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The Diet-Body Offset in Human Nitrogen Isotopic Values: A Controlled Dietary Study

T.C. O'Connell, C.J. Kneale, N. Tasevska and G.G.C. Kuhnle

1Department of Archaeology and Anthropology, University of Cambridge, UK
2McDonald Institute for Archaeological Research, University of Cambridge, UK
3MRC Dunn Human Nutrition Unit, Wellcome Trust/MRC, Building, Cambridge, UK
4Department of Food and Nutritional Sciences, University of Reading, UK
5Department of Public Health and Primary Care, MRC Centre for Nutritional Epidemiology in Cancer Prevention and Survival, University of Cambridge, UK

ABSTRACT The “trophic level enrichment” between diet and body results in an overall increase in nitrogen isotopic values as the food chain is ascended. Quantifying the diet–body Δ15N spacing has proved difficult, particularly for humans. The value is usually assumed to be +3–5‰ in the archaeological literature. We report here the first (to our knowledge) data from humans on isotopically known diets, comparing dietary intake and a body tissue sample, that of red blood cells. Samples were taken from 11 subjects on controlled diets for a 30-day period, where the controlled diets were designed to match each individual’s habitual diet, thus reducing problems with short-term changes in diet causing isotopic changes in the body pool. The Δ15N_diet-RBC was measured as +3.5‰. Using measured offsets from other studies, we estimate the human Δ15N_diet-keratin as +5.0–5.3‰, which is in good agreement with values derived from the two other studies using individual diet records. We also estimate a value for Δ15N_diet-collagen of ≈6‰, again in combination with measured offsets from other studies. This value is larger than usually assumed in palaeodietary studies, which suggests that the proportion of animal protein in prehistoric human diet may have often been overestimated in isotopic studies of palaeodiet. Am J Phys Anthropol 149:426–434, 2012.

KEY WORDS collagen; keratin; blood; nutritional biomarker; trophic level; discrimination factor

Light element isotopic analyses of human and animal body tissues are increasingly used to elucidate dietary patterns in past and living populations, with applications in archaeology, ecology, and nutritional epidemiology. However, the full potential of those analyses remains constrained by our limited understanding of the mechanisms involved in the transfer of the isotopic signature to the body during the absorption and incorporation of food. This is particularly the case with nitrogen isotopes, where there is observed enrichment between diet and body (the “trophic level effect” or Δ15N_diet-body), resulting in an increase in δ15N as the food chain is ascended (DeNiro and Epstein, 1981; Minagawa and Wada, 1984; Schoeninger and DeNiro, 1984). Despite its clear empirical success as a dietary indicator, we do not yet know metabolically how and where the 15N enrichment between diet and body occurs. Ecological studies suggest that mammals, fish, birds, reptiles, and insects all have similar enrichments (Caut et al., 2009), so it seems to be independent of the mode of nitrogen excretion, but there has been little exploration of the cause. Quantifying the enrichment has proved difficult: large-scale ecological studies suggest that the enrichment associated with each trophic level is ≈+3–4‰, while small-scale animal feeding experiments show values anywhere between +1.5 and +6‰ (see review in Caut et al., 2009). In addition to being poorly quantified and understood, the trophic level effect also seems capable of quite large variation under a range of environmental conditions (temperature, altitude, aridity), as well as being potentially affected by physiological factors such as water stress, starvation and growth, digestive physiology and diet composition (for a review see McCue and Pollock, 2008).

For isotopic studies of human diet, the resolution of our interpretations is limited because we do not know what value to use for the 15N enrichment in humans (see Hedges and Reynard, 2007). While broad-scale changes in diet are easily observed in human isotopic values (Vogel and van der Merwe, 1977; Tauber, 1981; Buikstra and Milner, 1991; Lubell et al., 1994; Bonsall et al., 1997; Richards et al., 2003), our lack of knowledge of the Δ15N_diet-body value, and of influencing factors on this parameter, means that we cannot with confidence identify isotopic shifts resulting from small-scale dietary changes. For this, we need to quantify better the Δ15N_diet-body in humans.

QUANTIFYING THE ENRICHMENT

It has been generally assumed that the nitrogen isotopic enrichment in mammals, including humans, is broadly similar, with a Δ15N_diet-body value initially taken

*Correspondence to: T.C. O’Connell, McDonald Institute for Archaeological Research, Downing St, Cambridge, CB4 3DZ, UK. E-mail: tco21@cam.ac.uk

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to be about 3% (DeNiro and Epstein, 1981; Minagawa and Wada, 1984; Schoeninger and DeNiro, 1984; Hare et al., 1991), but more recently values of up to 5% have been postulated (Ambrose, 2000; Jenkins and Partridge, 2001; Bocherens and Drucker, 2003; Sponheimer et al., 2003; Robbins et al., 2005; Caut et al., 2009). General reviews of the ecological literature for animals ranging from invertebrates to large mammals and aquatic and terrestrial species give overall mean $\Delta^{15}N_{\text{diet-body}}$ values of 2.5–3.5%, with a high degree of variability, based on analyses of a range of body tissues (Post, 2002; McCutchan et al., 2003; Vanderklift and Ponsard, 2003). A value of around 3% fits with numerous predator–prey relationships in terrestrial ecological situations (see a summary in Bocherens and Drucker, 2003).

A large number of controlled animal feeding studies have been carried out, to attempt to quantify the offset (see summary in Caut et al., 2009). But for humans, the situation is more complicated, as there are significant difficulties in obtaining reliable data on which to base an estimate of human $\Delta^{15}N_{\text{diet-body}}$. A number of human studies have looked at isotopic variation within populations depending on self-reported diet type (O'Connell and Hedges, 1999a; Bol and Pfieger, 2002; Petzke et al., 2005b), or compared human isotopic variation to estimated diets, either at a population level (Minagawa et al., 1986; Schoeller et al., 1986; Minagawa, 1992; Thompson et al., 2011; Valenzuela et al., 2011) or on a household basis (Yoshinaga et al., 1996). A few studies have compared individuals’ isotopic values to self-reported dietary records (Petzke et al., 2005a; Hedges et al., 2009; Huelsmann et al., 2009; O'Brien et al., 2009; Nash et al., 2012). Most studies of humans have used hair keratin, and some have used blood proteins (RBC, plasma, serum). Some short term feeding studies have measured other samples (such as urine and feces: Kuhnle et al., in press).

A significant problem with controlled diet isotopic studies is that of tissue turnover rates. When measuring the $\Delta^{15}N_{\text{diet-body}}$ the tissues usually of interest (e.g., bone collagen, hair keratin, blood proteins) isotopically reflect medium or long-term diet (months or years), so that a short-term dietary intervention study is not possible, due to issues with tissue turnover and isotopic equilibration (Jones et al., 1981; Tieszen et al., 1983; O’Connell and Hedges, 1999a; Ayliffe et al., 2004; Huelsmann et al., 2009; Petzke and Lemke, 2009). This has long been recognized, and all robust published controlled animal feeding studies are of animals raised on a single diet over a long time period of several years, if not a lifetime. Such a study is not ethically or practically possible in humans.

Here we report isotopic analyses from humans on known and controlled diets for a short period, where the controlled diets were designed to match each individual’s habitual diet, thus reducing problems with short-term changes in diet causing isotopic changes in the body pool. We measured dietary intake and a body tissue sample, red blood cells (RBCs).

**MATERIALS AND METHODS**

Samples were collected from healthy subjects taking part in a 30-day dietary intervention study to develop dietary biomarkers during the period of October 2002 to June 2003. Participants were provided with their habitual diet under controlled conditions for 30 days; blood samples and duplicate diets were collected. Details of the study protocol can be found in Tasevska et al., (2005, 2006). The study was approved by the Cambridgeshire Local Research Ethics Committee (LREC No 02/232) and all participants gave their full informed written consent. Samples were archived in a controlled storage facility (Fisher Bioservice, Bishop’s Stortford, UK) at −80°C for RBC and −20°C for all other specimens, and analyzed for this study in 2009–2010.

**Subjects**

A total of 13 healthy subjects from Cambridgeshire, UK, were recruited with advertisements. All participants were medically examined before the beginning of the study, including an assessment of the individual’s past and family medical history, details of recent and current medications, vitamin supplements, and tobacco/alcohol intake, and a cardiovascular examination. Blood analysis of fasting plasma glucose and glycated hemoglobin (HbA$_{1c}$) was undertaken and all subjects were within the normal range (fasting plasma glucose <6.1 mmol/l, HbA$_{1c}$ <6%). For this study, only samples from 11 participants (five males and six females, aged 23–66 y (59.7 ± 14.7 y), with a mean BMI of 25.8 ± 4.6 kg/m$^2$; Table 1) were suitable, as the 30-day study period for the remaining two was not continuous (a brief break for Christmas).

**Study design**

For the duration of the study, participants lived in the volunteer suite of the MRC Dunn Human Nutrition Unit (Cambridge, UK), where all food provided was prepared by trained technicians, and all specimens collected and processed. Participants followed their normal daily routine but were only allowed to consume foods prepared by the diet technicians. Subjects weighed themselves daily on an electric balance without shoes and in light clothing and recorded their body weight in the study diary. Physical activity was assessed using a questionnaire validated by the EPIC study (Wareham et al., 2003). Physical activity was recorded in the study diary on a daily basis as time (minutes) engaged in different type of exercise. A four-level score (inactive, moderately inactive, moderately active, and active) was assigned by combining occupational physical activity together with time participating in higher-intensity physical activities such as cycling, aerobics, swimming, jogging, exercising at a gym on a regular basis, etc.

**Diets**

Prior to the study, participants were asked to keep 7-day food diaries for 4 weeks while living at home. Weekly interviews with one of the investigators provided additional information, such as brand names. These data were used to replicate the habitual diet of each participant for the duration of the study. From approximately two-and-a-half times the amount of food expected to be eaten by the participant, one-half was prepared and one-half was kept for the preparation of a duplicate meal. The prepared half was weighed to the nearest gram, labeled with the name and the day, and left in a separate refrigerator for each individual. During the day, participants helped themselves and returned the uneaten food to the containers in the refrigerator. The next day, the
uneaten food was weighed out and the amount of food eaten was calculated.

Dietary intake was calculated from the UK food-composition tables using DINER (Data Into Nutrients for Epidemiological Research) (Welch et al., 2001). Tea and coffee were consumed freely during the course of the study, but participants were asked to keep their intake consistent and estimated intake was included in the data analysis. Five of the participants occasionally consumed alcohol; as this was not permitted in the volunteer suite, participants consumed alcohol outside the premises and recorded amount and type. The calculated dietary intake for alcoholic drinks was also added into the consumption data obtained in the study.

Duplicate diets were prepared daily for each participant. All food and drink items (excluding coffee, tea, alcoholic drinks, water, added salt, and pepper) were weighed to the nearest 1 g, chopped up and crushed, mixed with a weighed amount of boiling deionized water, and homogenized with a Magimix 5100 automatic food processor, usually for 10–15 min, until a smooth emulsion was obtained. Aliquots of each duplicate were stored at –20°C for analysis.

Blood collection, handling, and storage

Blood was sampled twice from each subject, at the start and in the last week of the study, by a trained phlebotomist. For one subject (V12), only blood collected at the end of the study was available for analysis. Fasting venous blood was collected into 10 ml lithium heparin monovettes. Within 1 h, the monovettes were centrifuged, the red blood cells removed from below the LiHep beads, washed thrice in chilled physiological solution, and then stored at –80°C prior to analysis.

Isotopic analyses

Duplicate diet samples were analyzed as liquid homogenates representative of 24 h food intake for each individual’s diet. Eight to twelve days’ diets were analyzed per subject, from the last half of the study. Samples were lyophilized and weighed into tin capsules (0.8 mg per aliquot). Red blood cell samples (0.2 ml) were lyophilized and then weighed into tin capsules (0.8 mg per aliquot). Diet samples were isotopically analyzed in duplicate, while blood samples were run in triplicate.

Isotopic analyses were performed using a Costech (Valencia, CA) automated elemental analyzer coupled in continuous-flow mode to a Thermo Finnigan MAT253 (Bremen, Germany) mass spectrometer at the Godwin Laboratory, Department of Earth Sciences, University of Cambridge. Stable isotope concentrations are measured as the ratio of the heavier isotope to the lighter isotope relative to an internationally defined standard, AIR (Hoefs, 1997). Isotopic results are reported as 15N values in parts per 1000 or “permil” (‰) values, where

\[ \delta^{15}N = \left( \frac{^{15}N_{\text{sample}}}{^{15}N_{\text{standard}}} - 1 \right) \times 1000. \]

Based on replicate analyses of international and laboratory standards, measurement errors are less than ±0.2‰ for δ15N.

Statistical analysis

Because of the sample size and distribution of the data, nonparametric tests were conducted to investigate differences. The main objective of this study was to investigate differences in δ15N between diet and blood;
assuming a standard deviation of 10% (higher than observed in this study) and a sample size of 11, changes of 15% can be detected with a power (1-β) of 0.9 at a significance level of α = 0.05. Power calculations were performed with G*Power 3.1.2 (Faul et al., 2009). Data analyses were conducted using Stata 11.2 (Statacorp, College Station, TX). The bivariate boxplot (bagplot: Rousseeuw et al., 1999) was prepared in R 2.12.1 (Team, 2009). Unless indicated otherwise, data are given as mean ± standard deviation.

**RESULTS**

Results are shown in Table 1. Overall, the body weight remained constant throughout the study (75.6 ± 15.7 kg at start vs. 75.8 ± 15.6 kg at end; Wilcoxon signed rank test, P = 0.56) which suggests that the intake achieved in the study was a valid reflection of the usual dietary habits in these volunteers. Weight changed by less than 2% in 10 participants; in one participant, the weight increased from 63.1 kg to 64.8 kg. However, this can be explained by normal fluctuations in the body weight, and changes in activity patterns during the study. Thus we take this population as being in a good approximation to steady state. True steady-state conditions are rarely achieved in free-living individuals, because abrupt changes in nitrogen balance occur from day to day, related to changes in dietary intake. Net accumulations and loss in nitrogen can be as much as ±2SE for free-living individuals, largely due to day-to-day variations in dietary nitrogen intake which can take several days to be reflected in excreted nitrogen (Bingham and Cummins, 1985). Of the 11 subjects, three of the subjects were physically inactive, three moderately inactive, four moderately active, and one active. They mostly practiced cycling, swimming, exercising at the gym, and jogging.

The median diet nitrogen isotopic value for all subjects was 4.7% (range in subject medians of 4.3–5.2%). The mean diet nitrogen isotopic value for all subjects was 4.8 ± 0.4% (range in subject means of 4.4–5.5%). We investigated whether daily variation in dietary nitrogen content would affect the average dietary nitrogen isotopic value for each subject, since individuals did not consume the same amount of protein on each of the 30 days of the study. For nine of the subjects, the difference between the arithmetical mean δ¹⁵N and the mean δ¹⁵N of each subject’s diets weighted by the nitrogen contribution from each day’s diet was less than 0.1‰, and for two individuals, the difference was less than 0.2‰; overall there was no statistically significant difference (Wilcoxon signed rank test, P = 0.37) between the two means (Table 1), so we consider that varying nitrogen intake had little if any quantifiable effect. Total protein intake and total nitrogen intake were inversely correlated with diet δ¹⁵N, although this correlation was only marginally significant (Spearman rank correlation: ρ = −0.59, P = 0.05, and ρ = 0.57, P = 0.07, respectively).

The range of BRC nitrogen isotopic values for all subjects was 7.6–8.9% at the start of the study and 7.4–8.8% at the end of the study. The median δ¹⁵N for all subjects was 8.2% (IQR = 7.9–8.6%) at the start of the study, and 8.1% (IQR = 8.0–8.4%) at the end of the study; the mean δ¹⁵N for all subjects was 8.3 ± 0.5‰ at the start of the study, 8.2 ± 0.4‰ at the end of the study, and 8.2 ± 0.4‰ for the two values averaged. Comparison of the δ¹⁵N of blood taken at the start and end of the study shows a small decrease (comparison possible for 10 of the 11 subjects; median difference = −0.1%, Wilcoxon test, P = 0.02; Table 1).

The overall difference between blood RBC and diet δ¹⁵N (Δδ¹⁵N) in the population can be calculated in several ways, depending on whether the mean or median for the population is used (Table 2). The range of individual Δδ¹⁵N is between 2.7 and 4.4‰, whichever way is used, and the average Δδ¹⁵N for the group is between +3.3 and +3.6‰, with the statistically most parsimonious value (using the final blood sample δ¹⁵N and the median diet δ¹⁵N of +3.5‰). We did not observe any statistically significant difference between men and women, and no significant correlation with age or physical activity. The study was carried out over a period of months, but the sample size was too small to investigate the possible effects of seasonal changes in metabolic activity. However, Δδ¹⁵N—correlated significantly with BMI (Spearman rank correlations, respectively: ρ =

<table>
<thead>
<tr>
<th>Subject</th>
<th>Arith mean δ¹⁵N (‰)</th>
<th>Median δ¹⁵N (‰)</th>
<th>Blood 2 δ¹⁵N (‰)</th>
<th>Mean δ¹⁵N (‰)</th>
<th>Δδ¹⁵N (blood - mean diet) (‰)</th>
<th>Δδ¹⁵N (blood - median diet) (‰)</th>
<th>Δδ¹⁵N (blood - median RBC) (‰)</th>
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<tr>
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<td>3.3</td>
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<tr>
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but that range of diets for several months. Therefore we suggest that the
using the mean bloods taken at the end of the study may be an overesti-
jects with BMI correlations are non-significant, thus it is possible that
If we exclude those who are obese (BMI $\geq 30$), all three
the two identified outliers are shown in gray.

The assumption underlying the premise of this study
is that the controlled diet consumed by subjects over the
30-day study was isotopically similar to their habitual
diets. The study for which these samples were collected
was not designed as an isotopic study, so no considera-
tions were made of isotopic variability in foods. However,
the diets were carefully designed so as to match the com-
position of habitual diets, including the matching of
brands consumed. A small but significant average
decrease of 0.1\% in $\delta^{15}N_{RBC}$ suggests that the study
diets were not isotopically identical to habitual diets
(bearing in mind that each subject’s study diet was spec-
tic to them, so some may have been different and
others not). Red blood cells have a mean in vivo life span
of 120 days (Landaw, 1991), so a median change of
$-0.1\%$ in $\delta^{15}N_{RBC}$ over the duration of the 30-day study
suggests that there could be a median difference of
$-0.4\%$ over 120 days. Thus the measured $\delta^{15}N_{RBC}$ of
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$+3.6\%$ (Table 2), should be expanded to $+3.0$--$3.6\%$, but
that $\Delta^{15}N_{diet}$ is highly likely to be larger than
$+3.0\%$. For the further discussion in this paper, we use
the value of $+3.5\%$, based on the most parsimonious
value of $\Delta^{15}N_{diet-RBC}$, with the recognition that it may be
a slight overestimate.

Studies have shown that isotopic differences between
diet and animal tissues can vary under different condi-
tions (e.g., Ambrose and DeNiro, 1986; Heaton et al.,
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here will not be universally constant for all humans on
all diets. However, this is the first quantified isotopic
study of the diet to body enrichment in humans on con-
trolled diets, and therefore gives an indication of the
magnitude of the offset that we can expect. We found no
effect of sex or age on $\Delta^{15}N_{diet-body}$ offset in these
subjects. The observed positive correlation with BMI, driven
by the two obese subjects, is intriguing and requires fur-
ther investigation: the possibility of an effect of differen-
tial bioavailability of nutrients and differential uptake
between individuals may be a factor here, and one that
should be considered further.

**DISCUSSION**

The assumption underlying the premise of this study
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effect of sex or age on $\Delta^{15}N_{diet-body}$ offset in these
subjects. The observed positive correlation with BMI, driven
by the two obese subjects, is intriguing and requires fur-
ther investigation: the possibility of an effect of differen-
tial bioavailability of nutrients and differential uptake
between individuals may be a factor here, and one that
should be considered further.

**Offsets from diet to keratin and collagen**

To be able to use this measured diet-body offset for
humans in palaeodietary studies, we must estimate
what it equates to in terms of tissues analyzed in other
studies, such as keratin or collagen. We can combine our
data with that of three other studies, all on North Amer-
ican residents, to derive a value for $\Delta^{15}N_{diet-keratin}$ (Ta-
ble 3). Nash et al. (2009) showed a mean increase of
$+1.5 \pm 0.6\%$ from RBCs to hair keratin. Kraft et al.
(2008) showed that blood plasma has a higher $\Delta^{15}N$ than
red blood cells by 1.5\% on average. Schoeller et al.
(1986) showed a mean increase of +0.3 ± 0.7\% from
plasma protein to hair keratin. Combining the plasma/
RBC/keratin results from these two latter studies, we
get an estimated offset of +1.8\% from RBCs to hair ker-
atin, in fairly good agreement with the value of +1.5\%
observed by Nash et al. Our measured $\Delta^{15}N_{diet-RBC}$ value
of +3.5\% equates to a $\Delta^{15}N_{diet-keratin}$ of +5.0\% using the
Nash offset, and to +5.3\% using the Kraft-Schoe-
ller combined offset (no errors propagated).

Our derived $\Delta^{15}N_{diet-keratin}$ value can be compared to
estimates from two studies specifically examining the
offset from diet to hair keratin, based on estimates of di-
etary intake combined with food and hair isotopic anal-
ysis (Table 3). Yoshinaga et al. (1996) analyzed 49 males
dietary surveys, food isotopic analysis, and hair isotopic analysis, they derived an estimated value of +5.0–6.9\% for $\Delta^{15}N_{diet-keratin}$ based on a cal-
culated diet for each individual. Hedges et al. (2009) ana-
alyzed 20 females in Fiji sampled in 1999. Through diet
surveys, food isotopic analysis, and hair isotopic analysis, they derived an estimated value of +4.1 ± 0.7\% for
$\Delta^{15}N_{diet-keratin}$ based on a calculated diet for each individ-
ual. Our measured data with a combination of the Nash-
Jahren-Schoeller offsets gives an estimate of $\Delta^{15}N_{diet-ker-
atin}$ of +5.0–5.3\%, which falls between the estimated values
from Yoshinaga and Hedges. Studies estimating die-
tary intake at the population level have estimated a
$\Delta^{15}N_{diet-keratin}$ of ca. +4.3\% (Minagawa et al., 1986;
Schoeller et al., 1986).
Table 3. Nitrogen isotopic values of tissues, diet and calculated diet-tissue offsets in published human studies (all given in units of %)

<table>
<thead>
<tr>
<th>Population</th>
<th>Sex</th>
<th>Hair δ¹⁵N</th>
<th>Bone δ¹⁵N</th>
<th>RBC δ¹⁵N</th>
<th>Plasma δ¹⁵N</th>
<th>Diet δ¹⁵N</th>
<th>Δ¹⁵Ndiet-collagen</th>
<th>Δ¹⁵Ndiet-keratin</th>
<th>Δ¹⁵Ndiet-RBC</th>
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<tr>
<td>Rural PNG</td>
<td>M</td>
<td>9.1 ± 0.5</td>
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<td>8.8 ± 0.3</td>
<td>8.8 ± 0.3</td>
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<tr>
<td>Radal PNG</td>
<td>M</td>
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<td>Wonie PNG</td>
<td>M</td>
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<td>2.0</td>
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<tr>
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To consider how our data would translate to a Δ¹⁵N_diet-collagen offset, we must then consider the offset between human hair keratin and bone collagen. Three published studies have measured this in humans, one in a modern population (+0.9 ± 0.2‰; O’Connell et al., 2001) and two in archaeological individuals (+1.0 ± 1.1‰; O’Connell and Hedges, 1999b; +1.0 ± 1.4‰; Richards, 2006) (Table 3). There are problems in using such data (such as the small sample sizes and the large standard deviations in the two archaeological studies) but it is noteworthy that all studies have similar mean offsets for the Δ¹⁵Nkeratin-collagen offset. Adding these corrections to the estimated Δ¹⁵N_diet-keratin of +5.0–5.3‰ derived from our data and the offsets measured by Nash/Kraft/Schoeller et al., we derive a range of +5.9–6.3‰ for the Δ¹⁵N_diet-collagen offset (again no errors propagated).

As we discuss earlier, the measured δ¹⁵N_RBC may be an overestimate, and thus the derived values of Δ¹⁵N_diet-keratin and Δ¹⁵N_diet-collagen may also be overestimated. Possible problems with studies comparing keratin to diet include issues with growth cycle errors (Williams et al., 2011). Problems with studies comparing collagen and keratin include differential time periods represented in the two tissues (O’Connell et al., 2001; Hedges et al., 2007). However, even with a very conservative approach, assuming a Δ¹⁵N_diet-RBC value of +3‰, and using minimum offset values to keratin (Nash study, +0.9‰, i.e., 1σ less than the mean), and to collagen (O’Connell 2001 modern study, +0.7‰, i.e., 1σ less than the mean), our results suggest a Δ¹⁵N_diet-collagen offset of +4.6‰, which is at the upper end of the currently accepted range. These data suggest therefore a larger offset than commonly assumed.

We can place the limited human data in the context of that from other animal studies. All controlled feeding studies on animals so far have observed isotopic inhomogeneity in different tissues, and such isotopic differences can be substantial (Caut et al., 2009). Other mammalian studies have shown a similar pattern to that summarized above for humans: whole blood and red blood cells generally have low nitrogen isotopic values relative to other tissues, or at the low end of the range, and in comparisons of plasma and red blood cells, plasma always has a higher nitrogen isotopic value, often by more than 1‰ (Table 4). As regards the magnitude of the offsets, similar values to our estimates are found for a range of species in the literature. A number of animal studies have found Δ¹⁵N_diet-body differences of greater than 4‰ for a variety of tissues (DeNiro and Epstein, 1981; Hilderbrand et al., 1996; Roth and Hobson, 2000; Sponheimer et al., 2003; Arneson and MacAvoy, 2005; Miron et al., 2006; Caut et al., 2008), and studies of goat, alpaca, seal and bear have shown differences larger than 5‰, up to 6.4‰ (Kurle, 2002; Felicetti et al., 2003; Sponheimer et al., 2003).

Implications of this study for palaeodietary work

Overall, our data suggest that the Δ¹⁵N_diet-collagen offset in this group is ca. +6‰, larger than that usually assumed in the archaeological literature, typically around +3.5‰ (Bocherens and Drucker, 2003). Using a very conservative approach to the data, the estimate is still ca. +4.6‰, at the upper end of the currently accepted range. Such an observation has implications for the interpretation of human palaeodiet from isotopic data: an underestimation of the Δ¹⁵N_diet-collagen offset
will lead to an overestimation of the dietary importance of foods with higher nitrogen isotopic values, usually higher trophic level foods such as meat, milk and fish. As Hedges and Reynard (2007) note, using a \( \Delta^{15}N_{\text{diet-collagen}} \) value of 3-4% produces an estimate of dietary animal protein percentage (as a proportion of total protein intake) of 60% and sometimes up to 80% for prehistoric farmers in Europe, which is greater than animal protein dietary fraction of modern “developed” countries and twice that of modern “developing” countries (Sluijs et al.; Fras- setto et al., 2000; FAOSTAT, 2012), as well as being in excess of that consumed by most ethnographically documented hunter-gatherer populations (Cordain et al., 2000). If a value of +6% were used as \( \Delta^{15}N_{\text{diet-collagen}} \) offset, this would typically reduce the dietary animal protein intake estimate by about a third to a half, bringing such estimates for prehistoric farmers in line with dietary animal/plant protein ratios in living horticultural/ agricultural populations (Yoshinaga et al., 1996; Fras setto et al., 2000; Maclntyre et al., 2002; Muhammad Lawal and Balogun, 2007; Hedges et al., 2009; Iyangbe and Orewa, 2009; Baroudi et al., 2010).

**CONCLUSIONS**

In 11 subjects consuming their habitual diets under controlled conditions, we have measured the \( \Delta^{15}N_{\text{diet-RBC}} \) as +3.5%. This is the first study to measure the \( \Delta^{15}N_{\text{diet-body}} \) offset in humans on controlled diets of known isotopic composition. Using measured offsets from other studies, we estimate the human \( \Delta^{15}N_{\text{diet-keratin}} \) as +5.0–5.3%, which is in good agreement with estimates derived from the two other studies using individual diet records (Yoshinaga et al., 1996; Hedges et al., 2009). We also derive a value for \( \Delta^{15}N_{\text{diet-collagen}} \) of ≈6%, larger than usually assumed in palaeodiets studies. This larger value goes some way to resolving the conundrum of interpretations of very high animal protein intake in isotopic studies of prehistoric farmers—we suggest that this has often been overestimated. We advocate that dietary interpretations of previously published archaeological human isotopic data are reconsidered in the light of our work.

**ACKNOWLEDGMENTS**

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**LITERATURE CITED**


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