphasised. Newfoundland is added as a potential source for samples of 15th–16th century date.

11th–12th centuries – consistent with previous results from a pilot study of the Viking Age town of Haithabu in the Baltic Sea region (Barrett et al., 2008). This trade may have been of importance to northern producers, but was probably a ‘drop in the bucket’ for consumers in the southern North Sea region. Very few of the pre-13th-century target bones from England and Flanders have been attributed to a northern source (one of 12 specimens from York and one of 10 specimens from London).

By the 13th–14th centuries, conversely, more than half of the cod specimens from London are attributed to a source beyond the southern North Sea. Most are probably from very distant waters – the north-east Atlantic or Arctic Norway. The single specimen attributed to the eastern Baltic could be a statistical outlier, but preserved cod were produced in the region (Jonsson, 1986) and there was maritime traffic between England and eastern Europe at this time (Lloyd, 1991). Other settlements show a less profound draw on distant resources, still relying predominantly on North Sea fishing.

In the 15<sup>th</sup> to 16<sup>th</sup> centuries, the trend towards more imports continues. Almost all of the specimens from London are attributed to distant sources and Cambridge and Mechelen also yield higher proportions of imported cod bones than in previous centuries. The north-east Atlantic and/or Arctic Norway remain the most common sources, but some of the bones may also be from the north-west Atlantic and the Baltic Sea. In the 17th and 18th centuries, for which only Flemish samples were available, a mix of distant and local North Sea sources remains evident.

The introduction of prepared fish from non-local sources can also be summarised in a less precise way, without reliance on the attribution of individual specimens to a probable source and the level of uncertainty this procedure necessarily entails. Fig. 5 plots

the  $\delta^{13}\text{C}$  and  $\delta^{15}\text{N}$  results for target bones from 9th to 16th century London (the best time series) against the means  $\pm$  one standard deviation of the control data from each potential source region. Comparing this graph with Fig. 3, it is clear that the London targets differ from the southern North Sea controls (which include cranial bones from London) in having both greater dispersion and differing values. The higher variation suggests that the vertebrae and cleithra derive from a wider range of sources than the cranial elements (indicating the presence of preserved fish transported from multiple non-local fishing grounds). Moreover, although some target specimens match local southern North Sea isotope values, many instead match (within one or two standard deviations of the control data means) northern regions known to have exported dried cod. Through time, the trends towards greater dispersion and different isotope values (*vis-à-vis* the southern North Sea control data) first become clear in the 13th–14th centuries.

#### 4. Discussion

Our results suggest that the revolution in sea fishing at the turn of the first and second millennia AD resulted from a demand-driven intensification of local fishing rather than a supply-driven introduction of long-range trade goods. But what caused this change in demand for sea fish? The insufficiency or reduction of freshwater fish stocks, indicated by a concurrent decline in the relative importance of freshwater species in towns, must have played a role (Barrett et al., 2004a; Hoffmann, 1996; Van Neer and Ervynck, 2010). However, freshwater fish (increasingly cultivated in ponds) did continue to be available to the wealthy and in the countryside (Dyer, 1988; Van Neer and Ervynck, 2010), suggesting that their limited availability alone is unlikely to have driven fishermen to the



sea. Christian dietary practises (encouraging the consumption of fish during Lent and other fast days) must also have contributed to the sea fishing revolution, but are unlikely to be its primary cause because they were variable through time, by region and in different social groups (Barrett et al., 2004a; Eryvnyck, 1997; Woolgar, 2000).

A demographic imperative – from increasing numbers of urban consumers facing the need for new sources of inexpensive and easily preserved protein – may thus also be significant. Moreover, its impact on sea fishing continued to escalate. Already by the 13th–14th centuries the appetite of growing populations in urban centres like London became sufficiently strong that local fishing in the southern North Sea could no longer supply the demand. Marine fisheries thus began to ‘globalise’, with fish initially traded over long distances (from Arctic Norway, Iceland and/or the Northern Isles of Scotland to London via Bergen, for example) and then sought directly by fishermen from North Sea ports who travelled to increasingly distant waters. English boats were fishing off Iceland by the 1400s before moving on to the coast of North America in subsequent centuries (Hoffmann, 2002).

These conclusions illuminate the origins and development of the medieval sea fishing revolution. They also have implications for the chronology and magnitude of medieval urban expansion. Populous towns must have emerged early (to influence the rise of sea fishing in the 10th–11th centuries) and, in the case of London at least, grown quickly (to exceed the capacity of regional fish supplies by the 13th–14th centuries). Following this reasoning the continuing growth of demand in the 15th–16th centuries may reflect the increasing concentration of population in towns after the demographic crash of the 14th century (cf. Epstein, 2009). Concurrently, our findings increase the likelihood that medieval urban expansion impacted both freshwater and marine ecosystems – the start of an on-going process (Hoffmann, 2002; Jackson et al., 2001; Roberts, 2007).

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## Appendix. Supplementary data

Supplementary data related to this article can be found online at doi:10.1016/j.jas.2011.02.017

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