The domestication of South American camelids: a review

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Introduction

South American camelids are the only domesticated ungulates in the Americas, and the Andean region sustained the only pastoralist societies in the pre-Hispanic New World. South American Camelids are composed by two genera and four species, two of them wild (vicuñas *Vicugna vicugna* Molina, 1782, and guanacos *Lama guanicoe* Müller, 1776) and two domestic (llamas *Lama glama* Linnaeus, 1758 and alpacas *Lama pacos* Cuvier, 1800 suggested *Vicugna pacos* Linnaeus, 1758 by Wheeler et al., 2006) (Figure 1).

Guanacos have a broad geographic distribution across a variety of open habitats (arid, semiarid, hilly, mountain, steppe) and temperate forest environments. Their social structure reflects this wide distribution, with some plasticity in types of groups. In the breeding season, the guanaco social structure comprises three basic units: territorial family groups, nonterritorial male groups, and solitary individuals. In turn, mixed groups are common during the winter and in some populations migration occurs. Family group territoriality in the breeding season is correlated with stable food supply (Vilá, 2012: 64). Vicuñas live only in high-altitude Puna environments above 3,400 m in Peru, Bolivia, Argentina, and Chile. They are adapted to open grasslands and steppes; although they prefer to graze in the humid wetlands or marshes (vegas), due to the presence of livestock in these wetlands, vicuñas are usually found in the steppes. Vicuñas live in family groups consisting of one male, three to four females, and two offspring, and in bachelor groups. Family groups are stable and territorial all year round (Vilá, 2012: 42–43).

During pre-Hispanic times, the domestic llamas were circumscribed to the Andean regions of Peru, Bolivia, Chile, and Argentina, but alpacas had a more restricted habitat in the high and humid punas (bofedales) of Peru, Bolivia, and northern Chile.

The domestication of camelids was a complex process associated with the adaptations of hunter-gatherer groups to environmental fragmentation, caused by increased aridity during the Mid-Holocene and the consequent loss of productive habitats in the region (Yacobaccio et al., 2017). During this period, hunter-gatherer groups adopted a logistic strategy, reducing their residential mobility and introducing technological innovations. They developed communal hunts of wild camelids, made possible by population aggregation during their annual cycle, and opted for specialized hunting of camelids as their main source of food (Aschero and Martinez, 2001). Table 1 summarizes the correlated changes in the environment and in human populations by period. These changes did not happen all at once across the region; the geography of the origins of animal domestication and social complexity can be best described as a mosaic pattern.

Domestication is a process of interaction between an animal species and humans. Darwin (1868) made explicit that domestication includes the raising of animals in captivity that can occur without a conscious effort on the part of people and increases animal fertility, allowing them to have greater plasticity. According to Price (1984, 2002), domestication is an evolutionary process marked by the genotypic adaptation of animals to the captive environment.

Implications

- Appearance of individuals larger than the current llama (earliest evidence around 7100 cal. BP, increasing between 5800 and 4200 cal. BP).
- Detection of human impact due to environmental management practices that suggests more intensive human intervention in the environment since ca. 5300–4869 cal. AP.
- Detection of pathologies indicative of human handling in bones of the extremities and vertebrae since ca. 4900 cal. AP.
- First appearance of corrals in caves or in stone structures in deep ravines on the edge of valleys between 4500 and 3639 cal. AP.
Figure 1. The four camelids: taxonomy, subspecies, breeds, and original distribution. Salka and Uywa are the Quechua names for wild and domestic, respectively.

**Guanaco**  
*Wild-Salka*  
*(Lama guanicoe)*  
Two subspecies:  
*Lama guanicoe casicsensis* (north)  
*Lama guanicoe guanicoe* (south)  
Weight 100-120 kgs.  
Height at the withers 120 cms.

**Vicuña**  
*Wild- Salka*  
*(Vicugna vicugna)*  
Two subspecies:  
*Vicugna vicugna mensalis* (north)  
*Vicugna vicugna vicugna* (south)  
Weight 45 kgs.  
Height at the withers 90 cms.

**Llama**  
*Domestic-Uywa*  
*(Lama glama)*  
Two breeds:  
*a - Q’ara* (tall and strong, short woolled)  
*b - Tampulli* (hairly, long woolled)  
Weight 100-130 kgs.  
Height at the withers 110 cms.

**Álcpaca**  
*Domestic-Uywa*  
*(Lama pacos, suggested Vicugna pacos)*  
Two breeds:  
*Suri* (long straight fiber)  
*Huacaya* (wavy fiber)  
Weight 60 kgs.  
Height at the withers 90 cms.
animal is one whose mate selection is influenced by humans and whose docility and tolerance to humans are genetically determined.

The first step of the domestication process is based on the relationship of a phenotypically plastic species habituated to human presence, and occurs before any genotypic change; therefore, it can last a long time. This process, called the Baldwin effect, is an evolutionary transition from a facultative tolerance to humans toward a dependence on them; at this stage, the animal population becomes accustomed to the human presence and a selection mechanism for docility begins to function (Crispo, 2007; Francis, 2015).

For llama domestication, a multilocation model has been developed that includes two phases: herd protection and captivity-selective breeding (Yacobaccio and Vilá, 2016). Herd protection refers to human intervention in guanaco populations, or population subgroups, whose individuals are protected from its nonhuman predators and are facilitated access to feeding areas. A second step is the captivity and selective breeding of certain individuals. In the phase of herd protection, channeled by the Baldwin effect, people are a neutral stimulus. In the second step, when people become a positive stimulus—usually associated with the presence of food or shelter—an associative kind of learning emerges that goes beyond habituation and generates the taming process. Tameness is a condition for reproductive manipulation, as well as for the isolation of individuals in confinement or captivity. This step involves a greater degree of handling and isolation, meaning the existence of a physical barrier between wild population and captive herds. The space constraint increases animal density, resulting in changes in the social structure of the group of camels and triggering genetic adaptation to captivity (Yacobaccio and Vilá, 2016: 10–11).

Of the four pathways of domestication process, camelds are models for the “prey pathway,” which includes medium to large ungulates targeted as prey (Larson and Fuller, 2014). In the transition from game management to herd management, hunter-gatherers changed their hunting strategies to maximize the availability of the prey (Larson and Fuller, 2014). In the Southern Andes, this is suggested by several proxy data. The representation of camelid bone remains in archeofaunal assemblages increased through time from 29.7% to nearly 90%, whereas other taxa, generally small fauna, are markedly reduced (Figure 2).

The composition of zooarcheological assemblages and their temporal change is reflected in their diversity (Shannon $H$) and dominance ($D$) indices. The Shannon $H$ index accounts for the abundance and evenness of species in an assemblage, and $D$ is a measure of dominance that reveals the most conspicuous and abundant species. From the Early to the Mid-Holocene II (12890–3770 cal. BP), diversity decreased from 0.93 to 0.29, whereas $D$ increased from 0.52 to 0.84 (Yacobaccio, 2013; Yacobaccio et al., 2017). These two measures can be used as proxies of generalized vs. specialized use of fauna, in this particular case, camels. The most ancient radiocarbon dates associated to this change in diversity and dominance indices have a pooled mean of 6177 ± 39 cal. BP. This suggests that before that date, hunter-gatherers had a mostly generalized subsistence base on a regional scale in which different habitats were used according to gross species abundance, with the exception of patches that offered a limited range of resources in great quantities, like some rodent colonies or bird nesting places. After ca. 6200 cal. BP, a specialized economic system emerged and resilient habitats were used mainly to increase the use of the focal resource. The emergence of this economic system based on the intensification of the use of camels coincided
Animal Frontiers

published by Barreta et al. (2013) confirmed that guanacos are of the mitochondrial genome in Bolivian llamas and alpacas; these studies concluded that the llama was derived from the guanaco lineage when microsatellites were analyzed (Kadwell et al., 2001). Microsatellites notwithstanding, this study found exclusive haplotypes shared between alpacas and vicuñas, but a significant number of alpacas (51%–63% of the samples) were found to belong in the guanaco clade. This indicates a high degree of hybridization, suggesting that alpacas had a mixed origin, or alternatively, that an introgression occurred during or after domestication. If the first hypothesis were confirmed, it would provide strong proof that alpacas were domesticated after the llama. The mitochondrial control region indicates that all the haplotypes shared between guanacos and alpacas also exist in llamas. This could indicate that hybrids between domestic forms were common. The model derived from the results of Barreta et al.’s article lend support to the idea that alpacas resulted from interbreeding between vicuñas and llamas. Likewise, from the confirmation of the existence of two lineages of guanacos (northern and southern) and the finding that some llamas share haplotypes with southern guanacos, they conclude: “The present study would support also at the genetic level and taking into account the archaeological evidence, the existence of additional llama domestication centres in Argentina and Bolivia” (Barreta et al., 2013: 8). The hybridization produced since the 16th century AD, after the Spanish conquest of the Andes, produced mitochondrial lineages shared between different species, limiting the power of this line of inquiry to clarify the origins of domestic cameldids.

Figure 2. Use of animal resources during the Holocene in the high Andean region of Northwestern Argentina and northern Chile. Box and jitter graph of data represented in percent of identified bones from 28 archeological sites. See location of some of them in Figure 3.

Genetic Evidence

Genetic studies have shown that the two wild species belong to two distinct groups and are therefore good taxonomic genera. Vicuñas are differentiated into two clades; that is, they are two parts of a group that descend from a common ancestor, according to their subspecific assignment (the northern, Vicugna vicugna mensalis and the southern Vicugna vicugna vicugna). Guanacos can also be divided into two subspecies: the northern L. g. caesilensis and the southern L. g. guanicoe, the latter having a greater distribution that includes the southern Andes and Patagonia (Kadwell et al., 2001; Marin et al., 2017). Llamas and guanacos form a monophyletic group in a clear ancestor-derived species process. Alpacas are associated with vicuñas from mitochondrial genome (Marin et al., 2017), and also associated with the guanaco lineage when microsatellites were analyzed (Kadwell et al., 2001). Microsatellites notwithstanding, these studies concluded that the llama was derived from the guanaco and the alpaca from the vicuña (Wheeler et al., 2006; Marin et al., 2017).

An analysis of genetic diversity in the hypervariable region of the mitochondrial genome in Bolivian llamas and alpacas published by Barreta et al. (2013) confirmed that guanacos are the ancestors of llamas, but the origin of alpacas remains unclear (Barreta et al., 2013). The article found exclusive haplotypes shared between alpacas and vicuñas, but a significant number of alpacas (51%–63% of the samples) were found to belong in the guanaco clade. This indicates a high degree of hybridization, suggesting that alpacas had a mixed origin, or alternatively, that an introgression occurred during or after domestication. If the first hypothesis were confirmed, it would provide strong proof that alpacas were domesticated after the llama. The mitochondrial control region indicates that all the haplotypes shared between guanacos and alpacas also exist in llamas. This could indicate that hybrids between domestic forms were common. The model derived from the results of Barreta et al.’s article lend support to the idea that alpacas resulted from interbreeding between vicuñas and llamas. Likewise, from the confirmation of the existence of two lineages of guanacos (northern and southern) and the finding that some llamas share haplotypes with southern guanacos, they conclude: “The present study would support also at the genetic level and taking into account the archaeological evidence, the existence of additional llama domestication centres in Argentina and Bolivia” (Barreta et al., 2013: 8). The hybridization produced since the 16th century AD, after the Spanish conquest of the Andes, produced mitochondrial lineages shared between different species, limiting the power of this line of inquiry to clarify the origins of domestic cameldids.

For this reason, it is of fundamental importance to carry out paleogenomic studies. Of the various studies carried out in South American cameldids (Weinstock et al., 2009; Westbury et al., 2016; Diaz Marotto, 2018; Abbonna et al., 2020), we will comment here on Diaz Marotto (2018) for its relevance. The author analyzed samples from three archeological sites in the Salar de Atacama area (Chile) dated between 2750 and 2500 cal. BP. This work does not refer to the earliest moments of domestication, but rather to the moment when pastoralism became the predominant economic strategy among local human groups. Diaz Marotto studied the complete genome of 77 bone samples, obtaining conclusive evidence that llamas were domesticated derivatives of guanacos. In this case, a distinction between subspecies of guanacos was made, confirming that both L. g. guanicoe and L. g. caesilensis were the ancestors, in contrast with previous studies of current genomes (Marin et al., 2007) that had proposed L. g. caesilensis as the only ancestor. In turn, the complete mitochondrial genome of alpacas shows that this species, also domesticated before the Spanish conquest, has a much closer relationship with the guanaco/llama lineage than with vicuñas. Diaz Marotto determined a clade of domestic species where llamas and alpacas are grouped very closely, suggesting that the llama was the first domesticated species, followed by the emergence of alpacas as a result of sustained interbreeding of female llamas with male vicuñas. Likewise, she agreed with Barreta et al. (2013) that the South-Central Andes was a domestication center independent from that of the Central Andes, based on the evidence of the domestication of the L. g. guanicoe subspecies. This conclusion is in line with more general arguments about the number of...
domestication events in ungulates. The use of genetic sequences has led numerous authors to conclude that animal domestication was a great deal more frequent and evenly distributed than previously thought. This claim is based “on the affinity between DNA sequences of domestic animals and their wild counterparts and the assumption that branching patterns on phylogenetic trees reflect independent domestication episodes. This rationale has been used to support claims for multiple and independent domestications of genetically and geographically divergent populations” (Larson and Fuller, 2014: 214). This argument takes into consideration pigs, goats, sheep, horses, cows, and now, we may add, llamas.

**Main Archeological Evidence**

Osteometry has been used as a proxy to study the domestication process by detecting changes in the size of individuals. Indeed, South American camelids have a size gradient from smallest to largest: vicuña—alpaca—northern guanaco—llama. This defines two groups: the small one—vicuñas and alpacas—and the large one—guanacos and llamas. As can be seen, both groups contain wild and domestic camels. In the large group, there is a complicated fact, and it is the variability of the guanaco size; *L. guanicoe* has a high clinal dispersion, which influences its size. Patagonian or southern guanacos are much larger than northern or North Andean ones. That is why we have to be very careful with the reference measurements used to compare with archeological samples. These issues have been widely debated in Andean zooarcheology (Izeta, 2008; Cartajena, 2009; Gasco et al., 2014; Hernández and L’Heureux, 2019). In the large group, the variation in guanaco size imposes certain restrictions on the determination of species (guanaco vs. llama) based on osteometry alone. There is a “zone of uncertainty” defined by an overlap in size between small llamas and North Andean guanacos. However, the larger measurements beyond this overlap zone can be determined as *L. glama* without a doubt.

As far as we know today, the first osteological evidence of a change in size in camels that could reveal the modifications produced by a domestication process made its first appearance in the Southern Andes at approximately 7100 yr cal. BP. This evidence is an increase in the width of the distal metacarpus, along with an increase in the size and robustness of other bones, such as phalanges, scapulae, and humerus. These specimens are larger than the known sizes for North Andean guanacos, and are comparable to modern llamas, or even larger (Cartajena et al., 2007). This points to the emergence of a camelid similar in size to the largest among current llamas. In the herd protection phase (7100–4500 cal. BP), there was an increase in size variability, especially with the emergence of very large specimens that were first detected in the archiological record at the Hornillos 2 site (layer 2), but later increased their distribution significantly between 5800 and 4200 cal. AP, when their presence was noted in numerous archeological sites in the region (Izeta, 2010, Figure 3).

The variation in size that we refer to here is summarized in Figure 4. From the variation in breadth of the phalanx *I* facies articularis proximalis, it can be inferred that between 4900 and 4700 cal. BP, there was a significant proportion of camelids that were larger than current llamas, followed by a reduction in size by 2600 cal. BP, setting the median and interquartile range equal to that of current llamas. From that date on, bone remains equivalent to the size of current llamas were found in other ecosystems, such as mesothermal valleys (between 1500 and 2900 masl) and the lowlands of the Chaco, in numerous archeological sites (Izeta, 2010, Mercolli, 2019, Del Papa, 2020). At 580 cal. BP, during the Inka period, the size of llamas completely coincides with that of today.

Harbers et al. (2020) found that “mobility reduction induces a plastic response beyond the shape variation of wild boars in their natural habitat, associated with a reduction in the range of locomotor behaviours and muscle loads” and that produces changes in the calcaneus shape due to captivity. This is extremely interesting because identifies osteological modification as a consequence of confinement. The case of llamas seems to have been different because there was no such a reduction in mobility on that scale. Moreover, as llamas were used as pack animals, their mobility remained to be high. However, the zooarcheological record shows a reduction of size variability after pens appeared as can be seen in Figure 4, where size became stabilized since 2600 cal. BP. Recent research on morphometric analysis made on guanaco and llamas first phalanges are promising, but no conclusive yet because the two species have not been differentiated in these analyses (Hernández and L’Hereux, 2019).

Other paleoenvironmental and archeological evidence accompanied the change in size of the camels and point toward a change in the camelid-human relationship. Pollen analysis in several localities of the Puna has identified a greater abundance of Chenopodiaceae–Amaranthaceae and *Pennisetum* from ca. 5300 to 4869 cal. BP, which is consistent with more intensive human impact on wetlands due to environmental management practices, perhaps including periodic burning to increase patch productivity.

This change in the human–camelid relationship is also seen in the appearance of bone pathologies. Cartajena et al. (2007) recorded periostitis in distal phalanges and metaphasia from the Tulán 52 (ca. 4900 cal. BP), and Puripica 1 (ca. 4700 cal. BP) sites. These exostoses are due to the proliferation of the bone, possibly caused by long-term irritation of the periosteum. They also observed osteophytes produced by arthropathies due to the living conditions of the animals, perhaps related to the long periods of exercise that are characteristic of pack animals. Labara Encina and Gallardo (2015) analyzed 14 bones with pathologies from the Topater 1 cemetery (Calama, Chile) dated between 2517 and 1868 cal. BP. Most of the pathologies occur in limb bones that, due to their size, have been assigned to *L. glama*. At this site, most of the phalanges show exostoses, an abnormal formation of new tissue on the outside of the bone. Its manifestation is mild to moderate and is located in the diaphysis and, to a lesser extent, on the dorsal and lateral faces.
of the epiphyses. These pathologies were attributed to constant trauma to the joints and could result from environmental factors, such as walking on uneven ground, as well as the excessive use of the animal due to cultural practices, as would be the case of pack animals. A llama head with articulated vertebrae recovered from an inhumation at the Huachichocana III site (3170 and 2867 cal. BP, Figure 3) also provided information on bone pathology. The cervical vertebra C2 (axis) presents periostitis due to direct trauma caused by a tie rope or due to an excessive use of this articular section of the neck. The use of a muzzle to restrain the animal is the most probable explanation for this pathology. In addition, the analysis of stable carbon and nitrogen isotopes reinforces the idea that this specimen's diet was strongly determined by human intervention (Yacobaccio et al., 2018).

The use of muzzles at such an early time would not be uncommon, as ropes had already begun to appear in the archaeological record. These ropes can be interpreted as a technological innovation related to the onset of pastoralism. Indeed, at Alero Unquillar (Figure 3), in which two metapodia were determined to belong to a llama by osteometry, a rope made with local grasses dated at 3989–3570 cal. BP was recovered. Likewise, the grave goods of human burial number 4 in the Huachichocana III site included remains of ropes made with palm leaf fibers and local bromeliads (*Tillandsia usneoides* or *Deuterocohnia*; Lema, 2017). This burial is relatively contemporary with burial 3—the one containing the llama head—and it could be associated with two radiocarbon dates from ca. 3360 to 3170 cal. BP. The appearance of ropes has direct implications for the development of gripping or restraining technology associated with herd management.

A change in phase of the domestication process occurred with the emergence of confinement technology, that is, pens. The first corrals were small caves, such as Inca Cave 7, whose entrance was covered by a wall of stone boulders. Given its size, it was probably used to keep young camelids in confinement. A layer of guano that covered the ground within the cave was helpful to determine its function as a corral (Figure 5).
As mentioned above, two metapodia determined as belonging to very large llamas were recovered from this site. The corralling episode has been dated between 4635 and 4232 cal. BP. In the Tulán gorge, Cartajena et al. (2007: 168) observed that the “high and rocky slopes are used as natural boundaries on both borders; on softer slopes big regular boulders have been arranged to enclose pens. Boundary lines that cross the ravine are interrupted in the stream part to avoid holding back the water; palisades or similar solutions were probably used to bound areas close to the stream, allowing the enclosure of pens[…].] Pen dimensions (ca. 300 × 80 m) suggest a considerable amount of labour.” This corral can be dated at around 2600 cal. BP.

Discussion and Conclusion

The transition from hunting to herding has been a complex one. Herd protection involved changes in camelid behavior, but also modifications in the human strategies used to approach wild camelids. This phase lasted for a long time, from 7100 to 4500 cal. BP, when corrals made their first appearance in the region.

Environmental fragmentation promoted the aggregation of humans and wildlife in resilient habitats, such as wetlands, thus creating the conditions for the development of a closer and more stable relationship between people and camelids. Then, as a condition for herd protecting and habituation, human communities reduced their mobility, stabilizing their residence in these areas where grazing resources were more concentrated. The entire time span characterized by herd protection also was accompanied by an increase in human population. The archaeological sites show evidence of more intensive occupation at this time, and about 4900 cal. BP, the first site with several stone enclosures could have functioned as places where human populations gathered periodically (Núñez and Perlès, 2018).

Moreover, the grouping of radiocarbon dates from archaeological sites in the region has been analyzed as indicators of human demography or anthropogenic signal in the Holocene (Muscio and López, 2016). This study suggests that after 6177 cal. BP, the anthropogenic signal increased, reaching its maximum at 4700 cal. BP. This date correlates quite well with the
appearance of corrals or courtyards in the archeological record, which points toward an intensification of the camelid domestication process. This is a significant finding because, for the diffusion of an innovation to take place—in this case, domesticated animals—there need to be long-reaching interaction networks sustained by large, interconnected populations. Evidence of such interaction networks can be provided by the existence of domestic llamas in the microthermal valleys and lowlands of the Chaco region since at least 2100 cal. BP, which reveals a relatively rapid dispersion of this innovation.

This process is contemporary with what happened in the Central Andes, in which early domestication indicators in the Puna de Junín occurred between 5470 and 3480 cal. BP (Moore, 2016). In southern Peru, in the Osmore Valley, evidence of domestication appeared between 4090 and 3677 cal. BP (Aldenderfer, 1998).

Taking the Andes as a whole, the knowledge that we have about the domestication of camelids is still very fragmentary, since there are portions of territory with no archeological evidence on the period that concerns us. Likewise, more genetic studies on ancient materials are needed, although according to what we know today, llamas would have been domesticated first and later, alpacas, which resulted from introgressions of llamas with vicuñas (Barreta et al., 2013; Díaz Marotto, 2018).

There are still many unresolved issues and unanswered questions concerning the domestication of South American camelids. The next few years will hold many new and exciting insights for researchers in this challenging field.

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**Literature Cited**

Abbonna, C.C., G. Neme, J. Jonhson, T. Kim, A.F. Gil, and S. Wolverton. 2020. Were domestic camelds present on the prehispanic South American agricultural frontier? An ancient DNA study. PeerJ 15:e0240474. doi:10.7317/journal.pone.0240474

Aldenderfer, M.S. 1998. Montane foragers. Asana and the south-Central Andean Archaic. Iowa City (IA): University of Iowa Press.

Aschero, C.A., and J. Martinez. 2001. Técnicas de caza en Antofagasta de la Sierra, puna meridional Argentina. Relac. Soc. Argent. Antropol. 26:215–241.

Barreta, J., B. Gutierrez-Gil, V. Iniguez, V. Saavedra, R. Chiri, E. Latorre, and J.J. Arranz. 2013. Analysis of mitochondrial DNA in Bolivian llama, alpaca and vicuna populations: a contribution to the phylogeny of the South American camelids. Anim. Genet. 44:158–168. doi:10.1111/j.1365-2052.2012.02376.x

Cartajena, I. 2009. Explorando la variabilidad morfométrica del conjunto de camélidos pequeños durante el Arcaico Tardío y el Formativo Temprano en Quebrada Tulán, norte de Chile. Rev. Museo Antropol. 2:199–212.

Cartajena, I., L. Núñez, and M. Grosjean. 2007. Camelid domestication on the western slope of the Puna de Atacama, northern Chile. Anthropozoologica 42:155–173.

Crispo, E. 2007. The Baldwin effect and genetic assimilation: revisiting two mechanisms of evolutionary change mediated by phenotypic plasticity. Evolution 61:2460–2479. doi:10.1111/j.1558-5646.2007.00203.x

Darwin, C.R. 1868. The variation of animals and plants under domestication. 2 vols. London (UK): J. Murray.

Del Papa, L.M. 2020. Diferenciación de camélidos en la región Chaco-Santiaguëla (Argentina) para el periodo Agroalfarero (350–1550 dC). Latin American Antiquity. https://www.cambridge.org/core/terms. doi:10.1017/laq.2019.104

Díaz Marotto, P. 2018. ADN antiguo y los camélidos sudamericanos: Un aporte metodológico para la identificación taxonómica de restos óseos de camélidos durante la Fase Tilocalar (3.200–2.300 A.P.) en los sitios Tulán-54 y Tulán-85, Norte de Chile. Chile: Tesis para optar al grado de Doctora, Universidad de Tarapacá/Universidad Católica del Norte.

Francis, R.C. 2015. Domesticated: evolution in a man-made world. New York (NY)/London (UK): Norton & Company Inc.

Gasco, A., E. Marsh, and J. Kent. 2014. Clarificando variables osteométricas para la primera falange de camélidos sudamericanos. Intersecc. Antropol. 15:131–138.

Harbers, H., D. Neaux, K. Ortiz, B. Blanc, F. Laurens, I. Baly, F. Laurens, I. Baly, F. Laurens, I. Baly, and N. Tripcevich, editors. The archaeology of Andean pastoralism. 2 vols. London (UK): J. Murray.

Izeta, A. 2008. Late Holocene camelid use tendencies in two different ecological zones of Northwestern Argentina. Q Int. 180:135–144.

Izeta, A. 2010. Variabilidad osteométrica de sitios arqueológicos del NOA. In: Gutiérrez, M.A., M. De Nigirs, P.M. Fernández, M. Giardina, A. Gil, A. Izeta, G. Neme and H. Yacobaccio, editors. Zooarqueología a principios del siglo XXI. Buenos Aires (Argentina): Ediciones del Espíritu; p. 29–38.

Kadwell, M., M. Fernandez, H.F. Stanley, J.C. Wheeler, R. Rosadio, and M. Bruford. 2001. Genetic analysis reveals the wild ancestors of the llama and alpaca. Proc. R. Soc. Lond. B 268:2575–2584.

Labarca Encina, R., and F. Gallardo. 2015. The domestic camels (Cetartiodactyla: Camelidae) from the middle formative cemetery of
Topater 1 (Atacama Desert, Northern Chile): osteometric and palaeo-pathological evidence of cargo animals. Int. J. Osteoarchaeol. 25:61–73.
Larson, G., and D.Q. Fuller. 2014. The evolution of animal domestication. Annu. Rev. Ecol. Evol. Syst. 66:115–36.
Lema, V. 2017. Del Objeto al Contexto: La(s) capa(s) “E” de Huachichoca III. Chungara 49:209–226.
Marín, J.C., C.S. Casey, M. Kadwell, K. Yaya, D. Hoces, J. Olazabal, R. Rosadio, J. Rodriguez, A. Spotorno, M.W. Bruford, et al. 2007. Mitochondrial phylogeography and demographic history of the vicuña: implications for conservation. Heredity (Edinb) 99:70–80. doi:10.1038/sj.hdy.6800966
Marín, J.C., K. Romero, R. Rivera, W.E. Johnson, and B.A. González. 2017. Y-chromosome and mtDNA variation confirms independent domestica-
tions and directional hybridization in South American camelids. Anim. Genet. 48:591–595. doi:10.1111/age.12570
Mercollí, P.H. 2019. La importancia de los recursos silvestres y el manejo de los rebaños de llamas por parte de las sociedades humanas en la Quebrada de Humahuaca, Jujuy, Argentina (0-1536 d.C.). Tesis para optar por el grado de Doctor. Facultad de Filosofía y Letras, Universidad de Buenos Aires.
Moore, K. 2016. Early Domesticated Camelids in the Andes. In: Capriles, J.M. and N. Tripevich, editors. The archaeology of Andean pastoralism. Albuquerque: The University of New Mexico Press; p. 17–38.
Muscio, H.J., and López, G.E.J. 2016. Radiocarbon dates and anthropogenic signal in the South-Central Andes (12,500–600 cal. years BP). J. Archaeol. Sci. 65:93–102.
Núñez, L., and C. Perlés. 2018. Ceremonialismo y congregación durante la transición arcaico tardío y formativo temprano en la circumpuna de Atacama (norte de Chile). Rev. Geogr. Norte Gd. 70:183–209.
Price, E.O. 1984. Behavioural aspects of animal domestication. Q. Rev. Biol. 59:1–32.
Price, E.O. 2002. Animal domestication and behavior. Wallingford (UK): CABI Publishing.
Vilá, B. 2012. Camélidos Sudamericanos. Buenos Aires (Argentina): EUDEBA.
Weinstock, J., B. Shapiro, A. Prieto, J.C. Marin, B.A. González, M. Thomas, P. Gilbert, and E. Willerslev. 2009. The Late Pleistocene distribution of vicuñas (Vicugna vicugna) and the “extinction” of the gracile llama (“Lama gracilis”): new molecular data. Q. Sci. Rev. 28:1369–1373.
Westbury, M., S. Prost, A. Seelenfreund, J.M. Ramirez, E.A. Matisoo-Smith, and M. Knapp. 2016. First complete mitochondrial genome data from an-
cient South American camels – the mystery of the chilihuesques from Isla Mocha (Chile). Sci. Rep. 6:38708. doi:10.1038/srep38708
Wheeler, J.C., L. Chikhi, and M.W. Bruford. 2006. Genetic analysis of the ori-
gins of domestic South American camels. In: Zeder, M.A., D.G. Bradley, E. Emswiller, and B.D. Smith, editors. Documenting domestication. New genetic and archaeological paradigms. Berkeley (CA): University of California Press; p. 329–341.
Yacobaccio, H.D. 2013. Towards a human ecology for the middle Holocene in the southern Puna. Q. Int. 307:24–30.
Yacobaccio, H.D., M.R. Morales, and R. Hoguin. 2017. Habitats of ancient hunter-gatherers in the Puna: resilience and discontinuities during the Holocene. J. Athropol. Archaeol. 46:92–100.
Yacobaccio, H.D., C.T. Samec, G. Marcoppido, and B. Vilá. 2018. Domesticación de camélidos y patologías óseas: un caso de estudio en la puna Argentina. Cuadernos Inst. Nac. Antropol. Pensam. Latinoam. Ser. Especiales 6:71–79.
Yacobaccio, H.D., and B. Vilá. 2016. A model for llama (Lama glama Linnaeus, 1758) domestication in the southern Andes. Anthropozooologica 51:5–13.