



Diversification in Feeding Pattern of Livestock in Early Bronze Age Northwestern China

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Animal husbandry has been an indispensable part of human subsistence since the origin of agriculture. Along with the prehistoric cultural exchange, several kinds of major domestic animals diffused and gradually got popularized across the Eurasia. The specific geographic setting makes the Hexi Corridor in northwestern China one of the key regions to converge various types of major crops and livestock, and to witness the deep impact of novel species on local subsistence. Archeological evidence reveals an emergence of early oriental-occidental cultural communication at the opening of the local Bronze Age in Hexi Corridor, resulting in a significant shift of local subsistence. However, due to the lack of more detailed archeological evidence, the initial timing and trajectory of the transformation of livestock feeding patterns remain unclear. In this study, we reported systematic stable isotope and precise radiocarbon dating analyses on faunal remains unearthed from Huoshiliang and Ganggangwa, two Early Bronze Age settlements (ca. 4,000–3,700 BP) in middle Hexi Corridor. Our results show distinct diversification in livestock feeding patterns at ~3,850 cal BP; in contrast with previous periods, some omnivorous livestock appear to have consumed mainly C₃ foodstuff and some herbivorous livestock primarily consumed C₄ plants. Combined with published stable isotope data and other archeological findings in the neighboring region, a clearer trajectory of the evolution of livestock feeding patterns has been revealed with diversified strategy amid the transformation during the Early Bronze Age in Hexi Corridor. We argued that the alteration of the local livestock feeding pattern reflects the attempt to achieve more efficient economy and sustainable society, in order to withstand the harsh arid environment in Hexi Corridor.

Keywords: Early Bronze Age, Hexi Corridor, stable isotopes, paleodiet, human subsistence, animal husbandry, diversified management

INTRODUCTION

The origin and spread of major crops and livestock markedly changed the pattern of human subsistence during prehistoric times, which generated deep impacts on the development of human society and the interaction between the environment and humans (Diamond and Bellwood, 2003; Liu et al., 2014; Spengler et al., 2017; Revelles et al., 2018; Dong et al., 2021a). During the Early Holocene (~11,700–8,200 BP, BP for year before present), various types of crops and livestock were domesticated in West Asia (especially wheat, barley, cattle, and sheep/goat; Bradley et al., 1998;

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Zeder and Hesse, 2000; Zeder, 2008; Riehl et al., 2013) and East Asia (especially foxtail millet, broomcorn millet, rice, pig, and dog; Fuller et al., 2009; Jones and Liu, 2009; Lu et al., 2009; Zhao, 2011; Yuan, 2015; Ren and Dong, 2016), respectively. These botanic and zoic agricultural elements then diffused along with the early phase of trans-continental cultural exchange in the following several millennia (Sherratt, 2006; Anthony, 2010; Dong et al., 2017; Frachetti et al., 2017; Taylor et al., 2021). The spread and popularization of domestic livestock (Ren and Dong, 2016; Hermes et al., 2020; Honeychurch et al., 2021; Wilkin et al., 2021), together with the spread of major crops (also known as food globalization; Jones et al., 2011, 2016; Liu et al., 2019a), deeply transformed the human subsistence worldwide.

The introduction of novel crops and livestock, combined with heterogeneous natural environment across Eurasia, sculpted diverse patterns of human subsistence in different regions during the Neolithic and Bronze Ages. The analysis of stable carbon and nitrogen isotopes has been proved as an effective way to reconstruct paleodiet (van der Merwe and Vogel, 1978) and provide a unique perspective on human subsistence and related behaviors, and has therefore been extensively used in previous research of prehistoric Eurasia (e.g., Hu et al., 2008; Barton et al., 2009; Motuzaite Matuzeviciute et al., 2015; Chen et al., 2016; Ma et al., 2016; Miller et al., 2019; Li et al., 2020; Wilkin et al., 2020). In particular, human behavior related to animal husbandry practices has been frequently discussed via dietary perspective in recent years (e.g., Dai et al., 2016; Hermes et al., 2019; Ma et al., 2021; Vaiglova et al., 2021). These studies of paleodiet have offered valuable insights into human subsistence strategies in the past.

Acting as the passageway between the Yellow River Basin and the Arid Central Asia, the Hexi Corridor in northwestern China is regarded as a key region in cross-Eurasian cultural exchange (Dong et al., 2017; Yang, 2017), and has therefore received attention from an array of multidisciplinary researchers. The evolution and chronology of the prehistoric cultures in Hexi Corridor have been well established (Li, 2011; Yang, 2017; Yang et al., 2019a; Li et al., 2021). In addition, a significant body of Hexi Corridor research has focused on different aspects of the past relationship between humans and the environment (Sun et al., 2010; Li et al., 2011, 2017; Zhou et al., 2012; Yang et al., 2017, 2020; Zhang et al., 2017; Shen et al., 2018; Shi et al., 2019; Dong et al., 2021b). Furthermore, reconstruction of human subsistence has been finely reported *via* archaeobotanic (Zhou et al., 2016; Liu et al., 2019b,c, 2021; Shi et al., 2022), zooarchaeological (Flad et al., 2007; Song et al., 2016; Yang et al., 2019b), geochemical (Dodson et al., 2009), and genetic perspectives (Xiong et al., 2022).

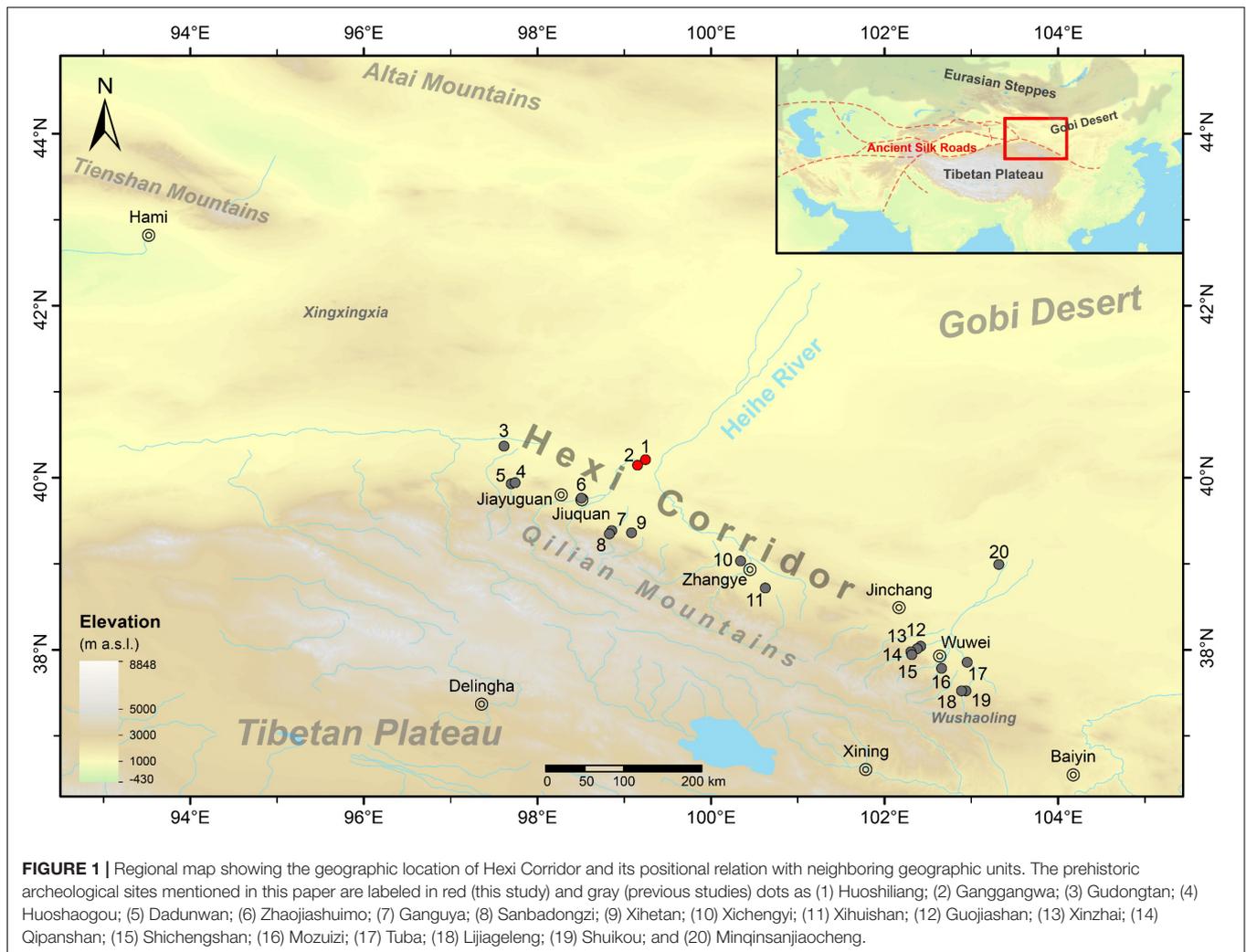
For dietary perspective, abundant stable isotope data from human and faunal remains in prehistoric Hexi Corridor have been published (Atahan et al., 2011; Liu et al., 2014, 2016; Zhang et al., 2015; Yang et al., 2019b; Ma et al., 2021; Vaiglova et al., 2021). Stable isotope evidence reveals a significant shift in human diet from C_4 millet crops to C_3 wheat and barley in Hexi Corridor during 4,000–3,000 BP (Yang et al., 2019b). The initial timing of this shift could be traced back to 4,000–3,800 BP (Liu et al., 2014), right after the introduction of these Western crops at the beginning of the local Early Bronze

Age (EBA; Dodson et al., 2013). Meanwhile, a similar pattern of dietary shift in domestic omnivores (pigs and dogs) has also been detected (Yang et al., 2019b). As the local domestic omnivore usually possess a C_4 dietary habit and the novel herbivorous livestock possess a C_3 dietary habit, the appearance of domestic individual showing opposite diet is abnormal and worth an extra attention. Recent studies with abundant stable isotope data from Hexi Corridor faunal remains have shed light on the practice of pastoralism and localized management strategy during EBA (Atahan et al., 2011; Yang et al., 2019b; Ma et al., 2021; Vaiglova et al., 2021). However, in this study, we sought to resolve several outstanding issues regarding livestock feeding patterns, namely, (1) the implications of individual livestock with abnormal $\delta^{13}C$ -values and (2) the initial timing and trajectory for the shift in livestock feeding patterns during EBA in the Hexi Corridor. Based on materials acquired from archeological survey, a previous study on stable carbon and nitrogen isotopes from Huoshiliang and Ganggangwa sites has effectively contributed to the investigation of paleodiet and human subsistence during EBA in Hexi Corridor (Atahan et al., 2011). However, systematic excavation is still of necessity to provide a dynamic perspective to reveal the issues mentioned above.

This study reports systematic stable isotope analysis and precise radiocarbon dating results from faunal remains unearthed during the excavation of these two EBA archeological settlements. Carbon and nitrogen isotope values were acquired for a total of 56 specimens from a range of both livestock and wild animals from these two sites. In addition to 6 dates previously reported by Ren et al. (2022), a total of 15 direct radiocarbon dates were obtained to provide precise chronology in this study. We aimed to give clearer insight into the initial timing of the previously observed shift in livestock feeding patterns, as well as the implications of abnormal $\delta^{13}C$ -values. By combining these newly acquired data with previously published studies, we were able to use faunal carbon isotope data to more precisely reconstruct the trajectory of the transformation of livestock feeding patterns and give novel insight into the transformation of local subsistence and animal management in the Hexi Corridor.

STUDY REGION

Being located in northwestern China, the Hexi Corridor is a passageway bordered by the Tibetan Plateau on the southwest and the Gobi Desert on the northeast, which connects the Yellow River Basin at Wushaoling Mountain on the east and the Arid Central Asia at Xingxingxia Gorge on the west (**Figure 1**). With modern mean annual precipitation lower than 200 mm, the dominant climate type in modern Hexi Corridor involves temperate semi-arid (BSk) and temperate arid (BWk) according to the Köppen classification (Chan et al., 2016). In addition, C_3 plants make up the majority of modern flora of Hexi Corridor while a few taxa of C_4 plants mainly distributed in the arid zone act as supplementary (Jiang et al., 2019). Rivers originating from the Qilian Mountains of northeastern Tibetan Plateau flow into Hexi Corridor, developing bunches of oases, making the region a relatively decent habitat and transitable passageway among the



arid deserts in the communication between the East and the West, especially for the ancient Silk Road. However, restricted by the low precipitation and poor land productivity, this region has always suffered from the harsh natural environment when developing economy. The Heihe River forms its channel across the middle Hexi Corridor and streams toward the hinterland of Gobi Desert, raising a bunch of oases along its valley, which provide a possible avenue linking with the Eurasian Steppes (Jaang, 2015).

Huoshiliang (99.2446°E, 40.2116°N, 1,198 m a.s.l.) and Ganggangwa (99.1524°E, 40.1469°N, 1,203 m a.s.l.) are located in Jinta County, Jiuquan Region, northwestern Gansu Province (Figure 1). Situated at Jinta Basin in the middle reaches of Heihe River, these two archeological sites are identified as settlements and smelting locations of the EBA in middle Hexi Corridor (Figure 2). Standing on the intersection among the Yellow River Basin, Arid Central Asia, and Eurasian Steppes, Huoshiliang and Ganggangwa sites witnessed the emergence of prehistoric East-West cultural exchange and the opening of EBA in Hexi Corridor. In particular, the river valley of Heihe provides critical connection between Hexi Corridor and the Eurasian

Steppes, making these sites pivotal stations during the prehistoric cultural exchange in northwestern China (Jaang, 2015). The geographic settings, landscapes, as well as previous archeological-environmental surveys and related studies of these two sites have been finely reported (Dodson et al., 2009; Sun et al., 2010; Atahan et al., 2011; Li et al., 2011; Zhou et al., 2016; Yang et al., 2017). As indicated by radiocarbon dating results, the age of these two sites spans from ca. 4,100 to 3,700 cal BP (Dodson et al., 2009, 2013; Atahan et al., 2011). The starting age of Ganggangwa site was reported to be slightly earlier than Huoshiliang site, reaching the transition from Late Machang Culture to Xichengyi Culture (Gansu Provincial Institute of Cultural Relics and Archaeology, 2020).

In consideration of achieving more comprehensive understanding on Huoshiliang and Ganggangwa sites, Gansu Provincial Institute of Archaeology and Cultural Relics launched an exploratory excavation in 2017. According to the preliminary archeological prospection, an area of 32 m² in the core zone of each site was chosen for excavation by archeologists. During the excavation, plenty of pottery sherds, bronze slags, lithic tools, as well as animal bones, were unearthed from different

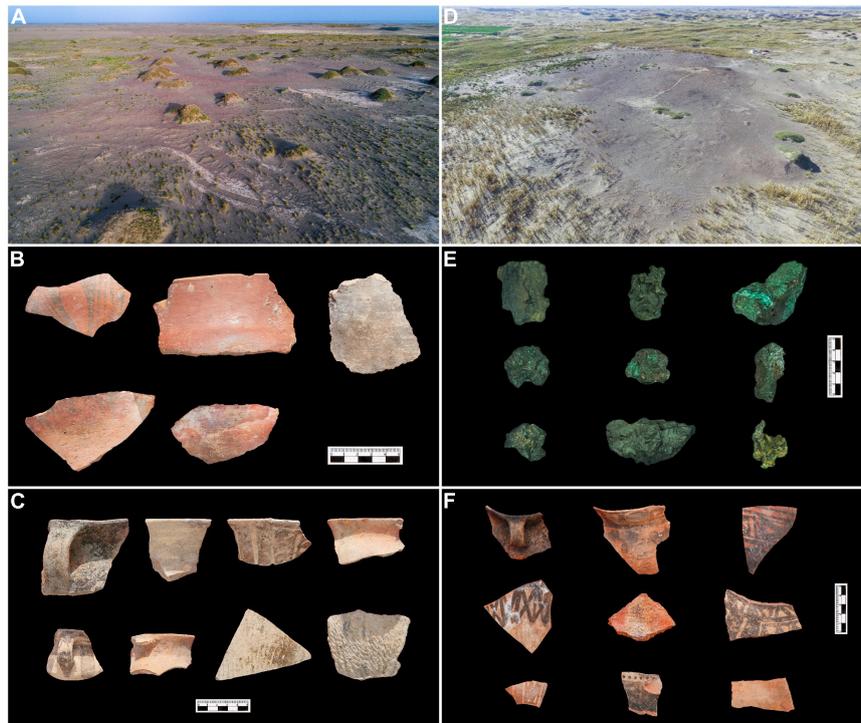


FIGURE 2 | Images of the archeological sites and their cultural relics. **(A)** Aerial photo of Ganggangwa site. **(B)** Pottery sherds of Late Machang Culture unearthed from Ganggangwa site (T1911F2). **(C)** Pottery sherds of Xichengyi Culture unearthed from Ganggangwa site (T1911L3). **(D)** Aerial photo of Huoshiliang site. **(E)** Bronze slags unearthed from Huoshiliang site (T1007L3). **(F)** Pottery sherds of Xichengyi Culture unearthed from Huoshiliang site (T1007H33).

archeological units, including house sites, ash pits, and burials. As indicated by the characters of typical pottery remains, the entirety of Huoshiliang and the majority of Ganggangwa belong to Xichengyi Culture of the local EBA (*ca.* 4,000–3,600 BP), while the early stage of Ganggangwa belongs to the Late Machang Culture of Late Neolithic Age in Hexi Corridor (*ca.* 4,100–4,000 BP) (Figure 2).

MATERIALS AND METHODS

Sample Selection

All unearthed animal bones were collected and cleaned for species identification. Dr. Lele Ren from Lanzhou University performed the species identification; the results were reported in another manuscript in detailed, which was submitted to the *Journal of Archaeological Science* and has currently been accepted (Ren et al., 2022). On this basis, we sampled bone specimens of different kinds of species from various archeological units. Bones with clearer morphological features were preferentially sampled for stable isotope analysis, which can contribute to lower the risk of misidentification of species. Even so, it is still impossible to 100% avoid such issue. Since Huoshiliang and Ganggangwa are neighboring, homogeneous and almost coetaneous to each other, data from these two sites were reported combinedly. A total of 62 pieces of animal bones from different individuals were sampled from Huoshiliang ($n = 34$) and Ganggangwa ($n = 28$)

sites, including livestock (pig, $n = 11$; dog, $n = 9$; cattle, $n = 6$; and sheep/goat, $n = 16$) and wildlife (gazelle, $n = 5$; deer, $n = 7$; hare, $n = 4$; and bird, $n = 4$). Samples for radiocarbon dating were selected after the completion of stable isotope analysis such that livestock individuals with abnormal isotopic values could be targeted in addition to capturing the full stratigraphic range of the sites.

Collagen Extraction and Quality Examination

The method adopted to extract bone collagen in this study referred to Ma et al. (2016), which was modified from Richards and Hedges (1999). Bones were water washed and polished *via* a handheld electric grinder to remove the sediments adhered on the surface. Dense part on the bone was cut off as small fragment that weighed around 1 g for extraction. Then, the bone fragments were demineralized in 0.5 mol/L hydrochloric (HCl) solution at 5°C for more than 10 days, with refresh of the HCl solution every 1 or 2 days. When the samples were softened and stopped producing bubbles, they were washed with purified water and further reacted in 0.125 mol/L sodium hydroxide (NaOH) solution at 5°C for 20 h. After being neutralized again with purified water, the samples were placed in an acidic solution (HCl, pH = 3) at 75°C for 48 h to gelatinize, and were filtered afterward. For the final step, the samples were frozen in the refrigerator and freeze-dried in a lyophilizer, with concentrated solid collagen remained.

Rate of yield and atomic C/N ratio was utilized to examine the quality of bone collagen. With five samples experienced failure during the collagen extraction, the rest of the samples ($n = 57$) were weighed to calculate the rate of yield (Table 1). Content of carbon and nitrogen of all collagen samples was measured at the State Key Laboratory of Applied Organic Chemistry in Lanzhou University via an Elemental Analyzer (Vario EL Cube) to calculate atomic C/N ratios as described below.

Stable Isotope Analysis

For stable carbon isotope, $\delta^{13}\text{C}$ -value varied significantly between C_3 and C_4 plants due to the different CO_2 absorption during the photosynthesis. Among the major crops, wheat, barley, and rice are C_3 plants while foxtail millet, broomcorn millet, corn, and sorghum are C_4 plants. Such signal further transmits to animals and humans through the food chains, thus can be used as detector for diet and human subsistence (van der Merwe and Vogel, 1978), as well as the human behavior about animal husbandry practices (e.g., Dai et al., 2016; Hermes et al., 2019; Ma et al., 2021). Referring to previous study in Hexi Corridor (Yang et al., 2019b), $\delta^{13}\text{C}$ -values of -18 and -11‰ are adopted as boundary lines for C_3 ($< -18\text{‰}$), mixed C_3/C_4 (-18 to -11‰), and C_4 diets ($> -11\text{‰}$) in this study. For stable nitrogen isotope, the proportion of $^{15}\text{N}/^{14}\text{N}$ enriches along the food chains, presenting distinct $\delta^{15}\text{N}$ -values in different trophic levels (Lee-Thorp, 2008). However, the background value of $\delta^{15}\text{N}$ could be raised by draught stress in arid zones, which results in notable interregional discrepancy (Ambrose and DeNiro, 1986; Schwarcz et al., 1999; Hartman, 2011).

Stable carbon and nitrogen isotope analyses were carried out at the Key Laboratory of Western China's Environmental Systems (Ministry of Education) in Lanzhou University. Collagen samples were proceeded in an automated carbon and nitrogen analyzer linked to a Thermo Finnigan Flash DELTA-plus XL mass spectrometer. The ratios of carbon and nitrogen isotopes were calculated following the V-PDB and AIR standards using a two-point linear normalization, respectively. Graphite (IAEA code: USGS 24, $\delta^{13}\text{C} = -16.0 \pm 0.1\text{‰}$) and caffeine (IAEA code: IAEA-600, $\delta^{15}\text{N} = 1.0 \pm 0.2\text{‰}$) standards were used to calculate the results, with the precision of final data not exceeding $\pm 0.2\text{‰}$. Full information is available in Table 1. Since the $\delta^{15}\text{N}$ measurement of five samples (specially labeled in Table 1) was missed during the analysis, remedial analysis was carried out by the Beta Analytic Testing Laboratory with instruments of Costech ECS and Thermo Electron DeltaPlus IRMS; and the results were reported with the precision of $\pm 0.5\text{‰}$. $\delta^{13}\text{C}$ of these five samples were also measured as contrast to confirm that data from these two labs are comparable.

Radiocarbon Dating

In total, 15 collagen samples (including 6 samples first reported in Ren et al., 2022) from various archeological units were selected for radiocarbon dating in different batches to construct the chronology of these two sites precisely. Samples were sent for accelerator mass spectrometry (AMS) radiocarbon dating at Key Laboratory of Western China's Environmental Systems (Ministry of Education) in Lanzhou University or Beta Analytic Testing

Laboratory, respectively. Conventional ^{14}C ages were calibrated with the IntCal20 calibration curve (Reimer et al., 2020) in the OxCal 4.4.4 online program (Bronk Ramsey, 2021).

RESULTS

Bone Preservation

Excluding one sample that experienced a low collagen yield ($< 0.5\%$), the remaining 56 samples yielded between 2.8 and 12.5% collagen. The yield of sample HSL-34 was too low (0.2%); thus, the data of this sample was rejected due to its poor reliability. The quality of bone after the long-term preservation in soil was tested by atomic C/N ratio. As shown in Table 1, these 56 samples (excluded five samples experienced failure during extraction and one sample with too low yield) are well preserved. The carbon content for all samples range between 41.5 and 44.1% while the nitrogen content range between 14.6 and 15.9%, generating atomic C/N ratios from 3.2 to 3.5, which fall into the acceptable range established by previous studies (DeNiro, 1985; Ambrose, 1990; van Klinken, 1999).

Stable Isotopes

The stable carbon and nitrogen isotope data for all the individual animal samples are shown in Table 1 and Figure 3A. Data were further classified and analyzed by species (Table 2 and Figure 3B). The fauna utilized in Huoshiliang and Ganggangwa sites ranges widely in both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ -values, from -22.7 to -7.2‰ with mean and standard deviation (SD, 1σ) of $-15.3 \pm 4.4\text{‰}$ for $\delta^{13}\text{C}$, and from 3.0 to 10.8‰ with mean and SD at $7.7 \pm 1.7\text{‰}$ for $\delta^{15}\text{N}$, respectively. The $\delta^{15}\text{N}$ -value of livestock ranges relatively closely, from 6.1 to 10.8‰ with mean and SD at $8.3 \pm 1.0\text{‰}$. However, the $\delta^{13}\text{C}$ -value of livestock varied significantly between two groups consuming primarily C_3 and dominantly C_4 foods, respectively. For omnivorous livestock, more than half of the pigs (7/10) and all dogs consumed predominantly C_4 diets according to their $\delta^{13}\text{C}$ -values, while a small group of pigs (3/10) consumed mainly a C_3 diet, which was analyzed separately. For herbivorous livestock, most of the cattle and sheep/goat consumed C_3 or mixed C_3/C_4 but primarily C_3 diets, with only one individual of each category consumed a C_4 diet. The $\delta^{13}\text{C}$ -value of wildlife mainly fitted with C_3 diet, indicating a consuming reliance on the natural ecosystem, except for gazelles. Most of the gazelles (4/5) have mixed C_3/C_4 - $\delta^{13}\text{C}$ -values, indicating considerable intake of C_4 plants. The $\delta^{15}\text{N}$ -values of deer ($4.2 \pm 0.6\text{‰}$) were notably lower than that of other species (Figure 3B).

Chronology

The radiocarbon dating results for the 15 directly dated animal collagen samples are listed in Table 3. Among these results, six dates with special labels were first reported in Ren et al. (2022) to represent the integral age of the faunal remains, while the other nine dates were reported for the first time in this study. Bone collagen was regarded as better material than charred wood for radiocarbon dating due to the potential influence of "old wood"

TABLE 1 | Stable carbon and nitrogen isotope and quality indicator measurements of animal bone collagen from Huoshiliang and Ganggangwa sites.

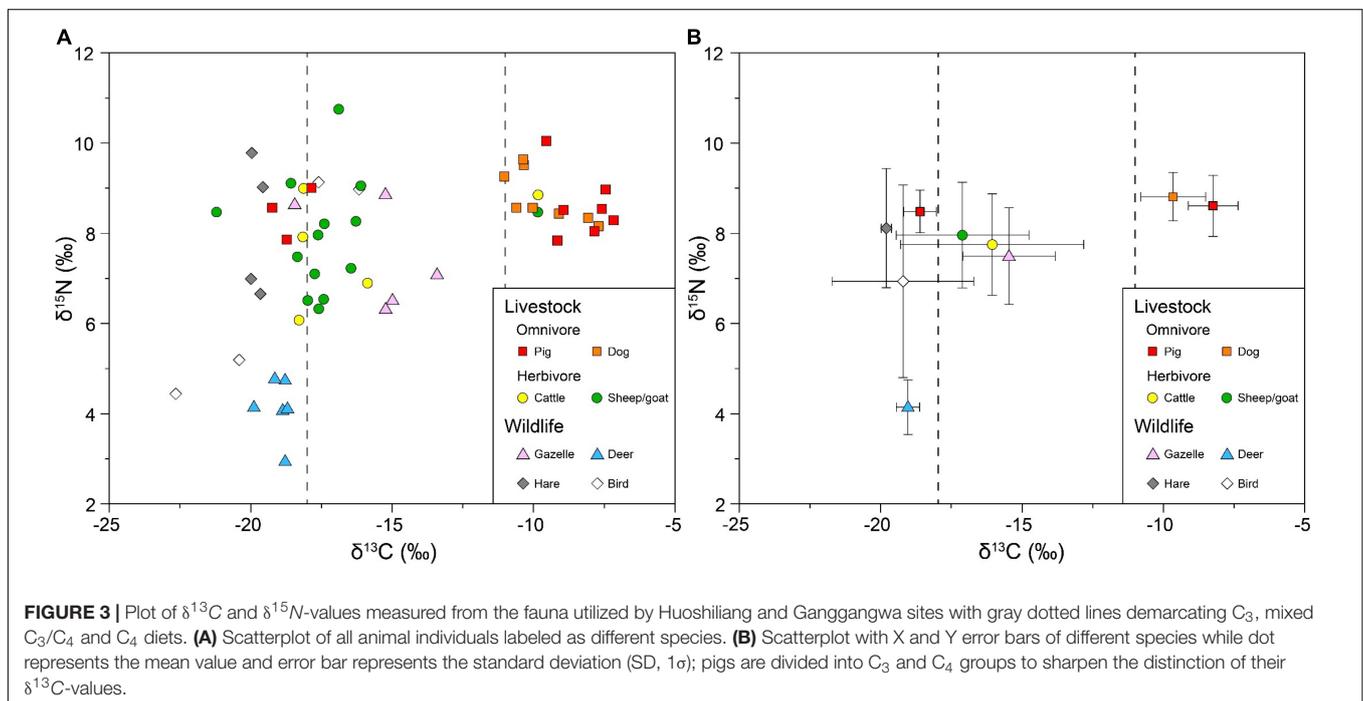
Lab ID	Site name	Context ^a	Taxon	Collagen (%)	C (%)	N (%)	C/N Ratio	$\delta^{13}\text{C}$ (‰)	$\delta^{15}\text{N}$ (‰)
HSL-15	Huoshiliang	2017JHIT1007M2	Pig (<i>Sus domestica</i>)	6.3	42.7	15.4	3.2	-17.8	9.0
HSL-25	Huoshiliang	2017JHIT1007L10	Pig (<i>S. domestica</i>)	6.4	43.4	15.8	3.2	-7.9	8.0
HSL-18	Huoshiliang	2017JHIT1007H26L2	Pig (<i>S. domestica</i>)	7.7	42.6	15.4	3.2	-9.2	7.8
HSL-19	Huoshiliang	2017JHIT1007H5L3	Pig (<i>S. domestica</i>)	5.0	42.7	15.6	3.2	-8.9	8.5
HSL-11	Huoshiliang	2017JHIT1007H9	Pig (<i>S. domestica</i>)	5.6	42.2	15.3	3.2	-18.7	7.9
HSL-27	Huoshiliang	2017JHIT1007H35	Pig (<i>S. domestica</i>)	5.0	42.4	15.3	3.2	-9.6	10.0
HSL-33	Huoshiliang	2017JHIT1007H33	Pig (<i>S. domestica</i>)	0	-	-	-	-	-
HSL-29	Huoshiliang	2017JHIT1007H40	Dog (<i>Canis familiaris</i>)	10.9	42.9	15.6	3.2	-10.3	9.5
HSL-21	Huoshiliang	2017JHIT1007L6	Dog (<i>C. familiaris</i>)	6.5	42.6	15.6	3.2	-9.1	8.4
HSL-17	Huoshiliang	2017JHIT1007H19	Dog (<i>C. familiaris</i>)	3.5	42.2	15.3	3.2	-8.1	8.3
HSL-31	Huoshiliang	2017JHIT1007L3	Dog (<i>C. familiaris</i>)	0	-	-	-	-	-
HSL-12	Huoshiliang	2017JHIT1007M2	Cattle (<i>Bos taurus</i>)	6.3	42.2	15.4	3.2	-9.8	8.9
HSL-28	Huoshiliang	2017JHIT1007H38	Cattle (<i>B. taurus</i>)	9.8	43.0	15.6	3.2	-18.1	9.0
HSL-9	Huoshiliang	2017JHIT1007H17	Cattle (<i>B. taurus</i>)	0	-	-	-	-	-
HSL-22	Huoshiliang	2017JHIT1007H12L1	Sheep/goat (Ovicaprid)	6.1	41.7	15.0	3.2	-18.4	7.5
HSL-16	Huoshiliang	2017JHIT1007F4	Sheep/goat (Ovicaprid)	8.5	43.1	15.8	3.2	-18.6	9.1
HSL-10	Huoshiliang	2017JHIT1007L7	Sheep/goat (Ovicaprid)	8.6	42.7	15.5	3.2	-18.0(-17.6)	6.5 ^c
HSL-20	Huoshiliang	2017JHIT1007H38L2	Sheep/goat (Ovicaprid)	8.7	43.0	15.7	3.2	-17.4	6.5
HSL-14	Huoshiliang	2017JHIT1007M2	Sheep/goat (Ovicaprid)	8.8	43.0	15.7	3.2	-9.8	8.5
HSL-32	Huoshiliang	2017JHIT1007H21	Sheep (<i>Ovis aries</i>)	8.5	43.5	15.8	3.2	-16.3	8.3
HSL-26	Huoshiliang	2017JHIT1007L11	Sheep (<i>O. aries</i>)	3.5	43.6	15.9	3.2	-16.1	9.1
HSL-30	Huoshiliang	2017JHIT1007L8	Sheep (<i>O. aries</i>)	0	-	-	-	-	-
HSL-5	Huoshiliang	2017JHIT1007L9	Gazelle (<i>Gazella</i> sp.)	6.7	43.9	15.8	3.2	-15.2	8.9
HSL-24	Huoshiliang	2017JHIT1007L3	Gazelle (<i>Gazella</i> sp.)	9.6	43.3	15.7	3.2	-15.0	6.5
HSL-13	Huoshiliang	2017JHIT1007L6	Deer (Cervidae)	9.2	43.4	15.7	3.2	-18.8	4.8
HSL-34	Huoshiliang	2017JHIT1007H18	Deer (Cervidae)	0.2 ^b	39.3	13.1	3.5	-22.6	-
HSL-4	Huoshiliang	2017JHIT1007H2	Deer (Cervidae)	3.5	42.5	15.3	3.2	-18.9	4.1
HSL-3	Huoshiliang	2017JHIT1007H19	Deer (Cervidae)	2.7	42.5	14.9	3.3	-18.7	4.1
HSL-8	Huoshiliang	2017JHIT1007L3	Deer (Cervidae)	9.2	42.8	15.5	3.2	-19.9(-19.9)	4.2 ^c
HSL-6	Huoshiliang	2017JHIT1007L10	Hare (<i>Lepus capensis</i>)	7.7	42.5	15.4	3.2	-20.0(-20.0)	7.0 ^c
HSL-7	Huoshiliang	2017JHIT1007H26L2	Hare (<i>L. capensis</i>)	12.5	43.1	15.7	3.2	-19.7(-19.8)	6.7 ^c
HSL-2	Huoshiliang	2017JHIT1007M2	Middle bird (Aves)	5.6	41.5	15.0	3.2	-17.6(-17.6)	9.1 ^c
HSL-23	Huoshiliang	2017JHIT1007H30	Middle bird (Aves)	3.6	42.2	15.0	3.3	-22.6	4.4
HSL-1	Huoshiliang	2017JHIT1007H9	Middle bird (Aves)	6.1	43.0	15.6	3.2	-16.2	9.0
GGW-51	Ganggangwa	2017JGT0101H3	Pig (<i>S. domestica</i>)	6.5	42.7	15.4	3.2	-7.2	8.3
GGW-53	Ganggangwa	2017JGT0101L6	Pig (<i>S. domestica</i>)	8.2	42.9	15.3	3.3	-7.5	9.0
GGW-57	Ganggangwa	2017JGT1911L3	Pig (<i>S. domestica</i>)	5.0	43.3	15.5	3.3	-7.6	8.5
GGW-62	Ganggangwa	2017JGT1911L2	Pig (<i>S. domestica</i>)	5.7	43.3	15.7	3.2	-19.2	8.6
GGW-49	Ganggangwa	2017JGT0101H1	Dog (<i>C. familiaris</i>)	4.8	43.1	15.5	3.2	-11.0	9.3
GGW-48	Ganggangwa	2017JGT0101L5	Dog (<i>C. familiaris</i>)	6.6	42.2	15.0	3.3	-7.7	8.2
GGW-56	Ganggangwa	2017JGT1911L3	Dog (<i>C. familiaris</i>)	4.9	42.6	15.1	3.3	-10.4	9.6
GGW-59	Ganggangwa	2017JGT1911L2	Dog (<i>C. familiaris</i>)	7.2	41.6	14.9	3.3	-10.0	8.6
GGW-61	Ganggangwa	2017JGT1911L2	Dog (<i>C. familiaris</i>)	5.9	41.8	15.3	3.2	-10.6	8.6
GGW-36	Ganggangwa	2017JGT0101F1L2	Cattle (<i>B. taurus</i>)	7.0	44.1	14.6	3.5	-18.3	6.1
GGW-54	Ganggangwa	2017JGT1911L5	Cattle (<i>B. taurus</i>)	5.3	41.6	15.0	3.2	-18.2	7.9
GGW-60	Ganggangwa	2017JGT1911L6	Cattle (<i>B. taurus</i>)	6.7	43.4	15.6	3.2	-15.9	6.9
GGW-38	Ganggangwa	2017JGT0101L6	Sheep/goat (Ovicaprid)	6.2	42.2	15.3	3.2	-17.4	8.2
GGW-40	Ganggangwa	2017JGT0101L8	Sheep/goat (Ovicaprid)	9.2	43.3	14.7	3.4	-17.6	8.0
GGW-45	Ganggangwa	2017JGT0101L2	Sheep/goat (Ovicaprid)	5.2	41.7	15.0	3.2	-17.7	7.1
GGW-44	Ganggangwa	2017JGT0101L5	Sheep/goat (Ovicaprid)	7.1	42.2	15.2	3.2	-21.2	8.5
GGW-46	Ganggangwa	2017JGT1911L3	Sheep/goat (Ovicaprid)	5.7	43.3	15.6	3.2	-16.5	7.2
GGW-55	Ganggangwa	2017JGT1911L3	Sheep/goat (Ovicaprid)	5.9	42.2	15.3	3.2	-17.6	6.3
GGW-58	Ganggangwa	2017JGT1911F2	Sheep/goat (Ovicaprid)	3.0	43.3	14.8	3.4	-16.9	10.7

(Continued)

TABLE 1 | (Continued)

Lab ID	Site name	Context ^a	Taxon	Collagen (%)	C (%)	N (%)	C/N Ratio	δ ¹³ C (‰)	δ ¹⁵ N (‰)
GGW-47	Ganggangwa	2017JGT0101H2	Sheep/goat (Ovicaprid)	0	–	–	–	–	–
GGW-35	Ganggangwa	2017JGT0101L1	Gazelle (<i>Gazella</i> sp.)	6.8	42.8	15.5	3.2	–13.4	7.1
GGW-50	Ganggangwa	2017JGT0101H3	Gazelle (<i>Gazella</i> sp.)	5.4	41.9	15.1	3.2	–15.2	6.3
GGW-37	Ganggangwa	2017JGT0101L6	Gazelle (<i>Gazella</i> sp.)	4.5	42.5	15.3	3.2	–18.4	8.6
GGW-43	Ganggangwa	2017JGT1911L3	Deer (Cervidae)	6.0	43.2	15.6	3.2	–19.2	4.8
GGW-42	Ganggangwa	2017JGT1911L2	Deer (Cervidae)	6.3	42.2	15.3	3.2	–18.8	3.0
GGW-52	Ganggangwa	2017JGT0101L6	Hare (<i>L. capensis</i>)	6.2	41.8	15.2	3.2	–19.6	9.0
GGW-41	Ganggangwa	2017JGT1911L3	Hare (<i>L. capensis</i>)	6.3	43.4	15.6	3.2	–20.0	9.8
GGW-39	Ganggangwa	2017JGT0101L6	Middle bird (Aves)	7.4	43.1	15.6	3.2	–20.4	5.2

Some issues are specially labeled: (a) archeological context of each sample confirmed during the excavation while T stands for trail trench, L for layer, H for ash pit, F for house site, and M for burial; (b) this sample was excluded from further data analysis due to its extremely low collagen yield (< 0.5%); (c) δ¹⁵N measurement of these five samples were missed during the analysis in Lanzhou University, and was supplemented at Beta Analytic while δ¹³C measured by Beta Analytic are listed with brackets after as contrast.



on the latter (Gavin, 2001; Dong et al., 2014; Yang et al., 2019a). The conventional age of these 15 samples ranges from 3,610 to 3,480 BP, while the calibrated age 2σ ranges between 4,070 and 3,650 cal BP, basically consistent with previous studies (Dodson et al., 2009; Atahan et al., 2011).

The radiocarbon dating results reported in Ren et al. (2022) and this study focused on not only the starting and ending ages of the faunal remains, but also the precise age of the individuals with abnormal stable carbon isotope values. The cattle and sheep/goat individuals with C₄ values, as well as two of the three pig individuals with C₃ values, were chosen for radiocarbon dating. As the result shows, all radiocarbon dates of samples with abnormal δ¹³C-values fall into the latter half (generally later than ca. 3,850 cal BP) of the sites. Within the context of archeological features, the age of the only C₃ pig individual without direct radiocarbon date falls into the latter half of the sites' occupation,

as decided by the two radiocarbon dates derived from the same archeological unit (the back fill of burial M2 in Huoshiliang).

DISCUSSION

Diversified Livestock Feeding Pattern in Huoshiliang and Ganggangwa Sites

As indicated by radiocarbon dating results and consecutive archeological deposits, animal resources were continuously utilized in Huoshiliang and Ganggangwa sites during 4,070–3,650 cal BP (calibrated age, 2σ range) (Table 3). Although the upper limit of the calibrated age 2σ approaches nearly 4,100 cal BP, the majority of the faunal remains are concentrated between 4,000 and 3,700 cal BP (Table 3), which falls into the time period of Xichengyi Culture as refined by a previous study

TABLE 2 | Statistics of stable carbon and nitrogen isotope data reported in this study.

Sort	Sample size (n)	$\delta^{13}\text{C}$ Max (‰)	$\delta^{13}\text{C}$ Min (‰)	$\delta^{13}\text{C}$ Mean (‰)	$\delta^{13}\text{C}$ SD 1 σ (‰)	$\delta^{15}\text{N}$ Max (‰)	$\delta^{15}\text{N}$ Min (‰)	$\delta^{15}\text{N}$ Mean (‰)	$\delta^{15}\text{N}$ SD 1 σ (‰)
Fauna	56	-7.2	-22.7	-15.3	4.4	10.8	3.0	7.7	1.7
Livestock	37	-7.2	-21.2	-13.8	4.5	10.8	6.1	8.3	1.0
Omnivore	18	-7.2	-19.2	-10.6	3.8	10.1	7.8	8.7	0.6
Herbivore	19	-9.8	-21.2	-16.8	2.7	10.8	6.1	7.9	1.2
Wildlife	19	-13.4	-22.7	-18.3	2.2	9.8	3.0	6.5	2.1
Pig	10	-7.2	-19.2	-11.4	4.8	10.1	7.8	8.6	0.6
Pig C ₃	3	-17.8	-19.2	-18.6	0.6	9.0	7.9	8.5	0.5
Pig C ₄	7	-7.2	-9.6	-8.2	0.9	10.1	7.8	8.6	0.7
Dog	8	-7.7	-11.0	-9.7	1.2	9.6	8.2	8.8	0.5
Cattle	5	-9.8	-18.3	-16.1	3.2	9.0	6.1	7.8	1.1
Sheep/goat	14	-9.8	-21.2	-17.1	2.4	10.8	6.3	8.0	1.2
Gazelle	5	-13.4	-18.5	-15.5	1.6	8.9	6.3	7.5	1.1
Deer	6	-18.7	-19.9	-19.0	0.4	4.8	3.0	4.2	0.6
Hare	4	-19.6	-20.0	-19.8	0.2	9.8	6.7	8.1	1.3
Bird	4	-16.2	-22.7	-19.2	2.5	9.1	4.4	6.9	2.1

TABLE 3 | Radiocarbon dating results of animal bones excavated from Huoshiliang and Ganggangwa sites.

Lab code	Site name	Context	Material	CollagenLab ID	Taxon	AMS ¹⁴ C age(BP)	Calibrated age 2 σ (cal BP)	Mean and σ (BP)
LZU20279	Huoshiliang	2017JHIT1007H9	Bone collagen	HSL-11	Pig	3,500 ± 20	3,840–3,690	3,770 ± 40
LZU20280 ^a	Huoshiliang	2017JHIT1007M2	Bone collagen	HSL-12	Cattle	3,550 ± 20	3,910–3,720	3,830 ± 50
Beta-567009 ^a	Huoshiliang	2017JHIT1007M2	Bone collagen	HSL-14	Sheep/goat	3,500 ± 30	3,870–3,650	3,770 ± 50
LZU18134	Huoshiliang	2017JHIT1007F4	Bone collagen	HSL-16	Sheep/goat	3,495 ± 25	3,840–3,690	3,770 ± 40
Beta-567430	Huoshiliang	2017JHIT1007L3	Bone collagen	HSL-24	Gazelle	3,530 ± 30	3,900–3,690	3,800 ± 50
LZU20404	Huoshiliang	2017JHIT1007L10	Bone collagen	HSL-25	Pig	3,480 ± 20	3,840–3,690	3,760 ± 40
LZU18135 ^a	Huoshiliang	2017JHIT1007L11	Bone collagen	HSL-26	Sheep	3,520 ± 20	3,880–3,700	3,780 ± 40
Beta-567010 ^a	Ganggangwa	2017JGT0101L8	Bone collagen	GGW-40	Sheep/goat	3,610 ± 30	4,070–3,830	3,920 ± 50
Beta-567431	Ganggangwa	2017JGT0101L2	Bone collagen	GGW-45	Sheep/goat	3,510 ± 30	3,880–3,690	3,780 ± 50
LZU20281	Ganggangwa	2017JGT0101L5	Bone collagen	GGW-48	Dog	3,600 ± 20	3,980–3,840	3,910 ± 40
LZU20405	Ganggangwa	2017JGT0101L6	Bone collagen	GGW-53	Pig	3,580 ± 20	3,970–3,830	3,880 ± 40
Beta-567011 ^a	Ganggangwa	2017JGT1911L3	Bone collagen	GGW-55	Sheep/goat	3,510 ± 30	3,880–3,690	3,780 ± 50
LZU20282	Ganggangwa	2017JGT1911L2	Bone collagen	GGW-59	Dog	3,600 ± 20	3,980–3,840	3,910 ± 40
LZU20283	Ganggangwa	2017JGT1911L6	Bone collagen	GGW-60	Cattle	3,580 ± 20	3,970–3,830	3,880 ± 40
LZU20284 ^a	Ganggangwa	2017JGT1911L2	Bone collagen	GGW-62	Pig	3,530 ± 20	3,890–3,710	3,800 ± 50

Conventional ages were calibrated by the IntCal20 curve (Reimer et al., 2020) via the OxCal 4.4.4 online program (Bronk Ramsey, 2021). Data with special label (a) have been previously reported in Ren et al. (2022).

(Yang et al., 2019a). Moreover, the two earliest conventional ages dated from surficial charcoal in Ganggangwa site reported in Dodson et al. (2009) and Atahan et al. (2011), which are nearly 100 years earlier than other results, may have been influenced significantly by the “old wood” effect (Gavin, 2001; Dong et al., 2014; Yang et al., 2019a), thus should be regarded more cautiously.

For wildlife, the relatively wide distribution in both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ -values reflects their diverse habitats and diets (Figure 3 and Table 2). Stable isotope values of birds were the most distributed among the wildlife (Figure 3B), consistent with the diversified living space and varied diets of these creatures. Among wild herbivores, hare and deer present consistent diets in consuming prominent C₃ plants as revealed by $\delta^{13}\text{C}$ -values, which can reflect the C₃ plant-dominated botanic situation of their habitat. However, the significant distinction in $\delta^{15}\text{N}$ -values between hare

and deer could be attributed to different extents of forest cover in their respective habitats and/or the metabolic variation under different aridity (Ambrose and DeNiro, 1986; Sealy et al., 1987; Schwarcz et al., 1999; Hartman, 2011; Hofman-Kamińska et al., 2018), since the trophic levels of these species are similar. These hares are more likely to inhabit arid grassland, while the deer are more likely to inhabit moist forest in the piedmont of the Qilian Mountains. The gazelles presented generally mixed C₃/C₄ signal in $\delta^{13}\text{C}$ -values ($-15.5 \pm 1.6\text{‰}$), which is different from other herbivorous wildlife reported in this study (Figure 3). The relatively positive $\delta^{13}\text{C}$ -values of these gazelles (about 3.5‰ higher than the deer and slightly higher than the cattle and sheep/goat) demonstrate notable intake of C₄ plants and/or ecological influences such as the canopy effect and watering effect (e.g., Drucker et al., 2008; Wallace et al., 2013). Though some scholars argued that wild C₄ plants in the arid desert constitute

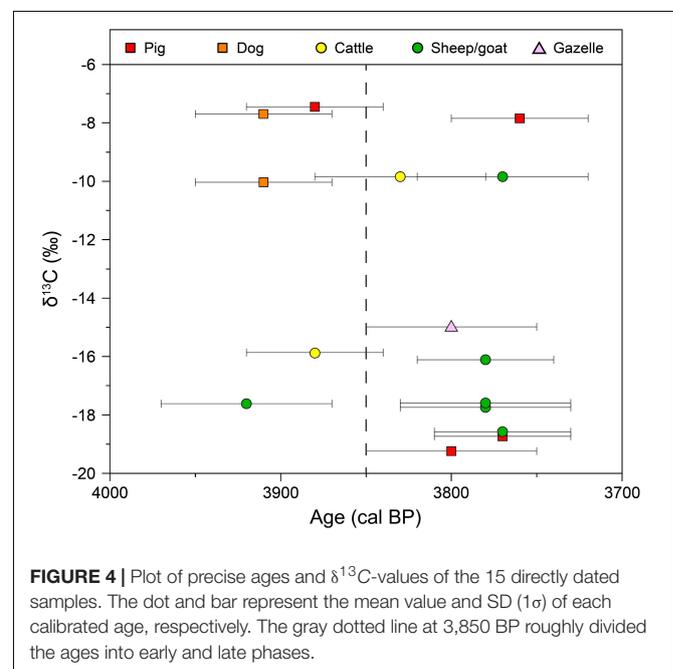
the C_4 intake of wild herbivores in Hexi Corridor (Ma et al., 2021), it is impossible to eliminate the potential contribution of C_4 crops in these gazelles' diets. Moreover, the similarity of $\delta^{15}N$ -values between gazelles and herbivorous livestock (Figure 3B) may also indicate their overlapped living space. A case study on African grasslands revealed that prehistoric human activity had shaped the savanna landscapes *via* enriching nutrient for the local ecosystem (Marshall et al., 2018), which indicates that human settlement in arid area could make the perimeter zone more nutritious for wildlife to inhabit. Furthermore, the "inadvertent baiting" from human cultivated crops on wild herbivore was hypothesized and explored in eastern North America (Svizzero, 2016; Bonzani et al., 2021), which has shed light on the relation between agricultural groups and the neighboring wild animals. Thus, we presented a conjecture that the massive cultivation of millet crops during this period around the human settlements attracted the gazelles to forage around and accidentally got hunted. This speculation also describes a mutual beneficial relation between human and wildlife, though still requiring further validation.

For livestock, the $\delta^{13}C$ -values fall into two distinct groups, while the $\delta^{15}N$ -values are more concentrated relative to wildlife (Figure 3). Carbon isotope data suggest that all of the domestic dogs (8/8) and the majority of the domestic pigs (7/10) sampled in Huoshiliang and Ganggangwa consumed C_4 diets (Figure 3 and Table 1), indicating the consumption of millet, such as millet grains, millet by-products, and/or human leftovers, which is consistent with the tradition of Neolithic cultures in northwestern China (e.g., Barton et al., 2009; Chen et al., 2016; Ma et al., 2021). For the non-indigenous herbivorous livestock, most of the domestic cattle (4/5) and domestic sheep/goat (13/14) exhibited carbon isotope values indicative of a C_3 diet with a minor contribution of C_4 plants (Figure 3 and Table 1). With the exception of two individuals exhibiting high $\delta^{13}C$ -values (one cattle and one sheep/goat sample), the results are consistent with previous studies in the EBA of the Hexi Corridor and can be attributed to the consumption of wild C_3 plants during herding (Atahan et al., 2011; Ma et al., 2021; Vaiglova et al., 2021). Wild C_4 plants in arid desert and/or deliberate feeding of millet by-products were pointed out to constitute the intake of C_4 foodstuff of herbivorous livestock in Hexi Corridor (Su and Yan, 2008; Ma et al., 2021). The $\delta^{15}N$ -values for domestic pigs and dogs were more concentrated and slightly higher ($8.7 \pm 0.6\text{‰}$) than values for herbivorous livestock ($7.9 \pm 1.2\text{‰}$). The close $\delta^{15}N$ -values between omnivorous livestock and herbivores indicate limited protein consumption by these omnivorous livestock. The broader distribution of $\delta^{15}N$ -values for herbivorous livestock could result from grazing in different ecological zones (e.g., Pearson et al., 2007; Atahan et al., 2011) since the natural environment along the Heihe River valley as well as the surrounding regions in Hexi Corridor varies significantly from alpine meadow and coniferous forest to desert grassland and desert. The more concentrated $\delta^{15}N$ -values of domestic pig and dog indicate a relatively unified living environment and feeding pattern adopted by local people, which might involve captive breeding and deliberate feeding.

Notably, several individuals among the livestock have abnormal $\delta^{13}C$ -values; some herbivorous individuals (one cattle

and one sheep/goat) appear to have consumed a C_4 diet while some omnivorous individuals (three pigs) appear to have consumed a C_3 diet (Figure 3). The C_3 signal in omnivorous individuals of these two sites has been previously detected and was attributed to the consumption of wild foods (Atahan et al., 2011). The same phenomenon of C_3 -consuming pigs in coetaneous Xichengyi site was regarded as the result of free-ranging and/or inhabitation site raising with C_3 consumption (Ma et al., 2021). However, considering the open arid modern situation and the low productivity of the ecosystem around these two sites, the feasibility of free-ranging pigs remains questionable. Moreover, the abnormal C_4 signal of herbivorous livestock has not been well explained yet. It is likely that the existence of these abnormal livestock individuals relates to the transformation of animal husbandry strategies. However, some further investigation of geochemical tracing such as Sr isotope and trace element analyses on these abnormal individuals would be meaningful to examine the issue of local/extra-local provenance since the possibility of livestock's immigration could not be eliminated.

As indicated by direct radiocarbon dating results supplementary with stratigraphic evidence, a remarkable diversification in $\delta^{13}C$ -values of livestock during the latter half of the occupation at these two sites has been revealed. The relation between calibrated radiocarbon ages (shown as mean value and SD) and $\delta^{13}C$ -values of the 15 directly dated samples is plotted in Figure 4. The ages of all individuals consuming abnormal diet fall into the latter half of these two sites, basically later than *ca.* 3,850 cal BP. Radiocarbon dating results supplementary with stratigraphic evidence also reveal that livestock consuming conventional diets existed in both former and latter halves of these two sites' occupation (Figure 4 and Table 1). Thus, 3,850 BP is possibly the timing for the initial diversification in livestock



feeding pattern at Huoshiliang and Ganggangwa sites. According to archaeobotanical findings, the latter half of the occupation at the two sites (*ca.* 3,850–3,700 BP) witnessed the promotion of wheat and barley cultivation in Hexi Corridor, with the proportion of these non-indigenous crop seeds reached up to roughly a quarter among all crop seeds (Zhou et al., 2016; Yang, 2017). Stable isotope evidence also reveals the consumption of C₃ crops by humans since 4,000–3,800 BP (Liu et al., 2014). This evidence proves the potential for humans in the Huoshiliang and Ganggangwa sites to provide their livestock with these non-indigenous C₃ crops. Furthermore, pigs consumed a similar diet to humans (Liu and Jones, 2014), and the appearance of C₃ consumption in their diets was synchronous with the preliminary shift with increased C₃ intake in human diets. Additionally, the individuals with abnormal $\delta^{13}\text{C}$ -values nonetheless have $\delta^{15}\text{N}$ -values that are similar to the majority of omnivorous livestock (Figure 3), suggesting similar living environment and feeding patterns. Thus, we speculated that the domestic pigs with C₃ signal were more likely to be raised at their inhabitation site, instead of free-ranging, and were deliberately fed with C₃ foodstuff, such as wild plants and/or the products or human leftovers of wheat/barley. Similarly, herbivorous individuals with C₄ signal are likely to be adopted with the same management as the omnivorous livestock, including captive breeding and deliberate feeding of C₄ plants, most likely the products or human leftover of millet crops.

In summary, the absolute age of the faunal remains from Huoshiliang and Ganggangwa sites ranges from 4,070 to 3,650 cal BP (2σ), but are more concentrated between 4,000 and 3,700 cal BP (Table 3 and Figure 4). The appearance of both omnivorous and herbivorous livestock with abnormal $\delta^{13}\text{C}$ -values during the latter half of the sites' occupation reveals a diversified livestock feeding pattern adopted by humans. It is likely that some herbivorous livestock were raised at their inhabitation site and deliberately fed with millet products, while a small group of domestic pigs were deliberately fed with C₃ foodstuff, involving generally wild C₃ plants and/or products or human leftovers of wheat and barley. The initial timing for the diversification in livestock feeding pattern is circa 3,850 BP.

The Trajectory for the Transformation of Livestock Feeding Pattern in Prehistoric Hexi Corridor

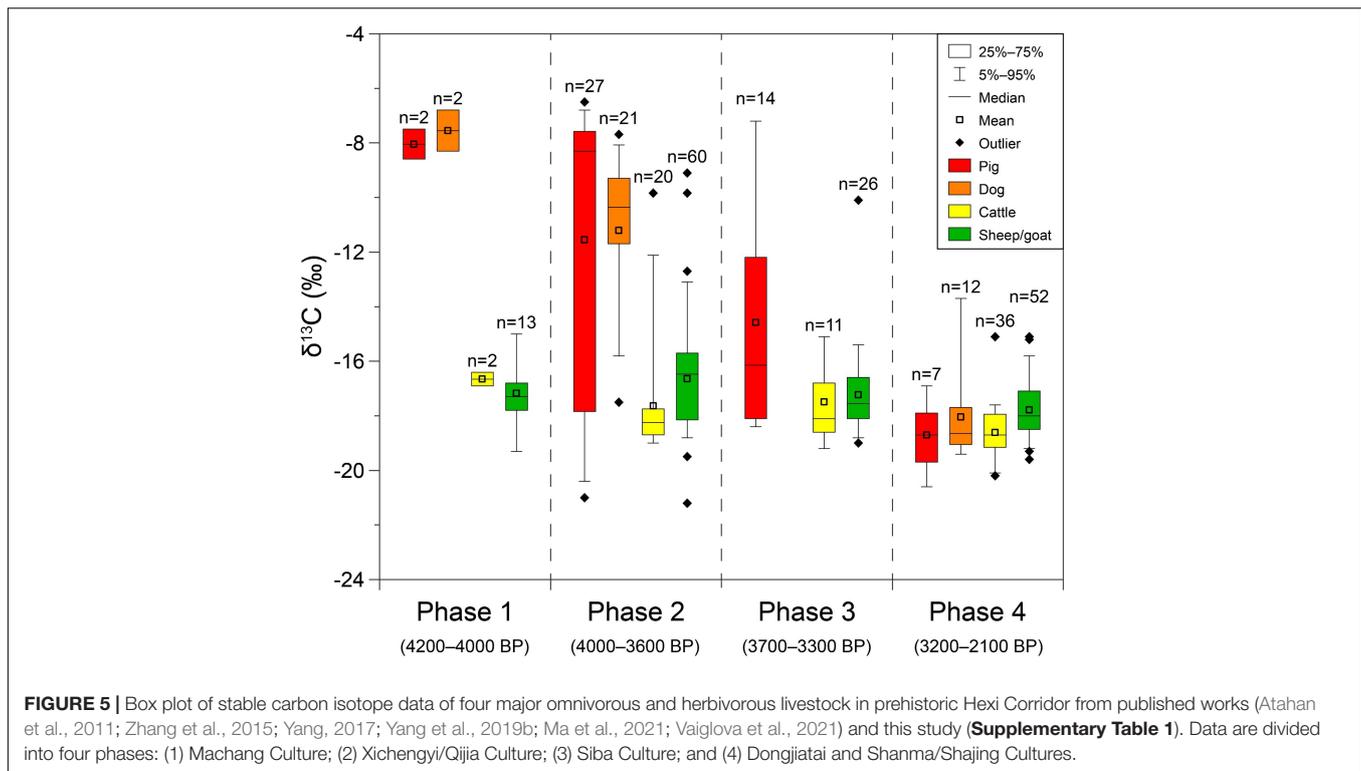
To obtain clearer insight into the temporal variation of livestock feeding pattern in prehistoric Hexi Corridor, published stable isotope data from faunal remains (Atahan et al., 2011; Zhang et al., 2015; Yang, 2017; Yang et al., 2019b; Ma et al., 2021; Vaiglova et al., 2021) were combined with data presented in this study (Supplementary Table 1). Stable carbon isotope data from four categories of major livestock involved in our discussion were charted as box plot and distinguished *via* cultural affiliations of different site and the material (Figure 5). The cultural affiliations of the materials reported in their original studies involved in Supplementary Table 1 and Figure 5 were reconfirmed through published archeological excavation reports, and the absolute age of each cultural period was determined

via their refined chronology (Yang et al., 2019a). According to the report of excavation in Xihetan site (Zhao, 2005) and recent archeological findings in other Xichengyi Culture sites, the cultural affiliation of this site has been refined as Xichengyi Culture (*ca.* 4,000–3,600 BP). The cultural affiliation of animal stable isotope data from Huoshaogou site (Vaiglova et al., 2021) has been refined as Shanma Culture referring to recently published excavation reports (Wang et al., 2019, 2021). Data collected in Supplementary Table 1 were then divided into four phases in accordance with their cultural affiliations to investigate a more detailed shifting trajectory of livestock $\delta^{13}\text{C}$ -values. An overlap of time between phase 2 and 3 occurred due to the overlap between Xichengyi/Qijia and Siba Cultures (Yang et al., 2019a).

The population of Majiayao, a type of Late Neolithic culture originated in the upper Yellow River Valley in northwestern China, is currently known as the earliest agricultural immigrants to expand westward and to enter Hexi Corridor at *ca.* 4,800 BP (Dong et al., 2018), brought in a pattern of subsistence that mainly engaged in millet cultivation and pig-dog feeding (Yang, 2017; Yang et al., 2019b). In the following eight centuries, the Late Neolithic culture in Hexi Corridor evolved into Banshan Culture (*ca.* 4,450–4,200 BP) and Machang Culture (*ca.* 4,200–4,000 BP) in succession (Yang et al., 2019a).

During the period of Machang Culture (*ca.* 4,200–4,000 BP), the eastern-originated foxtail millet and broomcorn millet were major crops and staple foods in the diets of humans and omnivorous livestock, as indicated by archaeobotanical and stable isotope data (Liu et al., 2014; Zhou et al., 2016; Yang et al., 2019b). This pattern of human subsistence was consistent with coetaneous and earlier societies in the neighboring Gansu-Qinghai region (Barton et al., 2009; Jia et al., 2013; Ma et al., 2016, 2021; Ren et al., 2020). Cattle and sheep/goat, the non-indigenous herbivorous livestock, were estimated to be introduced into Hexi Corridor during this period, though direct radiocarbon dating results were slightly younger (Yang, 2017; Yang et al., 2019b; Ren et al., 2022). The $\delta^{13}\text{C}$ -values of these early cattle and sheep/goat revealed a C₃ plant-dominated consumption, which is consistent with the natural vegetation (Yang et al., 2019b; Figure 5). The feeding patterns of both indigenous omnivorous livestock and novel herbivorous livestock during this period in Hexi Corridor are revealed to be consistent with their traditions, respectively.

The period of time during Xichengyi/Qijia Cultures (*ca.* 4,000–3,600 BP) witnessed the phenomenal diversification in livestock feeding patterns in Hexi Corridor, with significant expansion of $\delta^{13}\text{C}$ -values among both herbivorous and omnivorous livestock (Figure 5). As revealed by archaeobotanical evidence, wheat and barley were introduced into Hexi Corridor and widely spread during this period (Dodson et al., 2013; Zhou et al., 2016; Yang, 2017). Charred seeds of these Western-originated crops were commonly unearthed from contemporaneous archeological sites, reaching a proportion up to circa a quarter among all crop remains (Zhou et al., 2016; Yang, 2017). Correspondingly, the contribution of C₃ crops in human diet emerged and gradually increased during this period (Liu et al., 2014, 2016). The importance of herbivorous livestock also increased (Yang, 2017; Yang et al., 2019b), together with the increased practice of pastoralism (Atahan et al., 2011;



Ma et al., 2021). As shown in **Figure 5**, the consumption of C_3 foodstuff in omnivorous livestock enhanced significantly. The presence of several domestic omnivorous individuals that mainly consumed C_3 plants (**Supplementary Table 1**) is suggestive of deliberate feeding with C_3 crop by-products and/or allowing the animals to free range (Ma et al., 2021). In contrast, the general dietary shift of herbivorous livestock was not that prominent (**Figure 5**). Domestic herbivorous individuals with dominant intake of C_4 plants are likely to be raised at inhabitation site and deliberately fed with millet products, similar to the tradition of raising pigs and dogs (e.g., Barton et al., 2009; Chen et al., 2016; Ma et al., 2021). This bilateral transformation of feeding patterns between domestic omnivores and domestic herbivores jointly constructed the diversification in livestock feeding patterns during EBA in Hexi Corridor. Combined with the result shown in this study, the initial timing for this transformation could possibly be *ca.* 3,850 BP (**Figure 3**). Moreover, the motivation behind such attempt to partly break with the feeding traditions for both indigenous and novel livestock may involve the exploration of new animal husbandry manners to realize a more efficient economy and better social sustainability, and to withstand the harsh arid living environment in Hexi Corridor.

When it came to the Siba Culture (*ca.* 3,700–3,300 BP), the diversification in livestock feeding pattern persisted (**Figure 5**). During this period, wheat and barley became the primary agricultural crops (Zhou et al., 2016; Yang, 2017; Yang et al., 2019b). Meanwhile, cattle and sheep/goat became primary livestock (Yang, 2017; Yang et al., 2019b). The consumption of C_3 foodstuff by humans and livestock also increased significantly

during this period (Liu et al., 2014; Yang, 2017; Yang et al., 2019b), consistent with the neighboring Gansu-Qinghai region in northwestern China (Ma et al., 2016; Vaiglova et al., 2021). As revealed by **Figure 5**, a reduction of C_4 food consumption in domestic pigs took place. The majority of domestic pigs shifted to C_3 -based diets, making the C_4 consumers the minority. The dietary tendency of domestic dogs during this period remains unknown due to the absence of stable isotope data. The diet of herbivorous livestock at this phase generally returned to the initial status with C_3 consumption dominated. Only a single sheep/goat individual remained consuming mainly C_4 plants (**Figure 5**) during this period, indicating the general termination of dietary diversification among herbivorous livestock.

The $\delta^{13}C$ -values of phase 4 (*ca.* 3,300–2,100 BP) demonstrate generally C_3 -based diets among all major livestock (**Figure 5**). This situation is consistent with the popularization of wheat and barley, as well as the fall of millet-based agriculture in northern China (e.g., Dodson et al., 2013; Chen et al., 2015, 2020; Liu et al., 2017). The dietary shift from C_4 -based to C_3 -based among omnivorous livestock seems to have been accomplished since *ca.* 3,200 BP (**Figure 5**). Meanwhile, the diet of herbivorous livestock returned to C_3 -dominated as well. For the last millennium before the common era, the rise and expansion of nomadism deeply influenced the Eurasian Steppe and northwestern China (Kuz'mina, 2008). The agro-pastoral residents in Hexi Corridor established more developed pastoral economy after *ca.* 2,800 BP, which are known as Shanma Culture (*ca.* 2,900–2,100 BP) in western Hexi Corridor and Shajing Culture (*ca.* 2,700–2,100 BP) in eastern Hexi Corridor (Li, 2011; Yang et al., 2019a).

In summary, the trajectory for the transformation of livestock feeding patterns in the prehistoric Hexi Corridor has been revealed as follows: (1) diets of both omnivorous and herbivorous livestock conformed with their original traditions during the Machang Culture (*ca.* 4,200–4,000 BP); (2) during the Xichengyi/Qijia Culture (*ca.* 4,000–3,600 BP), the range of $\delta^{13}\text{C}$ -values exhibited by both omnivorous and herbivorous livestock expanded dramatically, indicating a diversified livestock feeding pattern during EBA; (3) during the Siba Culture (*ca.* 3,700–3,300 BP), herbivorous livestock generally returned to a C_3 diet, but domestic pigs continued to consume a wide array of C_3 – C_4 food products; and finally (4) the transformation of livestock feeding patterns seems to have fully completed during the Dongjiatai and Shanma/Shajing Cultures (*ca.* 3,200–2,100 BP), with all four major livestock groups in the Hexi Corridor consuming C_3 diets.

CONCLUSION

In this study, systematic stable carbon and nitrogen isotope analysis and precise radiocarbon dating were performed on faunal remains acquired through archeological excavation of two EBA settlements in middle Hexi Corridor, Huoshiliang, and Ganggangwa sites. In total, stable carbon and nitrogen isotope data were acquired for 56 faunal samples and 15 direct radiocarbon dates were obtained. The stable isotope results of domestic animals reveal a significant diversification in livestock feeding pattern, with some domestic pigs consuming mainly C_3 foodstuff and some herbivorous livestock consumed mainly C_4 plants, respectively. It seems that humans in these two sites attempted to deliberately feed some of the herbivorous livestock following the tradition in feeding omnivorous livestock; and to deliberately feed some of the domestic pigs with C_3 foodstuff, possibly involving wheat/barley products. Supplemented by stratigraphic evidence, the precise radiocarbon dating results indicate that the initial timing for the diversification in livestock feeding pattern lies at *ca.* 3,850 BP.

Integrated with the summary of published animal stable isotope data, a vivid animation for the transformation of livestock feeding pattern during prehistoric times in Hexi Corridor was exhibited. By adopting more precise phase division, the trajectory of the transformation in livestock feeding pattern was further clarified. A notable diversification in livestock feeding pattern during EBA, which is consistent with the pattern revealed in Huoshiliang and Ganggangwa sites, has been detected. Such diversification existed in the transformation from millet-based agricultural society to wheat/barley-based agro-pastoral society in prehistoric Hexi Corridor, and possibly took place at *ca.* 3,850 BP. We argued that the motivation behind such attempt to partly break with the feeding traditions involves the exploration to innovate animal husbandry strategies. Such exploration may

be effective to improve the economic efficiency and the social sustainability, which can help the local people to withstand the harsh arid living environment in Hexi Corridor.

DATA AVAILABILITY STATEMENT

The original contributions presented in this study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding authors.

ETHICS STATEMENT

Ethical review and approval was not required for the animal study because this study only involves ancient animal skeletal remains (also known as a kind of fossil remain), which is totally distinct from the study of living animals. Thus, we confirm that there is no need of ethical review.

AUTHOR CONTRIBUTIONS

LR and GC designed the study. MQ, SZ, YY, and GC participated in the fieldwork. MQ, HL, ML, RL, and LR conducted the experiment and data analysis. MQ, SZ, YY, HL, and LR wrote the manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fevo.2022.908131/full#supplementary-material>

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