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Eneolithic subsistence economy in Central Italy: first dietary reconstructions through stable isotopes

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Abstract

The paper aims to point out the subsistence in Eneolithic Central Italian communities by Stable Isotope Analysis. This period marked a tipping point in the food strategies because it was characterized by economic changes and several technological improvements leading to enhance land exploitation and livestock breeding. Carbon and nitrogen stable isotope analysis has been used to analyze the food consumption of 54 people belonging to 5 Eneolithic communities scattered throughout Central Italy, where no data have yet been published. The estimation of the main protein intake has been achieved in order to quantify the differences among these communities. The results are consistent with a diet mainly based on terrestrial resources, with no exclusive marine sources consumption, although their occasional usage cannot be ruled out, especially for selected funerary contexts. The data suggest an overall subsistence based on greater local resource procurement, supported by regional productivity maximization. A roughly homogeneous landscape could be outlined in Tuscany and Marche communities witnessing a shared diet preference that could be modified by local preferences. The fully developed trade routes between the two sides of the Apennines could address the overall dietary homogeneity of the studied communities, especially between Fontenoce di Recanati and the southern Tuscan human groups such as Grotta del Fontino and Buca di Spaccasasso, with lesser influence for Le Lellere

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and Podere Cucule that seem to suggest a more locally based subsistence, even though the funerary affinities do not match this overall diet homogeneity.

Keywords Copper age \cdot Carbon and nitrogen stable isotopes \cdot Diet reconstruction

Introduction

Copper Age scenario

The Copper Age (or Eneolithic) is a period ranging ca. 4000–2100 cal. BCE and despite the radiometric dates are limited for each archeological context, the measurements shed light on the relationships between the Neolithic and Eneolithic (Manfredini et al. [2013](#page-15-0); Dolfini [2010;](#page-14-0) Leonini et al. [2013](#page-14-0); Cazzella et al. [2013](#page-13-0); Cocchi Genick [2014](#page-14-0)). The term Copper Age emphasizes the introduction of the first metal items and their use alongside lithic products: the period is also characterized by economic changes and several technological innovations leading to improved land exploitation and demographic increases. Thus, the Eneolithic marks the tipping point for the transformations in human groups: the emergence of copper processing corresponds to a significant increase in trade, both for the retrieval of the raw material and for manufactured objects (Kostov [2005](#page-14-0)). This trade accounts for the appearance in Europe of metal objects and exotic items, such as ivory and ostrich eggs, imported from North Africa and the Middle East (Chapman [1990](#page-14-0); García Sanjuán et al. [2013](#page-14-0)). The human groups become more and more complex, creating a variety of funerary practices for selected people (Leonini and Sarti [2006](#page-14-0); Cazzella and Guidi [2011](#page-13-0); Recchia and Baroni [2011\)](#page-15-0), and somewhere the funerary practices have linked to the megalithic structures, especially in Western European regions (Rubinetto et al. [2013](#page-15-0)). The Copper Age is also characterized by innovations in productive activities: livestock breeding increases, resulting into what has been called "The Secondary Products Revolution" (Greenfield [2013](#page-14-0)). This phenomenon is linked to changes in faunal exploitation: domestic animals were initially used solely for meat consumption, but in the Copper Age this management mode is replaced by the optimizing of their secondary products or applications. Secondary animal products can be obtained without slaughtering the animals, and the same animal can be repeatedly exploited over its lifetime (Greenfield [2010](#page-14-0)): thus, domestics are used for milk, wool, and textile productions. These products enable a significant shift in the economic development of Copper Age communities: the increase in both hunting and productivity leads to mobility, resulting in the occupation of a range of several environments (Sherratt [1993](#page-15-0)). The wide use of cattle as traction animals enables the cultivation of more land and the transportation of products for greater distances. These factors allow for the accumulation of wealth, and consequently for social differentiation (Champion et al. [2009\)](#page-13-0).

Copper Age in Italy

In Italy, the Copper Age exhibits itself in some cultural fractions (facies) having in common the use of natural cavities as funerary areas, though the sites may be manipulated (Leonini and Sarti [2006](#page-14-0)). These funerary spaces are distributed in all the peninsular areas, where complex funerary practices are scattered (artificial and natural caves, open air spaces, and tomb enclosures could be addressed). Even the rituals start to be more complex with the coexistence of single and multiple burials as well as violations or disturbance for secondary burial and several funerary goods left to the people. Though several archeological researches have been conducted in Italy about Copper Age communities (Martini [2006](#page-15-0); Barfield et al. [2010;](#page-13-0) Cazzella and Guidi [2011\)](#page-13-0), few studies have broadened the knowledge about how the communities used their alimentary resources, enabling an approach to the socioeconomic organization of human communities. Archeological human remains contain a wealth of information about not only the biology but also the cultural adaptations of past populations, including their dietary supply. It should be noted that to reconstruct the lifestyle of prehistoric communities, human bone analysis is sometimes the only available material and without it a holistic interpretation is not possible.

Thus, in the present paper, a bio-molecular approach based on stable isotope analysis (Katzenberg [2008\)](#page-14-0) will be used to analyze the food consumption of several Eneolithic communities scattered throughout Central Italy, where to the best of our knowledge no data have yet been published. The samples have been chosen to analyze putative cultural connections between the western and eastern regions in central Italy, as previously suggested for archeological records (Sarti [2005\)](#page-15-0): dietary patterns could be hypothesized as one of the most retained markers of the people and societies' cultural identity. This bio-molecular approach could support an anthropological point of view about an overall homogeneity of Copper Age populations, likely a consequence of extensive gene and cultural flows among human groups (Borgognini Tarli [1992](#page-13-0); Vargiu et al. [2009\)](#page-15-0).

Contexts: archeological areas

The Eneolithic sites selected for the present study are scattered throughout Central Italy and are mainly located in Tuscany and Marche: Fontenoce di Recanati (Marche), Podere Cucule, Le Lellere, Buca di Spaccasasso, Poggialti Vallelunga, and Grotta del Fontino (Tuscany) (Fig. 1). These funerary areas have been archeologically and chronologically evaluated (Silvestrini et al. [1993](#page-15-0); Pacciani [1993;](#page-15-0) Cencetti and Pacciani [1994](#page-13-0); Chilleri and Pacciani [2002;](#page-14-0) Vigliardi [2002](#page-15-0); Volante [2014\)](#page-15-0), but currently lack a bio-molecular characterization of the human remains that have been recovered there. The burials are extremely heterogeneous in every area, and the chronology of the funerary assets is reported in Table [1.](#page-3-0)

The selected samples pertain to not ritually homogeneous funerary contexts: it is well known that collective burial caves such as Grotta del Fontino and Buca di Spaccasasso hosted a huge number of bone remains in secondary burial deposition, while in rock cut tombs (for instance in Fontenoce di Recanati) the skeletons are found in primary deposition mode.

Stable isotopes background

The analysis of carbon and nitrogen stable isotopes of bone collagen over the last 35 years has been extensively used to study the diets of ancient communities, especially concerning their protein intake (Braun et al. [2013](#page-13-0)). Indeed, modeling studies based on the results of controlled animal feeding experiments proposed relatively fixed contribution of carbon from protein and energy fractions of diet to collagen (Fernandes et al. [2012](#page-14-0)). The method is based on the estimation of the ratios of 13^1 C/ 1^2 C and 1^5 N/ 1^4 N, detectable from the bone collagen that reflect the diet concerning the last decades of life

Fig. 1 Geographical location of every sampled community

Table 1 Chronological radiometric dates for the considered samples (personal communications refer to not yet published data generated in MIUR research project "EPIC" ID: 2010EL8TXP)

(Ambrose [1990;](#page-13-0) Hedges et al. [2007\)](#page-14-0). The natural abundance of ¹³C and ¹⁵N is expressed as per mil (%) deviation from international standards: ${}^{\delta13}$ C or δ^{15} N ($R_{\text{sample}}/$ $R_{standard}$ – 1) × 1000, where R in δ^{13} C or δ^{15} N is 13 C/¹²C or $15N/14N$, respectively.

The carbon standard is represented by V-PDB (Vienna PeeDee Belemnite), while atmospheric nitrogen (ambient inhalable reservoir, AIR) (Mariotti [1983](#page-15-0)) is used as the nitrogen standard.

Dietary reconstruction is based on the concept that the carbon and nitrogen isotope values in consumers' bone collagen are higher than the corresponding values of their prey (Schwarcz and Schoeninger [2012](#page-15-0)). For this purpose, the carbon isotope composition (δ^{13} C) is suitable to determine the consumption of plants with different photosynthetic pathways $(C_3$ and C_4) (Vogel [1980](#page-15-0); Gannes et al. [1998](#page-14-0)), even though this marker is also useful in distinguishing terrestrial and marine sources (Walker and DeNiro [1986;](#page-15-0) O'Brien [2015](#page-15-0)), and freshwater ecosystems are also identified because they often display lower δ^{13} C values than C_3 terrestrial plants (Dufour et al. [1999](#page-14-0); Doppler et al. [2011\)](#page-14-0). Additionally, the difference in values between two species at adjacent trophic levels is about 1‰ (Bocherens and Drucker [2003](#page-13-0); Lee-Thorp [2008](#page-14-0)).

The nitrogen signature $(\delta^{15}N)$ provides information on overall trophic levels because an average growth between 3 and 5‰ in the food net is observed (Tykot [2004](#page-15-0); Hedges and Reynard [2007](#page-14-0)). Thus, it is possible to detect low levels in plants, and greater levels in herbivores, omnivores, and carnivores (DeNiro and Epstein [1978](#page-14-0)). Moreover, the marine ecosystem sources could be ascertained through nitrogen stable isotope data: this phenomenon also accounts for the marine denitrification process that is associated with the fractionation of the heavy isotope and with the presence of 15 N-rich compounds (Dahnke and Thamdrup [2013](#page-14-0)).

Materials and methods

Sample

A total of 75 human bones were analyzed for the carbon and nitrogen stable isotopes. These samples reported in Table [2](#page-4-0) represent the whole human remains sample recovered in each funerary contexts. Despite the sample size in each community is heterogeneous and some of them seem to be trivial, they represent the first cumulative sample of Eneolithic Italian populations and its isotopic characterization comes to fill a critical gap for the region and the Eneolithic timeframe bringing new and reliable evidence regarding prehistoric human populations in the Italian peninsula. For each funerary context, information on sex and age at death were partially available from previous studies (Silvestrini et al. [1993;](#page-15-0) Pacciani [1993;](#page-15-0) Cencetti and Pacciani [1994;](#page-13-0) Chilleri and Pacciani [2002;](#page-14-0) Vigliardi [2002;](#page-15-0) Volante [2014](#page-15-0)).

The baseline for the terrestrial protein component in the diet has been accounted for through nine faunal remains from Fontenoce di Recanati in the Marche region: three are bone fragments of Canis familiaris, two are of Sus scrofa, and four are related to unknown herbivore species found in association with human skeletal remains.

No faunal remains were available for the Tuscany area, so we considered the values from a neighboring area (Latium) for a roughly coeval site (Casetta Mistici, in the area around Rome, De Angelis et al. [2016](#page-14-0)) that should be consistent with faunal values from Fontenoce di Recanati, supporting the use of the latter as reference ecologic data, while the baseline from the aquatic resources was estimated using information in literature related to different prehistoric time frames due to the lack of local fishbone findings (Richards and Hedges [1999;](#page-15-0) Drucker and Bocherens [2004](#page-14-0); Herrscher and Le Bras-Goude [2010](#page-14-0)).

Analytical methods

The extraction of collagen was performed following the Longin's protocol [\(1971](#page-15-0)) modified by Brown et al. [\(1988\)](#page-13-0). In addition, the extraction was performed simultaneously on a modern bovine sample used as a reference. To obtain a

Table 2 Samples sizes and
references

Table 2 Samples sizes and references	Funerary context	Geographic area	N	References	
	Poggialti Vallelunga	Tuscany		Chilleri and Pacciani (2002)	
	Podere Cucule	Tuscany		Pacciani (1993)	
	Le Lellere	Tuscany	8	Cencetti and Pacciani (1994)	
	Buca di Spaccasasso	Tuscany	4	Volante (2014)	
	Grotta del Fontino	Tuscany	25	Vigliardi (2002)	
	Fontenoce di Recanati	Marche	32	Silvestrini et al. (1993)	

satisfactory yield of collagen, the extraction must be performed on about 500 mg of bone powder. The ultrafiltration step, using > 30 kDa Amicon® Ultra-4 Centrifugal Filter Units with Ultracel® membranes (Millipore), was also performed for samples showing a low preservation appearance, in order to maximize the collagen concentration.

Each collagen extract was weighed 0.8–1.2 mg and analyzed in triplicate using continuous-flow isotope ratio mass spectrometry (CF-IRMS) in IGAG (Istituto di Geologia Ambientale e Geoingegneria, CNR) facilities.

To test reliability and exclude contamination from exogenous carbon and nitrogen sources, the samples were compared against established criteria to ascertain the percentages of carbon and nitrogen, atomic C/N ratios, and collagen yields (Ambrose [1990;](#page-13-0) DeNiro [1985;](#page-14-0) van Klinken [1999\)](#page-15-0). Analytical precision is $\pm 0.2\%$ for δ^{13} C, reported with respect to the PDB standard, and $\pm 0.3\%$ for δ^{15} N, reported with respect to AIR.

Past v.3.14 software (Hammer et al. [2001](#page-14-0)) was used to perform descriptive statistics and comparison tests (the Mann-Whitney U test and the Kolmogorov-Smirnov test).

The linear mixing model firstly proposed by Fraser et al. ([2013\)](#page-14-0) and recently developed also by Fontanals-Coll et al. ([2016](#page-14-0)) has been employed in order to quantify the fraction of animal protein exploitation out of total dietary protein consumption. As stated by the authors, this model quantifies the percentage of animal protein intake, starting from the midpoint of the usual $\delta^{15}N$ offset of + 3‰ and + 5‰ between consecutive trophic levels. Thus, the information based upon the plant, faunal, and human δ^{15} N values could be used to build up several models based on dissimilar trophic chains lying on selected resources δ^{15} N values. These models could proficiently use the plant δ^{15} N values to infer the faunal (herbivore) ones as well as the recorded herbivore data might be significant in achieving sympatric plants values: plant and herbivore values determine the starting point for the esteem of the percentage of animal protein in human diet. Specifically, a first attempt for estimating the animal protein fraction (percentage) of total dietary protein was performed starting from the herbivore $\delta^{15}N$ values recorded in Fontenoce di Recanati: the herbivore average $\delta^{15}N$

value allows to infer the plant $\delta^{15}N$ value by subtract 4‰. Afterwards, several attempts were focused on the single plant group contribution in food web and in the definition of the starting point of the model: the significance of wheat, barley, and pulses was evaluated, whose increasing of 4‰ allows to account for the herbivore $\delta^{15}N$ average value that is due to 0% animal protein–based diet. This δ^{15} N magnification was iteratively performed to detect the average human $\delta^{15}N$ value whether their diet would be totally due to herbivore-derived protein (100% animal protein–based diet). The interception of the line connecting herbivore- and human-estimated values with the real δ^{15} N values of each community matches their estimated percentage of animal protein in the diet. Unfortunately, few archaeobotanical investigations have been carried out in these areas, despite the fact that numerous archeological surveys have been performed. To the best of our knowledge, no isotopic data are currently available on coeval domestic plants in the considered areas, making hard the application the proposed model with local plant resources. Therefore, the plant δ^{15} N values to draw up the model come from Vaiglova et al. [\(2014](#page-15-0)), where floral and faunal isotope data have been determined for reconstructing early farming practices at

Results

modeling.

Collagen quality indicators

The collagen extraction was performed for all the samples, but the yield of the organic component was

Kouphovouno, a Late Neolithic village in southern Greece (ca. 5950–4500 cal. BC): this is the best time period and topographical comparison that could be used to constrain the "regional" plant isotopic features. Furthermore, a restricted sample of actual crops (5 grain samples) from central Italy has been isotopically analyzed by Brescia et al. ([2002](#page-13-0)) so as these data were recruited in IsoArcH repository (Salesse et al. [2018](#page-15-0)) to be used for properly obtained for only a fraction of the human bones (Table 3).

The results of the analysis of the human bones for carbon and nitrogen isotopes are listed in Table [4.](#page-6-0)

The C:N ratio and percent collagen yield are listed as an indication of the reliability of the δ^{13} C and δ^{15} N sample measurements. In order to assess the preservation state of the extracted collagen, different criteria were used: carbon content greater than or equal to 30%, nitrogen content greater than or equal to 10% (Ambrose [1990\)](#page-13-0), and C/N ratios between 2.9 and 3.6 (DeNiro [1985](#page-14-0)). Data from collagen with an elemental composition below these criteria and C/N ratios outside this range were excluded. The extraction yield was not used as a criterion (Ambrose [1990](#page-13-0)) because the ultrafiltration technique was used: only samples with yield of 0% were ruled out.

The faunal remains from Fontenoce di Recanati yielded enough collagen to be analyzed in five cases (Table [5\)](#page-7-0): a bone fragment of Canis familiaris, two pig samples, and two unidentified herbivore fragments. The obtained faunal $\delta^{13}C$ values are consistent with a C_3 European ecosystem (Schwarcz and Schoeninger [1991](#page-15-0)) where the $\delta^{15}N$ signature suggests the proper trophic level for the identified species. The compared faunal samples from Latium (Casetta Mistici, Latium, De Angelis et al. [2016\)](#page-14-0) were proven to be consistent with values from Fontenoce di Recanati, supporting the use of the latter as ecological reference data.

The higher trophic level of the humans (compared to the fauna) ensures quality results and suggests that the livestock might be considered prey stock for the humans.

Out of 75 samples, only 54 fitted the quality criteria (72%). Remarkably, Poggialti Vallelunga does not allow us to perform the analytical process due to the lack of suitable extracted collagen.

Isotopic results

Taking all 54 human individuals (Table [4\)](#page-6-0), δ^{13} C ranges from -17.5 to -21.2% , whereas δ^{15} N values are between 6.9 and 11.6‰. The general distribution indicates a certain

Table 3 Samples suitable to be evaluated by stable isotope analysis

Funerary context	N	Samples with satisfactory collagen yield	%
Poggialti Vallelunga	3	0	θ
Podere Cucule	3	3	100
Le Lellere	8	8	100
Buca di Spaccasasso	4	4	100
Grotta Fontino	25	24	96
Fontenoce di Recanati	32	15	47
Total	75	54	72

heterogeneity (Fig. [2\)](#page-8-0) both in the δ^{13} C and δ^{15} N values, which have a range of more than 4‰.

The sample stratification according to the site allows us to evaluate putative differences in food exploitation, even though they do not reach the level of statistical significance (coupled Mann-Whitney *U* test, Table [6](#page-8-0)).

The mean δ^{13} C and δ^{15} N for the five funerary areas were estimated (Table [7,](#page-9-0) Fig. [3\)](#page-9-0).

The values are consistent with a diet mainly based on terrestrial resources, and all the samples rely on a higher trophic level than the faunal samples, with no clear indication of exclusive marine source consumption. However, their occasional usage cannot be ruled out, due to the high $\delta^{15}N$ values in Buca di Spaccasasso and Grotta del Fontino ($\delta^{15}N = 9.6\%$) and their not-so-negative δ^{13} C data.

The previously accomplished anthropological evaluation of the remains (Silvestrini et al. [1993](#page-15-0); Pacciani [1993;](#page-15-0) Cencetti and Pacciani [1994;](#page-13-0) Chilleri and Pacciani [2002;](#page-14-0) Vigliardi [2002](#page-15-0); Volante [2014](#page-15-0)) allowed us to determine the age class of all individuals except for those at Grotta del Fontino, where no anthropological evaluation is available. This information led us to dissect the variability in food consumption between adults and children in Fontenoce di Recanati (8 adults and 7 children) and Podere Cucule (2 adults and 1 child).

The mean values of the infants in Fontenoce di Recanati $(\delta^{13}C = -19.7\%$ and $\delta^{15}N = 9.4\%$) are quite similar to the adult ones (δ^{13} C = − 19.4‰ and δ^{15} N = 8.9‰), suggesting a diet similar to that of the adult population, which means that in childhood no dietary differences caused by physiological growth-related processes (breastfeeding/weaning) can be ascertained (Mann-Whitney U test for $\delta^{13}C$, U = 95.5; Zscore = 0.67; $p > 0.05$ and Mann-Whitney U test for $\delta^{15}N$, $U = 101.5$; Z-score = 0.42; $p > 0.05$). Instead, despite the uniqueness, the sole child in Podere Cucule shows a lower δ^{15} N signature than the adults (7.9‰ vs 9.5‰), suggesting different food consumption, even though the δ^{13} C is quite similar (−18.4‰ vs −18.1‰). No differences could be ascertained between males and females in Le Lellere (3 females and 5 males) and Fontenoce di Recanati (4 females and 4 males) although the limited sample size could affect the reliability of this estimation (Le Lellere males/females δ^{13} C $U = 30$, Z-score = 0.05; $p > 0.05$; δ^{15} N $U = 22$, Z-score = 0.81; $p > 0.05$; Fontenoce di Recanati males/females $\delta^{13}C$ U = 31, Z-score = -0.10 ; $p > 0.05$; $\delta^{15}N$ U = 30, Z-score = -0.10 ; $p > 0.05$).

Based on the model proposed by Fraser et al. ([2013](#page-14-0)), we tried to distinguish the fraction of animal protein from the total protein intake. It seems to be significantly high in all the considered sites (δ^{15} N mean value, 9.3‰) except for Le Lellere (δ^{15} N value, 7.6‰). If the plain observation leads to an interpretation of these values as due to large amounts of animal protein, this conclusion might be due

Table 4 Available basic demographic parameters (sex and age at death), stable isotope measurements, and collagen quality control indicators of humans. F, female; M, male; NA, not available; y, years old; m, months old

Site	Lab code	Sex	Age	$\delta^{13}C\,\%o$	$\delta^{15}{\rm N}$ ‰	$\%C$	%N	C/N	Collagen ratio
Podere Cucule	PC62	M	Adult	-17.9	10.0	43.9	15.7	2.9	4.1
Podere Cucule	PC63	F	Young adult	-18.4	9.0	43.0	15.5	2.9	3.9
Podere Cucule	PC64	NA	Child $13 - 14y$	-18.4	7.9	43.0	15.3	3.0	1.2
Le Lellere	L65	M	Adult	-20.5	7.1	31.2	10.0	3.1	1.6
Le Lellere	L66	M	Adult	-20.2	7.9	37.3	12.8	2.9	1.1
Le Lellere	L67	F	NA	-20.2	7.5	40.1	12.0	3.6	1.3
Le Lellere	L68	M	Adult	-19.5	6.9	41.4	13.5	3.6	5.9
Le Lellere	L69	M	Adult	-19.6	8.0	39.6	12.6	3.3	1.7
Le Lellere	L70	F	Adult	-19.5	7.6	41.6	13.2	3.4	2.4
Le Lellere	L71	M	Adult	-19.8	7.9	39.6	12.4	3.2	1.3
Le Lellere	L72	F	Adult	-19.8	8.4	43.9	11.8	3.5	0.2
Buca di Spaccasasso	S1	M	Adult	-20.7	9.9	50.4	11.6	3.7	4.3
Buca di Spaccasasso	$\ensuremath{\mathrm{S2}}$	NA	Adult	-17.5	9.8	42.1	14.8	3.2	8.7
Buca di Spaccasasso	S ₃	$\mathbf M$	Adult	-18.9	9.4	44.0	14.3	3.1	7.4
Buca di Spaccasasso	S4	M	Adult	-19.2	9.5	43.6	13.2	3.3	7.3
Grotta del Fontino	GF ₁	NA	NA	-19.3	9.7	42.3	13.8	3.1	1.0
Grotta del Fontino	GF ₃	NA	NA	-20.0	8.7	45.4	13.1	3.5	1.1
Grotta del Fontino	GF4	NA	NA	-19.5	9.5	45.2	13.4	3.4	2.4
Grotta del Fontino	GF 5	NA	NA	-18.9	10.2	41.7	15.0	3.0	3.8
Grotta del Fontino	GF ₆	NA	NA	-18.9	10.9	41.5	14.1	2.9	1.0
Grotta del Fontino	${\rm GF}$ 7	NA	NA	-17.8	9.4	43.4	15.0	2.9	2.2
Grotta del Fontino	GF 8	NA	NA	-19.9	9.9	46.1	13.4	3.4	3.1
Grotta del Fontino	GF 10	NA	NA	-18.5	11.0	41.9	14.8	2.9	0.2
Grotta del Fontino	GF 12	NA	NA	-18.7	8.8	42.9	14.5	2.9	2.2
Grotta del Fontino	GF 14	NA	NA	-20.1	10.3	48.0	13.5	3.6	0.5
Grotta del Fontino	GF 17	NA	NA	-18.6	9.7	42.4	15.1	3.2	0.6
Grotta del Fontino	GF 19	NA	NA	-19.9	10.1	45.1	13.8	3.3	1.3
Grotta del Fontino	GF 20	NA	NA	-17.9	9.0	42.5	14.9	3.3	1.6
Grotta del Fontino	GF 21	NA	NA	-20.7	10.4	36.8	11.5	3.2	1.0
Grotta del Fontino	GF 22	NA	NA	-19.4	9.7	45.9	13.2	3.4	0.5
Grotta del Fontino	GF 23	NA	NA	-20.8	10.1	34.7	9.6	3.6	0.2
Grotta del Fontino	GF 25	NA	NA	-19.6	8.5	43.4	13.5	3.2	1.1
Grotta del Fontino	GF 26	NA	NA	-21.1	10.0	40.2	10.4	3.4	2.2
Grotta del Fontino	GF 27	NA	$\rm NA$	-21.2	9.0	40.0	10.3	3.5	0.3
Grotta del Fontino	GF 31	NA	$\rm NA$	-18.8	8.2	47.7	12.2	3.6	$0.7\,$
Grotta del Fontino	GF 32	NA	$\rm NA$	-19.6	9.3	38.5	12.5	3.1	$0.2\,$
Grotta del Fontino	GF 35	NA	$\rm NA$	-19.6	10.1	41.8	13.7	3.1	0.4
Grotta del Fontino	GF 37	NA	NA	-19.1	8.5	41.6	13.9	3.0	$0.2\,$
Grotta del Fontino	GF 39	NA	NA	-19.4	9.8	46.3	13.8	3.3	2.1
Fontenoce di Recanati	Re 74	NA	Child $12 - 14y$	-19.3	9.3	42.9	15.9	3.0	6.2
Fontenoce di Recanati	Re 76	NA	Infant $0-6$ m	-19.6	9.4	31.0	10.5	2.9	1.6
Fontenoce di Recanati	Re 77	NA	Child $10 - 11y$	-19.6	8.2	40.7	14.4	3.0	1.6
Fontenoce di Recanati	Re 78	M	Young adult	-19.8	9.3	44.1	16.2	3.1	1.2
Fontenoce di Recanati	Re 80	M	Young adult	-19.9	8.9	40.4	13.1	3.6	1.8
Fontenoce di Recanati	Re 86	М	Young adult	-19.2	9.6	42.3	13.6	3.6	2.9
Fontenoce di Recanati	Re 87	NA	Child	-20.3	9.2	41.0	13.1	3.5	2.5

Table 4 (continued)

to an overestimation of this consumption because these values could be ascribed also to a regular intake of plant proteins (Fraser et al. [2013\)](#page-14-0). We can define within this framework a model wherein the inferred plant δ^{15} N value has been determined by taking this offset from the herbivore collagen $\delta^{15}N$ values (4.3‰), assuming that humans and herbivores consumed the same plant resources. The standard model can thereby be drawn suggesting how the mean δ^{15} N values for every site indicate a diet exclusively (100%) based on animal protein consumption, and Le Lellere seems to be partially at odds with an animal protein consumption of $> 80\%$ of the total protein exploitation (Fig. 4).

Other scenarios could be designed using the real plant values instead of the inferred ones.

The aforementioned literature data for wheat, barley, and peas have been considered a starting point for the models used to quantify the animal protein fraction.

If only wheat (Fig. [5\)](#page-10-0) was the origin of the plant protein intake, every site would be characterized by a vegetarian eating program (fraction of animal protein in total dietary proteins = 0%).

Conversely, if barley (Fig. [6\)](#page-11-0) is set as the starting point of the model, the communities under consideration would have had a mixed diet composed of around 80–90% of animal proteins for almost all the Tuscan sites and Fontenoce di

Recanati, while Le Lellere would sustain a more equilibrated landscape (50% of animal proteins consumption).

A comparable scenario should be outlined with the pulses (Le Lellere lies below 60%, whereas the other communities exceed 90%; Fig. [7\)](#page-11-0).

The use of Central Italian diachronic sample from Brescia et al. ([2002\)](#page-13-0) allows us to reach a credible scenario where Podere Cucule, Buca di Spaccasasso, Grotta del Fontino, and Fontenoce di Recanati communities seem to be featured by a remarkable but not exclusive percentage of animal proteins in their nutritional habits. Indeed, they are characterized by 60–80% of the fraction of animal proteins out of total dietary ones whereas Le Lellere does not exceed 30% (Fig. [8\)](#page-12-0).

Discussion

The stable isotope analysis performed for Italian eneolithic human remains in five sites suggests the people consumed a diet mainly based on C_3 plant and C_3 consumer backbone resources.

The value distribution appears consistent with an overall subsistence based on greater local resource procurement, supported by regional productivity maximization, than the hunting commitments typical of the earlier populations (Lelli et al. [2012](#page-14-0); Mannino et al. [2015;](#page-15-0) Pickard [2016;](#page-15-0) Crittenden and

Table 5 Species, stable isotope measurements, and collagen quality control indicators of the faunal remains from Fontenoce di Recanati (Marche region). NA, not available

Schnorr [2017](#page-14-0); Scorrano et al. [2018](#page-15-0)). Even though a global land supply could be highlighted in all the considered areas, appreciable differences seem to exist at the local/regional level that could be ascertained by a single evaluation.

At first glance, there is no evidence of C_4 burning up in the whole sample, supporting the notion of widespread Bronze Age consumption in Italy moving from the northeastern regions (Varalli et al. [2016\)](#page-15-0). In spite of the lack of botanical records for these areas, we are able to identify wheat, barley, and some legumes as the plants normally consumed by these communities, on the basis of the archeological information.

The data allows us to state that there is no direct evidence of exclusive marine resource intake, although their occasional consumption (along with freshwater consumption) cannot be ruled out at all. In fact, up to 20% of the consumed protein could conceivably come from the marine ecosystem without any δ^{13} C shift in collagen-derived values (Milner et al. [2004](#page-15-0)); thus, a mixed diet could be easily misidentified as an exclusively terrestrial diet (Jim et al. [2006](#page-14-0)). The seashore vicinity of some sites, such as those at Grotta del Fontino and Buca di Spaccasasso, might suggest the putative role marine resources could play in the diet; this is further supported by the discovery of shellfish remains attributable to selective Tuscan coastal environments (Vigliardi [2002;](#page-15-0) Cavanna [2007\)](#page-13-0) even though the presence of shellfish remains alone cannot be taken to prove their edible usage due to their ornamental role (Micheli [2004](#page-15-0)). This suggests the existence of a process of collecting and supplying such resources from different coastal types in these areas. Buca di Spaccasasso is currently 10 km away from the current coastline, whereas Grotta del Fontino is located at the edge of the current Grosseto plain, but diachronic differences in the geographic conformation of the Maremma coast should be noted. The plain did not exist, the seaside slowly went into interruption, and the Ombrone and Bruna rivers flowed into the Prile coastal lake (Fig. [9;](#page-12-0) Ceccarelli and Niccolucci [2003](#page-13-0)).

Table 6 Coupled Mann-Whitney test for δ^{13} C and δ^{15} N values in the analyzed samples

p value	Podere Cucule	Le Lellere	Buca di Spaccasasso	Grotta del Fontino	Fontenoce di Recanati
U/Z -score					
$\delta^{13}C$					
Podere Cucule		0.4	0.8	0.5	0.4
Le Lellere	$36/- 0.85$		0.6	0.5	0.4
Buca di Spaccasasso	$22/- 0.19$	$54.5/-$ 0.55		0.8	0.9
Grotta del Fontino	$116.5/- 0.67$	334/0.65	$177/- 0.25$		1
Fontenoce di Recanati	$68/- 0.91$	203/0.84	$114.5/- 1.18$	$703.5/- 0.01$	
$\delta^{15}N$					
Podere Cucule		0.6	0.7	0.7	1
Le Lellere	$41.5/- 0.44$		0.1	0.2	0.3
Buca di Spaccasasso	21/0.32	$37.5/- 1.59$		0.6	0.2
Grotta del Fontino	131/0.34	302/1.26	169/0.53		0.7
Fontenoce di Recanati	89/0.02	195.5/1.01	$86/- 1.2$	676.5/0.44	

Table 7 Mean and standard deviation (SD) for the isotopic values

			Average $\delta^{13}C\%$ SD Average $\delta^{15}N\%$ SD	
Podere Cucule	-18.2	$0.3 \quad 9.0$		1.1
Le Lellere	-19.9	0.4 7.6		0.5
Buca di Spaccasasso	-19.1	$1.3 \quad 9.6$		0.2
Grotta del Fontino	-19.5	0.9 9.6		0.7
Fontenoce di Recanati -19.5		$0.5 \quad 9.1$		0.8

The standard linear mixing model (Fig. [4\)](#page-10-0) points out very high values that are unlikely to be considered because they entail vast quantities of edible meat and dairy products, and even though several faunal remains have been recovered, they cannot account for the sole animal product expenditure. In spite of the scanty information available, a general overview of plant gathering and cultivation could be roughly ascertained from the published list of findings and indirect evidence (Bellini et al. [2008\)](#page-13-0) suggesting how cereals and legumes were widely recovered in central Italian archeological surveys. Indeed, Triticum sp. (both glumen and naked wheats) and barley (Hordeum sp.) represented the most frequent occurrences, even though some Panicum miliaceum seeds are described in Tuscany in the Eneolithic/Bronze Age time edge (Bellini et al. [2008](#page-13-0); Mariotti-Lippi et al. [2017](#page-15-0)). This latter evidence seems to be consistent with other research (Tafuri et al. [2009](#page-15-0); Varalli et al. [2016\)](#page-15-0) into how millet could be an important crop during the Middle Bronze Age in Italy. The research shows that it could be cultivated from at least the Early Bronze Age onward, and perhaps before then, though no isotopic evidence suggests its edible usage in presented eneolithic communities.

As stated above, if only wheat (Fig. [5\)](#page-10-0) was the origin of the plant protein intake, every site would be characterized by a vegetarian eating program. This is unlikely, given the ecological and structural biological needs, as grains contain less protein than animal flesh, legumes, and seeds, at an average of about 10–12% of their dry weight (Shewry and Halford [2002](#page-15-0)). Moreover, wheat and other cereal proteins do not contain as many nutritionally essential amino acids as animal protein. Further, if consumed as a sole protein source, they are not utilized with the same efficiency as animal protein compounds to meet the physiological requirements for the people living there as a total protein, or to provide specific indispensable amino acids such as lysine, threonine, and tryptophan. However, when combined with other food such as legumes, oil seeds, and animal products, the proteins of wheat exhibit excellent nutritional complementarity (Young and Pellett [1985](#page-15-0)). The δ^{15} N values of wheat reported by Vaiglova et al. ([2014](#page-15-0)) seem too high for the Italian herbivore samples that sit at lower values (4.3‰) than their putative forages (5.6 ‰) and it should be noted that these high values could be ascribed to manuring practices that were employed as early as the Neolithic period in Europe to intensify arable production leading to increases in populations and to more opportunities for trading (Guttmann et al. [2005\)](#page-14-0). The manuring effect is well documented as a modifying factor for nitrogen values (Fraser et al. [2011;](#page-14-0) Kanstrup et al. [2011](#page-14-0), [2014](#page-14-0)), which could affect proper nitrogen identification in the recorded wheat. The result obtained through the modern crops appears to be more conceivable (Fig. [8](#page-12-0)): despite the known differences in δ^{15} N values due to differential degree of manuring and irrigation between modern and ancient horticultural practices, the proposed model fits the economic sustainability of ancient communities, whose subsistence should be based on a mixed animal and plant resource consumption.

The barley-based model (Fig. [6](#page-11-0)) seems to be even comparable: this cereal is usually cultivated in warm climates and complements wheat, which thrives in cooler climates. Their nutritional features are quite similar: it is considered high in

Fig. 3 Plot of the average values in the evaluated samples

Fig. 4 Modeled scenario estimating the animal protein fraction (percentage) of total dietary protein in the Central Italy human diet starting from the herbivore values recorded in Fontenoce di Recanati. The herbivore average δ^{15} N value allows to infer the plant $\delta^{15}N$ value by subtract 4‰. Avg, average value

carbohydrates, fat, dietary fiber, vitamin B, vitamin C, calcium, iron, magnesium, phosphorus, potassium, zinc, and folate but is deficient in the essential mineral selenium (Biel and Jacyno [2013](#page-13-0)).

A similar scenario would be outlined if we considered the pulses (Fig. [7\)](#page-11-0) the main plant resources. Pulses typically contain about twice the amount of protein found in whole grain cereals such as wheat and barley and can constitute a major source of protein in a diet. Moreover, when other foods are combined with pulses, their nutritional value is further enhanced, because other foods enable the body to better absorb the nutrients found in pulses (Trichopoulou et al. [2009](#page-15-0)). Thus, they could be a primary foodstuff for ancient people, an idea that is supported by evidence related to agronomy information regarding the legume cultivation attested in the Tuscany area (Oliva [1939\)](#page-15-0), and in general these plants gained greater edible magnitude during the Eneolithic (MariottiLippi et al. [2006\)](#page-15-0). All the proposed scenarios consider communities with mixed diets of plants, cereals, pulses, and animal proteins, and undoubtedly people from Le Lellere preferred a more mixed diet than the other peoples, mainly based on their arable resources. No differences could be outlined between the two sides of the Apennines, which makes sense, given the shared open trade net between Tuscany and Marche (Sarti [2005](#page-15-0)). Cultural contacts between the two sides of the Apennines have been extensively documented, as well as cultural swaps in material culture and funerary architectures (Cazzella and Silvestrini [2005;](#page-13-0) Leonini and Sarti [2006](#page-14-0)), especially between the eastern Marche region and the southern area of Tuscany. The northeastern area, meanwhile, seems to preserve a more autarchic trait: in this frame, the difference between Le Lellere and the other evaluated Tuscanian samples could mirror specific features in dietary habits. Of course, this scenario was

Fig. 5 Modeled scenario estimating the animal protein fraction (percentage) of total dietary protein in the Central Italy human diet starting from wheat values from Vaiglova et al. ([2014](#page-15-0)). Herbivore value is obtained by the increase of 4‰ of the starting point of the model

Fig. 6 Modeled scenario estimating the animal protein fraction (percentage) of total dietary protein in the Central Italy human diet starting from barley values from Vaiglova et al. ([2014](#page-15-0)). Herbivore value is obtained by the increase of 4‰ of the starting point of the model

sporadically intermixed, as suggested by the condition of Podere Cucule, which falls in the quite homogenous scenario proposed for southern Tuscany and Marche as to the exploited animal protein fraction highlighted by the $\delta^{15}N$ values. Although the sample sizes for some funerary contexts—such as Le Lellere, Podere Cucule, and Buca di Spaccasasso—are quite low and render any statistical evaluation unacceptable, the Kolmogorov-Smirnov test fails to identify significant differences between the Tuscan sites and the Fontenoce di Recanati isotopes values (δ^{13} C, D = 0.10; p = 0.98; δ^{15} N, D = 0.14; p = 0.75), confirming the overall dietary similarities between the two sides of the Apennines.

This homogeneity notwithstanding several peculiar elements could indicate some differences between these subsistence economies and could be ascribed to local environmental exploitation. Indeed, if the extremely small sample size could lead to misinterpretation, Podere Cucule shows very different δ^{13} C values. These values could be ascribed to several factors despite the lack of carpological elements, making C_4 plant consumption less plausible despite their presence in Tuscany in other surveys (Bellini et al. [2008\)](#page-13-0). Trade from the Adriatic regions suggests at least a sporadic and not continuous exploitation that could move the carbon values toward less negative results. In spite of this evidence, the changes in plant management strategies could address this heterogeneity. Indeed, many studies have shown that plants grown under water stress produce higher δ^{13} C values (Saugier et al. [1993;](#page-15-0) Ferrio et al. [2005\)](#page-14-0), allowing for the determination of a significant expected relationship between δ^{13} C and the environmental parameters related to water availability. This proposed approach has been pioneeringly used (Araus and Bux [1993\)](#page-13-0) in grains from archeological sites to gain insight into the environmental conditions of early agriculture, allowing for the relative quantification of water inputs and crop yields (Araus et al. [2002\)](#page-13-0). It could also be a proxy for obtaining further information about climate-derived and anthropogenic effects on plants' water status, as well as for distinguishing different strategies of water

Fig. 7 Modeled scenario estimating the animal protein fraction (percentage) of total dietary protein in the Central Italy human diet starting from pulses value from Vaiglova et al. [\(2014\)](#page-15-0). Herbivore value is obtained by the increase of 4‰ of the starting point of the model

Fig. 8 Modeled scenario estimating the animal protein fraction (percentage) of total dietary protein in the Central Italy human diet starting from actual grains values from Brescia et al. [2002.](#page-13-0) Herbivore value is obtained by the increase of 4‰ of the starting point of the model

management. Thus, the more positive δ^{13} C values could mask different strategies to obtain plant resources in Podere Cucule, even though the limited sample size makes it hard to properly understand the subsistence economy in such a community. Of course, this bias underlines a need to magnify the human sample size related to this community even though the current one represents all the human specimens recovered to date. Moreover, local seeds or plant macroremains and faunal individuals should be classified in order to confirm this subsistence role for the community and clarify inferences about the putative inter-site differences.

A local environmental constraint also seems to be responsible for the isotopic results at the Marche site of Fontenoce di Recanati. This area (Silvestrini and Pignocchi [1997\)](#page-15-0) is located close to the Potenza and Menocchia rivers, which could have represented valuable hunting and fishing grounds for the people living in their vicinity. Despite substantial introgression evidence from the eastern shores of the Adriatic sea (Cazzella and Silvestrini [2005\)](#page-13-0), there is no nutritional awareness of foods that could be attributed to Aegean/Balkan populations, such as C_4 metabolism plants, that were later introduced to Italy from the northeast (Tafuri et al. [2009](#page-15-0), [2018](#page-15-0)).

Fig. 9 Topographical location of ancient Prile coastal lake

Conclusions

This paper presents the first data about subsistence strategies for Copper Age communities in Central Italy, filling an existing gap in the bio-molecular analysis of the prehistoric Italian inhabitants. The dissection of the diet of people buried in the Tuscan and Marche areas during the Eneolithic seems to indicate a dietary pattern mainly based on terrestrial resources, with the greatest role played by animal protein. If the backbone dietary component for all the communities came from C_3 plants, it cannot be denied that there was a remarkable recourse to protein foodstuffs in every community, with consumption of animal sources between 60 and 100%, except in Le Lellere, where the animal protein input dropped to low extent (30–50%). C_3 plant sources such as wheat, barley, and pulses do not seem exclusively important to the diets of those samples, while the animal protein component would mainly come from the flesh of terrestrial herbivores (the livestock of the community), even though the use of milk and dairy products cannot be ruled out.

The archeological findings and the topographical locations of Buca di Spaccasasso, Grotta del Fontino, and Fontenoce di Recanati could represent evidence of the consumption of marine and freshwater resources, and their burning should not be ignored as merely occasional. These marine and freshwater protein resources could also be highlighted in Neolithic coastal sites in Italy (Lelli et al. [2012\)](#page-14-0) and in the Mediterranean region (Lightfoot et al. [2011](#page-14-0)), suggesting that consumption of this food clearly supported the rise of refined farming societies. These results support the hypothesis that the land exploitation maximization that occurred during the Eneolithic period had to contend with environmental constraints that, in such cases, provided beneficial habitats for spontaneous foraging. This might be corroborated by the evidence of putative eating support in the form of spontaneous C_3 plants in Podere Cucule, where the watering strategies could not be so finely tuned as to guarantee steady opportunities for cereal reaping. Despite a lack of carpological evidence, occasional C_4 plant consumption could be conceivable and a negligible amount of marine resource exploitation cannot be totally ruled out.

The existing fully developed trade routes between the two sides of the Apennines could address the highlighted overall dietary homogeneity of the studied communities, especially between Fontenoce di Recanati and the southern Tuscan human groups such as those at Grotta del Fontino and Buca di Spaccasasso, with lesser influence for Le Lellere and Podere Cucule, which seem to suggest a more locally based subsistence strategy, even though the funerary affinities do not match this overall diet homogeneity.

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