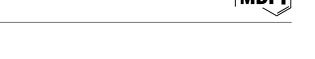


Review



A Review on the Latest Early Pleistocene Carnivoran Guild from the Vallparadís Section (NE Iberia)

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Abstract: The Vallparadís Section encompasses various geological layers that span a significant chronological range, extending from the latest Early Pleistocene to the early Middle Pleistocene, covering a timeframe from approximately 1.2 to 0.6 Ma. This period holds particular importance, as it coincides with a significant climatic transition known as the *Early–Middle Pleistocene Transition*, a pivotal phase in Quaternary climatic history. This transition, marked by the shift from a 41,000-year obliquity-driven climatic cycle to a 100,000-year precession-forced cyclicity, had profound effects on the Calabrian carnivorous mammal communities. Notably, the once diverse carnivore guild began to decline across Europe during this period, with their last documented occurrences coinciding with those found within the Vallparadís Section (e.g., *Megantereon* or *Xenocyon*). Concurrently, this period witnessed the initial dispersals of African carnivorans into the European landscape (e.g., steppe lions), marking a significant shift in the composition and dynamics of the region's carnivorous fauna.

Keywords: Early Pleistocene; carnivoran guild; Vallparadís Section; Epivillafranchian; Iberia

1. Introduction

1.1. Paleoenviromental Background

Throughout the latest Pliocene and the Early Pleistocene, the taphocenosis of the Northern Palearctic underwent profound transformations in response to the discernible cooling trend and heightened seasonality, a pattern intensified around 3 Ma. Moreover, there is widespread acknowledgment of a shift towards aridification across Europe during the Pleistocene, marking a transition from the tropical–subtropical ecosystems characteristic of the Pliocene to the contemporary environments we recognize today [1–3]. These environmental shifts prompted significant adaptations in large mammal assemblages, favoring cursorial species better suited to open habitats, a transition that coincided with the dispersals of the first hominins out of Africa [4].

Within this period, marked by glacial–interglacial dynamics forced by obliquity cycles with a periodicity of approximately 41 ka, pivotal events shaped the global climatic structure as we understand it. Notable among these were the intensification of cold and warm phases around 1.8 Ma, reflecting the dynamic of Pleistocene climate fluctuations [2,5]. Furthermore, a non-linear phase of the glacial–interglacial cycles, characterized by significant amplitude and asymmetry, began approximately 1.4–1.2 Ma: *the Early–Middle Pleistocene Transition* [6–8].



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Copyright: © 2024 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). The *Early–Middle Pleistocene Transition* (EMPT) refers to a critical period in Earth's history during the Pleistocene epoch, roughly spanning from around 1.2 to 0.7 Ma. This transition marks a significant shift in the dynamics of climatic and environmental conditions, as well as in the evolutionary trajectories of various species. Understanding the *EMPT* is crucial for unraveling the complex interactions between climate, ecology, and human evolution during this pivotal period in Earth's history. Specifically, in the North Mediterranean, the *EMPT* induced an increase in aridity and seasonality which could be related to the extinction of several groups of large mammals and the dispersal of new forms from Asia and Africa [4,6,8]

The end of the Early Pleistocene was characterized by even more drastic events, including a substantial glacial phase around 0.9 million years ago, known as the '900 ka event' [7,9,10], and a progressive transition towards forcing by precession cycles, with a periodicity of approximately 100 ka [11]. These transformative processes are intricately linked to shifts in global temperature and seasonality, often inferred through biological proxies within animal populations and communities [12–14].

1.2. The Early–Middle Pleistocene Transition Carnivoran Guild

Early Calabrian times were characterized mostly by open habitats [2], with evidence of aridity periods, exemplified on the site of Venta Micena by the presence of *Hemitragus* and *Praeovibos* [15]. Since the beginning of the Late Villafranchian (ca. 2.0 Ma), the carnivoran guild was dominated mostly by cursorial species adapted to open environments: the sabertoothed cats *Homotherium crenatidens* and *Megantereon adroveri*, the large pantherine *Panthera gombaszoegensis*, the feline *Viretailurus pardoides*, and the Mediterranean lynx *Lynx pardinus*. *Pachycrocuta brevirostris* emerged as the dominant large hyaena within these ecosystems, alongside canids such as *Xenocyon lycaonoides*, *Canis mosbachensis*, and *Vulpes alopecoides*. Additionally, a derived omnivorous form of large ursid, *Ursus etruscus*, was extensively recorded in Mediterranean Europe. Unfortunately, despite this rich carnivore diversity, the record of mustelids is relatively scarce; only badgers, represented by the extant species *Meles meles*, are abundantly recorded in Europe.

This situation started to change around 1.2 Ma with the record of more humid and forested habitats, precisely at the Vallparadís Section [16]. This trend has a break around the Jaramillo magnetostratigraphic subchron (MIS30), with another pulse of aridity recorded at the layer EVT12 of the Vallparadís Section but also in the Iberian site of Quibas [17,18], with new evidences of *Hemitragus* and *Praeovibos*. However, the increase in humidity and seasonality and the general harshening of the environmental conditions were perfectly recorded since the 0.9 Ma event. All the former changes had a significant impact on the carnivoran guild. The high diversity and intraspecific competition of Villafranchian species [19] led to the vanishing of the genus *Megantereon* around 1 Ma, followed later around MIS21 by *Xenocyon* and *Viretailurus* and the progressive decrease on the abundance of *Homotherium* and *P. gombaszoegensis*. These species were replaced by the African newcomers, *Panthera spelaea fossilis, Panthera pardus,* and *Crocuta crocuta,* who started being recorded as being present in several sites since 1.2 Ma, with the first leopard record from Vallonnet Cave [20].

1.3. The Vallparadís Section

The Vallparadís Section encompasses the paleontological sites of Cal Guardiola (CGR) and Vallparadís Estació (EVT), situated in the Vallès-Penedès Basin in Northeastern Iberia [21,22]. Over the course of emergency excavations conducted from 1997 to 2008, these sites yielded 32,000 vertebrate remains dating from the late Early to Middle Pleistocene. Both locations exhibit a distinctive depositional setting strongly influenced by the dynamics of an alluvial fan system and the geometry of the Miocene paleorelief.

Biochronological, magnetostratigraphic, and U-series–ESR evidence collectively suggests that the Vallparadís Section ranges from before the Jaramillo paleomagnetic subchron (approximately 1.2–1.1 Ma) to the early Middle Pleistocene (around 0.6 Ma) [21–24].

Analyzing the older part of the section (circa 1.2–1.1 Ma), pollen and wood studies from CGRD2 indicate a warm–temperate and humid ancient environment, likely a river or river–marsh ecosystem that supported diverse plant life, including aquatic macrophytes, deciduous trees, and grasses. The abundance of hippopotamuses that remains in this layer supports the interpretation of a fluvial primary depositional environment [25], while the high diversity of large ungulates suggests a landscape with both woodlands and more open, arid areas [16].

Meso- and microwear analyses conducted on a substantial sample of ungulate teeth from the section indicate a noteworthy shift in paleoenvironments since 0.9 Ma (MIS22). Initially dominated by open, dry grasslands with discernible seasonality (Layer EVT12, circa 1.0 Ma; MIS30), the environment transitioned to more humid woodlands, possibly with an even more pronounced seasonality (Layers EVT7 and CGRD7; circa 0.86 Ma; MIS21). These findings align with data from other contemporaneous Southern European sites [10]. Preliminary investigations into the stable isotopic signal from layers EVT12 and EVT7 also suggest a period of increased aridity during the Jaramillo subchron and more wooded environments during it. The study of enamel hypoplasia in hippopotamuses points in the same direction, highlighting increased seasonality in Europe during the Early Pleistocene [26].

2. Materials and Methods

The specimens reviewed in the present paper are housed at the Catalan Institute of Paleontology, Sabadell, Spain (ICP), and came from the Cal Guardiola and Vallparadís Estació emergency excavations. The code of the ICP collections is IPS. The detailed record of carnivoran species in each Vallparadís Section layer is displayed in Table 1.

Table 1. Occurrences of the different carnivoran species in each geological layer of the Vallparadís Section (see Supplementary Figure S1 for the stratigraphic scheme of the Vallparadís Section).

	Geological Layers of the Vallparadís Section					
_	Cal Guardiola	Local Section	Vallparadís Estació Local Section			
_	CGRD2	CGRD7	EVT12	EVT10	EVT7	EVT3
Homotherium crenatidens						
Panthera fossilis						
<i>Megantereon</i> sp.						
Panthera gombaszoegensis						
Puma pardoides						
Lynx pardinus						
Pachycrocuta brevirostris						
Canis (Xenocyon) lycaonoides						
Canis mosbachensis						
Vulpes alopecoides						
Vulpes vulpes						
Ursus deningeri						
Meles meles						

All specimens discussed in the text have been personally studied by the authors in the last years and compared with late Early Pleistocene assemblages they also personally studied, including Dmanisi (Georgia), Upper Valdarno, Olivola, Pirro Nord, Collecurti and Cava Redicicoli (Italy), Venta Micena, Barranco León, Fuente Nueva 3, Incarcal Complex, La Boella and Bòvila Ordis (Spain), and Le Vallonnet (France).

Bootstrapping Cluster Analysis and Non-Metric Multidimensional Scaling

To evaluate the similarity of the carnivore guild in Vallparadís with that from other Early Pleistocene localities in Eurasia and Africa, we conducted a bootstrapping cluster analysis (BCA) on specific composition matrices of the selected assemblages (Supplementary Table S1). The twenty-three localities are fairly evenly distributed geographically (six in Africa, eight in Asia, and nine in Europe) and date between approximately 2.0 and 0.7 million years ago (see Supplementary Figure S2). Similar to previous studies using comparable analyses [27,28], we selected localities with well-established chronologies and a substantial number of recorded carnivorans to avoid biased results. The occurrence matrices compiled are based on published data ([9,21,23,28–44]), with some cases revised and updated.

BCA is a segmentation technique that is valuable for evaluating the stability of clustering outcomes, also utilized to detect notable statistical resemblances among Pleistocene large mammal communities (refer to [27,28,45,46]). The process commences with an initial grouping of the dataset via the unweighted pair group method with arithmetic mean (UPGMA) algorithm [47]. This initial grouping acts as a baseline clustering of the chosen localities based on their taxonomic similarity, enabling us to gauge the likelihood of each branch in the baseline using a permutation approach. Following this, a random subset of the original data is selected to perform a new UPGMA cluster analysis, leading to a new clustering arrangement. This procedure is repeated numerous times. The newly generated clustering is then compared to the baseline one. A similarity index (G*) is computed between the baseline and the sample clustering, with values ranging from 0 (if the clusterings are entirely dissimilar) to 1 (if the clusters from the original and sampled data are identical). The G^{*} is subsequently contrasted with the expected similarity value (G°) under the null hypothesis that the sampled data are a true random subset of the original dataset. We performed the resampling procedure 1000 times, and in each instance, we conducted UPGMAs on the sampled data and determined similarity indices. If the likelihood that G^* is greater than or equal to G° surpasses the significance threshold (p($G^{\circ} \leq G^{*}$); $\alpha = 0.05$), the partitioning levels of the baseline cluster analysis are deemed robust (see [48]). We also explored ecological subdivisions within the analyzed guild, classifying carnivorans based on traditional dietary categories: hypocarnivores (diet containing less than 50% vertebrate meat), mesocarnivores (diet containing 50%–70% vertebrate meat), and hypercarnivores (diet containing more than 70% vertebrate meat), following [49,50]. Furthermore, we included an insectivore category (species specialized in consuming insects) and incorporated hunting strategies and dietary habits based on ecological categories proposed in the literature, e.g., [28,35,50–55].

Finally, we categorized the fossil carnivorans according to their habitat preference, particularly as taxa of open, mixed, closed, or aquatic environments. The same dataset was used to perform a non-metric multidimensional scaling (NMDS), an ordination method suitable for taxa occurrence matrices, as it uses distance matrices of the presence/absence of the taxa (but also relative abundances) to project the dataset in bi- or tridimensional spaces [56]. In the case at hand, NMDS on taxa occurrences was performed, taking into consideration the habitat preference per each site. To visualize the proportions of carnivoran abundance in the different sites, we used barplots of their ecological and environmental specifics. We used the software RStudio (v. 2023.12.1+402 'Ocean Storm' Release 4da58325ffcff29d157d9264087d4b1ab27f7204, 28 January 2024 [57]) in R environment (v. 4.3.2, [58]) to perform analyses and produce graphs. The BCA was obtained with the function *pvclust() ('pvclust'* v. 2.2-0 [59]), and for NMDS, the function *metaMDS() ('vegan'* package v. 2.6-4 [60]). The barplots were obtained with *barplot()*

('*graphics*' package v.4.3.2 [58]), and the plot of the NMDS was obtained with *ggplot()* ('*ggplot2*' package v.3.4.0; [61]).

3. Results

3.1. Felids

3.1.1. Homotherium crenatidens

Homotherium was a high-speed pursuit and slender predator adapted to open environments recorded in Eurasia, Africa, and America during the Plio-Pleistocene, and it became extinct around 20 ka [29,35,62–64]. In Europe, it is documented as having been present since the beginning of the Villafranchian period, around 3.0 Ma, with findings in locations such as Perrier-Les Etouaires, Saint Vallier, Senèze, Upper Valdarno, and the Incarcal complex [31,65–67].

Comparisons between *Homotherium*, other sabertoothed cats, and lions in terms of social behavior have been one of the most debated topics among Quaternary paleontologists in recent decades. While much attention has been given to the purported social behavior of *Smilodon*, particularly regarding its packs hunting similar to extant lions [68], the social dynamics of *Homotherium* remain less understood. A recent fossil DNA analysis suggests the social behavior of *Homotherium*; however, the extent of this behavior remains uncertain [63,64]. It is noteworthy that *Panthera spelaea* and *Homotherium* co-existed during more than half million years in Europe; however, the social behavior of both species was probably not similar to that of extant lionesses, according to available data [63,64]

Homotherium was only incidentally recorded at the Vallparadís Section, with a partial upper Canine from the layer CGRD7 [69], whereas it is very abundant, close to 500 specimens, in the Incarcal site of coeval chronology and only 100 km north [67].

3.1.2. Panthera spelaea fossilis

The decline in *Homotherium* records since the Early-to-Middle Pleistocene boundary is partially explained by the arrival of lions from Africa around the Jaramillo subchron, despite the fact that the sociality of both species was probably not very high the first European lions were larger and heavier built than the slender *Homotherium*. Three migration waves indicate the lion's arrival in Europe. The migration from Asia via Eastern Europe into Central Europe is documented by the remains from Bachatsk (western Siberia), dated at ca. 1.0 Ma [70], and later by the find from Kozi Grzbiet (750–700 ka, Poland [71]). The Southwestern Asia and Southeastern Europe route, via the Balkan Peninsula, is recorded at the Greek site Megalopolis–Marathousa [72,73] and Moldovan locality Sinjakovo 1 [74,75]. The earliest European appearance is documented by the Vallparadís Estació record and well supported by the oldest (750–700 ka) so-far known record from Pakefield [76]. Since then, lions were ubiquitously recorded during the Pleistocene of Europe and North America. P. s. fossilis remains were found in the Vallparadís Section in layers EVT10 and EVT7 (MIS30, MIS21; Figure 1W-P', corresponding to at least two different individuals. The most complete one includes a radius, and a practically complete forepaw came from layer EVT10. Additionally, a femur, a tibia, and a third metatarsal of slender appearance were recovered from layer EVT7. These remains represent the oldest records of lions outside Africa and exemplify the dispersal of African carnivorans into Europe during the Early-Middle Pleistocene Transition. The Vallparadís record of *P. s. fossilis* fits well into the Mediterranean migration route of many large carnivores of African origin, like P. pardus from Le Vallonnet Cave (1.2 Ma; [20]) or C. crocuta from Trinchera Dolina 4–6 (850–800 ka) [77]. The arrival in Europe of these three large carnivores of African origin was an important event for the faunal assemblages.

This was a main break in the hitherto stable carnivore paleoguild and led to some changes in its structure. Most of some far dominant species, like *H. crenatidens* and *P. gombaszogensis*, survived but were removed from the apex predator position in carnivore paleoguild. Their previously compact and wide geographic range shrunk into isolated and restricted areas. *H. crenatidens* had noticeably decreased in size and especially in massiveness [62,78–80]. Those lesser, gracile homotheres appeared at the same time in

different Eurasian areas. Lion pressure might play a significant role alongside the combination of climatic-induced changes in vegetation, exacerbated competition among large carnivores, and the increased pressure from *Homo* as a member of the predatory guild [81].

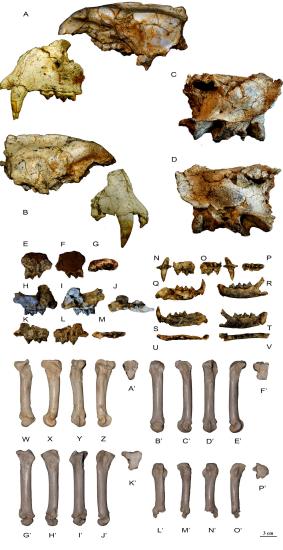


Figure 1. Felids recorded at the Vallparadís Section. Panthera gombaszoegensis cranium (EVT21072) from layer EVT7 in (A) left and (B) right lateral view; Neurocranium of Megantereon sp. (IPS125140) from layer EVT12 in (C) left and (D) right lateral views; maxillary with P3-P4 of Lynx pardinus from layer EVT12 (IPS84917) in (E) buccal, (F) lingual, and (G) occlusal views; maxillary with P3-P4 of Lynx pardinus from layer EVT12 (IPS84920) in (H) buccal, (I) lingual, and (J) occlusal views; maxillary with P3-P4 of Lynx pardinus from layer EVT12 (IPS84916) in (K) buccal, (L) lingual, and (M) occlusal views; maxillary fragment with C1-P4 of Lynx pardinus from layer EVT7 (IPS84915) in (N) buccal, (O) lingual, and (P) occlusal views; right hemi-mandible with c1-m1 of Lynx pardinus from EVT7 (IPS84914) in (Q) buccal, (S) lingual, and (U) occlusal views; left hemi-mandible with p3-m1 of Lynx pardinus from EVT12 (IPS60461) in (R) buccal, (T) lingual, and (V) occlusal views; right second metacarpal of Panthera spelaea fossilis from layer EVT10 (EVT14276) in (W) anterior, (X) lateral, (Y) posterior, (Z) medial, and (A') proximal views; right third metacarpal of Panthera spelaea fossilis from layer EVT10 (EVT16012) in (B') anterior, (C') lateral, (D') posterior, (E') medial, and (F') proximal views; right fourth metacarpal of Panthera spelaea fossilis from layer EVT10 (EVT14720) in (G') anterior, $(\mathbf{H'})$ lateral, $(\mathbf{I'})$ posterior, $(\mathbf{J'})$ medial, and $(\mathbf{K'})$ proximal views; right fifth metacarpal of *Panthera* spelaea fossilis from layer EVT10 (EVT16013) in (L') anterior, (M') lateral, (N') posterior, (O') medial, and (P') proximal views.

The early European lions were notable for their enormous size and robust build. Kurtén [82] observed that these lions made their entry into Europe as giant forms, stating the following: 'The lion entered the European scene (...) with a gigantic form. (...) The Cromerian lion in Europe may be the largest felid that ever existed'. Subsequent studies revealed a general trend of decreasing body size in the *P. spelaea* lineage over time ([82–87]). Throughout the Middle Pleistocene, variations in size among *P. spelaea* were minimal, with specimens dating from around 900 to 300 ka being only marginally larger than those from 300 to 100 ka. Significant size reductions did not occur until approximately 50 ka, coinciding with a marked decline in genetic diversity among lion populations [88]. Between 45 and 15 ka, much smaller lions emerged, comparable in size to modern African lions, *P. leo*. This reduction in body size, along with the eventual extinction of the steppe lion, may have been an ecological response to increasingly harsh environmental conditions and a decrease in prey availability.

3.1.3. Megantereon sp.

Megantereon was a stoutly built ambush predator adapted for stalk prey in closed environments, with a body size of a small jaguar (ca. 100 Kg) [89]. This genus was present in Eurasia and Africa during the Plio-Pleistocene. In Europe, as with *Homotherium*, *Megantereon* has been identified in several sites dating back to 3.0 Ma [44]. It is typically characterized as an ambush predator, and recent studies on Asian fossil assemblages suggest reconsidering *Megantereon*'s habitat preference, indicating a potential adaptation to more open environments rather than dense forests [89,90].

Debate has long surrounded the species diversity within the genus *Megantereon*. Some researchers advocate for the existence of only two primary species in the Old World [89], while others argue for a higher level of diversity [91].

In Europe, *Megantereon* fossils have been unearthed at various sites, including Venta Micena, Pirro Nord, Argentario, Apollonia-1, and, more recently, at Untermassfeld and Vallparadís Section EVT12 [21–23,89].

As an efficient hunter, *Megantereon* likely employed its robust forelimbs to overpower prey, utilizing its sharp and elongated canines (ca. 10 cm in length) to bite, suffocate, and exsanguinate its victims.

As with *Homotherium*, this genus is scarcely recorded in the Vallparadís Section. Only a basicranial fragment from the layer EVT12 is here attributed to *Megantereon* (Figure 1C,D) [92]. The fragmentary nature of the specimen does not permit taxonomical considerations. However, together with the remains from the German site of Untermassfeld, this element represents the last occurrence of this genus in Europe at MIS31-30.

3.1.4. Panthera gombaszoegensis

Several researchers have proposed an African origin for P. gombaszoegensis [93,94], suggesting its dispersal into Europe around 2.0 Ma, linked to the presence of a large pantherine form in the Late Pliocene of Laetoli Upper Beds (circa 3.7 million years ago). Other scholars have attributed the first pantherine fossils from Laetoli to P. leo [95] or, more recently, to a new species, P. principalis [80]. Ultimately, [35,96] offered a more parsimonious hypothesis, noting that African pantherine specimens older than 2 Ma are not identifiable at the species level. We concur with the latter authors, supporting the notion that there is no direct connection between the African specimens and the European P. gombaszoegensis s.l. Since around 2.0 Ma, early forms of P. gombaszoegensis (or P. toscana) have been common in the Late Villafranchian faunas of Europe, evidenced by records from Olivola, Upper Valdarno, Tegelen, Pirro Nord, Untermassfeld, and up to the Middle Pleistocene of Chateau [93,94,97]. In fact, the relationship between the early P. toscana and P. gombaszoegensis remains unclear, with several anatomical differences distinguishing the two forms from different time periods. Recently, Chatar et al. [98] studied Belgian specimens from the Middle Pleistocene and identified more similarities with tigers than with jaguars. It is true that *P. gombaszoegensis* was originally described based on mandibular specimens, whose robust morphology can resemble the ones of jaguars. However, in terms of cranial morphology, *P. gombaszoegensis* is more like the Asian lineage of tiger–snow leopard than to American jaguars. The post-cranial morphology of this species is largely unknown. No partial or complete skeletons were recorded up to now; the most complete record comes from the Middle Pleistocene of Château [94], pointing to a stout and large felid of more than 100 kg, with a similar morphology of a basal pantherine like *Panthera pardus*.

In the Vallparadís Section, *P. gombaszoegensis*, also known as 'The Eurasian Pantherine', was found only in the post-Jaramillo layers CGRD7 and EVT7, with few fragmented postcranial remains [69], isolated teeth, and a relatively complete cranium (Figure 1A,B). The preliminary study of the cranium morphological traits indicates more similarities with Middle Pleistocene forms than to *P. toscana* and suggests a putative second dispersal from Asia rather than to a local gradual evolution of characters. However, the virtual absence of remains from the period 1.8–1.2 Ma precludes accurate comparisons.

3.1.5. Viretailurus pardoides

The Eurasian puma-like cats are exceedingly rare in the fossil record: only one cranium and very few postcranial bones were identified. Consequently, the anatomy and ecology of these middle-sized carnivores remain poorly understood, and their taxonomic classification is a subject of considerable debate.

Panthera schaubi was initially described by Viret [65] at the French site of Saint-Vallier, dating back approximately 2.1 Ma. Viret suggested that these materials exhibited dimensions and morphological characteristics typical of a small pantherine. However, in 1964, Hemmer demonstrated that the skull morphology from St. Vallier was distinctly non-pantherine [99]. Instead, it shares numerous features with the American puma, *Puma concolor*. Consequently, Hemmer assigned the St. Vallier puma-like cat remains to a new genus, *Viretailurus*. Following this, Kurtén and Crusafont [100] described carnivore remains from the Iberian site of La Puebla de Valverde, identifying some as *Panthera* cf. *schaubi*, stressing similarities with specimens from the English Red Crags identified by Owen as *Felis pardoides* [101].

The first puma-like cats from the Iberian Peninsula were first recorded at the Puebla de Valverde site, dating to 2.2 Ma [100]. In Cueva Victoria, some remains were also identified belonging to a puma-like cat. Initial descriptions by Pons-Moyà and Moyà-Solà [102] attributed the specimens to the genus *Jansofelis* sp. Nevertheless, more recent publications attributed these remains to *Puma pardoides* [103–105]. The most recent Iberian puma remains were described in the site of El Chaparral [106].

Evidence from Central Europe includes the German site of Untermassfeld (ca. 1.0 Ma) and the Stránská skála site in the Czech Republic (ca. 0.6 Ma). In Bulgaria, the site of Varshets yielded a humerus fragment, dated to the early part of MN17. Additionally, maxillary fragments from the Georgian site of Kvabebi, previously attributed to *Lynx issiodorensis*, were later ascribed to *P. pardoides*. More recently, Werdelin et al. [107] attributed some new specimens from Graunceanu to a puma-like cat.

To summarize, recent researchers agree on including this taxon in the Felinae subfamily instead of Pantherinae felids. As in the case of *P. gombaszoegensis*, the puma affinities of the mandible are clear, whereas the cranial affinities are more difficult to ascertain. Notwithstanding, the taxonomy of this taxon is outside the scope of this paper, and despite the fact that *Puma pardoides* is the most commonly accepted, we prefer to use *Viretailurus pardoides* until the phylogenetic relationships of this taxon can be clarified.

Viretailurus is only recorded in the layer EVT7 of the Vallparadís Section based on a partial corpus with p4 and a fragment of m1 [105]. No further considerations are needed, except to remark that this specimen can be the last citation of the taxon in Europe.

3.1.6. Lynx pardinus

Small-to-medium-sized felids normally included in *Lynx issiodorensis* have been recorded in Europe since the Pliocene in sites like Serrat d'en Vaquer or Cuevas de Alzamora [108,109]. Nevertheless, the taxonomic adscription of these early felines remains controversial because of the scanty record, and we prefer to include it in *Lynx* sp. Since the beginning of the Villafranchian, true lynxes have been recorded at several sites, such as Perrier-Les Etouaires, Saint Vallier, Olivola, or Pantalla [65,110–112].

According to some scholars, *L. issiodorensis* was the ancestor of both *Lynx lynx* and *Lynx pardinus* [108,112–114], with *L. pardinus* first recorded in the Eastern Iberian Peninsula at ca. 1.6 Ma [113]. However, the morphology of *Lynx issiodorensis* and their similarities with the Mediterranean *lynx* suggests a closer relationship and an early split of *L. lynx* from the *L. issiodorensis–L. pardinus* lineage. The virtually absent Pliocene record in Asia precludes us from ascertaining the origin of the Boreal lynx. Recent studies of Early Pleistocene specimens suggested the inclusion of the specimens from Vallonnet, Apollonia, and Pirro Nord in *Lynx pardinus*, attesting the wide expansion of this taxon in Mediterranean Europe in the latest Early Pleistocene. The ecological role of the first Mediterranean Lynx and when they started their dietary specialization on lagomorphs are two of the questions remaining to be answered in the following years [113].

In the Vallparadís Section, the remains of lynx were recovered from the layers CGRD7, EVT7, and EVT12 [114], including a wide variety of morphological forms and size classes. Interestingly, the remains from the Jaramillo layer, with evidence of aridity and harsh climate [17], are considerably larger than those unearthed from the post-Jaramillo layers, reaching the size of a small *Viretailurus*. The observed variability in dental morphology also points to reconsider most of the previous citations of *Lynx issiodorensis* in the latest Early Pleistocene (Figure 1E–V) [114].

3.2. Hyaenids

Pachycrocuta brevirostris

Pachycrocuta brevirostris is a prevalent carnivore in the late Early Pleistocene faunal assemblages of Europe. Nonetheless, the origin of the genus *Pachycrocuta* has been a subject of intense debate in recent decades. Palmqvist et al. [115] advocated for an African origin of this genus, while other researchers, such as Werdelin [116] and Liu et al. [117], have convincingly argued for an Asian origin. These latter researchers highlight that the earliest records in both Asia and Africa are nearly concurrent: approximately 4.0–3.5 Ma [118].

Regardless of its origin, *P. brevirostris* spread into Europe around 2.2 Ma, with its earliest likely record at the French site of Senèze (J.M.-M unpublished data), where it emerged as one of the most prevalent elements in European faunas, frequently the primary agent responsible for bone accumulations. Its widespread presence persisted until the late Early Pleistocene in Europe, up to 0.86 Ma [21,22,118]. The influence of this species on the European Pleistocene taphocenosis and its potential competition with early hominins has been a topic of significant discussion in recent years [22,118].

One of the more debated topics around *Pachycrocuta* was their putative social behavior and strict scavenging behavior, as favored by several authors [115]. The social behavior of extinct hyaenas was recently studied in several works by Vinuesa et al. [119,120], who suggested that the social behavior of the extant spotted hyaena is probably a recently acquired trait, and the most parsimonious scenario is interpreting most of the extinct hyaenas as mostly solitary animals.

The strict scavenger behavior of *Pachycrocuta* has been deduced based on its fossil accumulations and the morphological traits of the only known skeleton from the Middle Pleistocene of Zhoukoudian-1 in China [121]. First, it is difficult to ascertain if these accumulations were the product of strict scavenging behavior or a combination of scavenging and hunting, as in the extant spotted hyena. Secondly, there are no certainties that this skeleton is really from the same individual and not a composite one. Additionally, all

the Middle Pleistocene Chinese *Pachycrocuta* are considerably larger and morphologically derived compared with their European counterparts from the Early Pleistocene.

Pachycrocuta was found in several layers of the Vallparadís Section from MIS35 to 21, including CGRD2, EVT12, EVT10, CGRD7, and EVT7. The most noteworthy specimen is a partial slender skeleton found in layer EVT7 (Figure 2A–B,D–V,X–Z). This skeleton displays several differences compared with the Zoukoudian-1 one. Firstly, the proportional shortening of the distal-limb elements (tibia and radius), which putatively support the scavenging behavior and cursorial locomotor behavior, is not accentuated in EVT7, being more like that of the brown hyaena. Secondly, the mean of body mass estimations for this skeleton is ca. 82 kg, much smaller than 140 kg, which was estimated for the Chinese specimens. Indeed, the EVT7 specimen is smaller than most of the *Crocuta spelaea* skeletons from the European Late Pleistocene and probably similar in size to the earliest European *Crocuta*. The in-depth study of this specimen will provide clues in the near future that we can use to discern the dietary behavior of this hotly debated taxon.

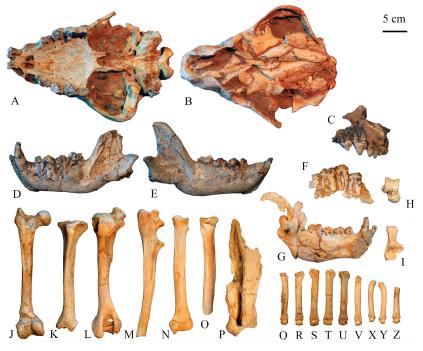


Figure 2. Pachycrocuta brevirostris remains from the Vallparadís Section. Partial cranium (EVT21863) from layer EVT7 in (A) ventral and (B) dorsal views; maxillary fragment with P2-P4 (IPS14521) from layer CGRD2 in (C) buccal view; left hemi-mandible with i3-m1 (EVT24222) from layer EVT12 in (D) buccal view; right hemi-mandible with i3-m1 (EVT24641) from layer EVT12 in (E) buccal view; fragment of right hemi-mandible with p3-m1 (EVT21980) from layer EVT7 in (F) buccal view; right hemi-mandible with i3-m1 (EVT23222) from layer EVT7 in (G) lingual view; right talus (EVT23130) from layer EVT7 in (H) anterior view; right calcaneum (EVT23066) from layer EVT7 in (I) anterior view; left femur (EVT23118) from layer EVT7 in (J) posterior view; right tibia (EVT23224) from layer EVT7 in (K) anterior view; right humerus (EVT23230) from layer EVT7 in (L) posterior view; right ulna (EVT23228) from layer EVT7 in (M) medial view; left radius (EVT23209) from layer EVT7 in (N) anterior view; right partial radius (EVT23213) from layer EVT7 in (O) anterior view; left scapula (EVT23226) from layer EVT7 in (P) dorsal view; right fifth metacarpal (EVT23233) from layer EVT7 in (Q) anterior view; left fourth metatarsal (EVT23144) from layer EVT7 in (R) anterior view; left fourth metacarpal (EVT21709) from layer EVT7 in (S) anterior view; right third metacarpal (EVT23184) from layer EVT7 in (T) anterior view; left third metacarpal (EVT23131) from layer EVT7 in (U) anterior view; left fifth metacarpal (EVT23060) from layer EVT7 in (V) anterior view; left fifth metatarsal (EVT23068) from layer EVT7 in (X) anterior view; right fifth metacarpal (EVT23027) from layer EVT7 in (Y) anterior view; indeterminate metacarpal (EVT23067) from layer EVT7 in (Z) anterior view.

3.3. Canids

3.3.1. Xenocyon lycaonoides

Adaptations for hypercarnivory behavior, characterized by a diet comprising 70% or more vertebrate meat, are relatively common in both fossil and contemporary Canidae lineages [122]. Molecular data suggest that the divergence between the extant species *Lycaon pictus* and *Cuon alpinus* from other wolf-like canids occurred between the Late Pliocene and the Early Pleistocene [123]. The sparse and scattered fossil records of large hypercarnivorous canids across the Old World add complexity to our understanding of the evolutionary history of these wild dogs. These hypercarnivorous dogs are well-documented during the Early Pleistocene in Europe, Asia, and Africa, though different taxa are attributed to them depending on the author [30,124–139].

Martínez-Navarro and Rook [128] proposed a hypothesis highlighting a gradual reduction in certain dental cusps, an increase in others, a tendency towards hypercarnivory behavior, and the loss of the first metacarpal (a unique feature of the modern *Lycaon pictus*). According to their hypothesis, all *Lycaon*-like dogs from the Early Pleistocene should be classified within the genus *Lycaon*, with three chronospecies: *Lycaon falconeri* for the earliest Eurasian forms of the Early Pleistocene, *Lycaon lycaonoides* for those from the latter part of the Early Pleistocene and the beginning of the Middle Pleistocene in Eurasia and Africa, and *Lycaon pictus* for the Middle–Late Pleistocene and contemporary African forms. However, other researchers [131,132,140] have argued that most or all of these forms should be placed in the genus *Xenocyon*, which is considered a sister genus to *Lycaon* and *Cuon*. Given the uncertainty regarding the generic classification of fossil wild-dog specimens and the scope of this paper, we prefer to refer to all large hypercarnivorous taxa with pronounced carnassial teeth and skull adaptations for hypercarnivory from the Late Ploicene and Early Pleistocene of the Old World as belonging to the genus *Xenocyon* until a more definitive hypothesis clarifies their taxonomy.

In the Vallparadís Section, remains attributable to *Xenocyon lycaonoides* were found in layers EVT12 and EVT7, corresponding to a complete foot and two hemi-mandibles, respectively. The large-sized hypercarnivorous canid *Xenocyon lycaonoides* was one of the most characteristic taxa of the second half of the Early Pleistocene ([141], FAD ca. 1.76 Ma [136]) persisting until the Middle Pleistocene age (LAD ca. 450–400 ka [139]).

The hemi-mandibles from layer EVT7 (Figure 3A–F) share several morphologic characteristics with other previously reported remains of *X. lycaonoides* form European assemblages [134].

The studied material exhibits minor differences compared to other specimens of the same species, specifically in the dimensions and sharpness of the main cusps of the m1 talonid and the m2 trigonid, as well as the reduction in the m1 and m2 entoconids. This variation is interpreted as a derived condition relative to the 'typical' *X. lycaonoides* material, indicating a small evolutionary step towards the hypercarnivory observed in this lineage [134]. This finding further supports the gradual nature of hunting-dog evolution. Additionally, the material studied represents one of the latest well chronologically constrained European records of the genus *Xenocyon* at 0.86 Ma, just before the Early–Middle Pleistocene boundary.

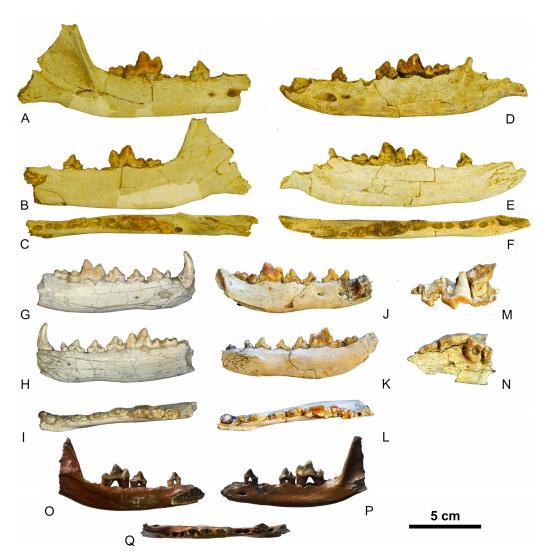


Figure 3. Canids recovered from the Vallparadís Section. Right hemi-mandible with p3, m1–m2 (EVT22049) of *Xenocyon lycaonoides* from layer EVT7 in (**A**) buccal, (**B**) lingual, and (**C**) occlusal view; left hemi-mandible with p2 and p4–m2 (EVT23434) of *Xenocyon lycaonoides* from layer EVT7 in (**D**) buccal, (**E**) lingual, and (**F**) occlusal view; right hemi-mandible with c1–m2 (EVT24342) of *Canis mosbachensis* from layer EVT12 in (**G**) buccal, (**H**) lingual, and (**I**) occlusal view; right hemi-mandible with p1–m2 (EVT13840) of *Canis mosbachensis* from layer EVT7 in (**J**) buccal, (**K**) lingual, and (**L**) occlusal view; left maxillary fragment with P3–M2 (EVT25504) of *Canis mosbachensis* from layer EVT12 in (**M**) buccal and (**N**) occlusal view; left hemi-mandible with p2, p4–m1 (IPS120168) of *Vulpes vulpes* from layer EVT3 in (**O**) lingual, (**P**) buccal, and (**Q**) occlusal view.

3.3.2. Canis mosbachensis

Historically, the appearance of *Canis* spp. marked the beginning of the biochronological event called the 'Wolf Event' around 2.0 Ma [142,143]. Recent discoveries pointed out an earlier European dispersal for canids. This evidence was based on scarce specimens attributed to *Canis* cf. *C. etruscus* from sites of Costa S. Giacomo FU (late Middle Villafranchian; [44]) and specimens of *Canis* sp. from the French site of Vialette [144]. The latter are dated back to the Early Villafranchian, around 3 Ma [144]. Therefore, these findings not only backdate the Wolf Event, but they show its diachronic nature (as noted by [145]), questioning its significance as a biochronological event [146]. In this sense, some authors [44,147] suggest using a different definition: the '*Pachycrocuta brevirostris* event', referring to the dispersal of the hyenid in Eurasia, would be a better designation of the faunal turnover around 2 Ma due to its ecological importance in Early Pleistocene habitats. The earliest well-documented evidence of the genus *Canis* from the Early Pleistocene of Europe has been recorded in the Italian sites of Olivola (Val di Magra, Tuscany, Italy; ca. 2 Ma; [44]) and Poggio Rosso (Upper Valdarno, Tuscany, Italy; ca. 1.9–1.8 Ma; [2,148,149]), with three species, *Canis arnensis, Canis etruscus*, and *Xenocyon falconeri* [38,125,142,150,151].

The taxonomical state of *Canis* species since their appearance in Europe has been widely studied [125–127,134,150,152–156], and different species have been described across Europe on the base of morphology and size. Nevertheless, their reciprocal phylogenetic relationships, as well as in comparison with extant canids, are still highly debated.

In the Iberian Peninsula, the first accounted finding of canids is that of Fonelas P-1 (Guadix Basin, Granada, Spain; approximately 1.9–1.7 Ma). From this site, four canids taxa have been recovered: *Vulpes alopecoides*, *C. etruscus*, *C. cf. falconeri*, and a new small species, *C. accitanus* [156,157]. Furthermore, the Guadix–Baza Basin has three of most important Early Pleistocene European sites, spanning approximately 1.4–1.2 Ma [158–160]: Venta Micena (around 1.4 Ma), Barranco León (around 1.3 Ma), and Fuente Nueva 3 (around 1.2 Ma). Along with remains of many Late Villafranchian mammal species, in all three of the sites remains, of the medium-sized *C. mosbachensis* have been found [23]. These findings are among the most ancient records of this taxon in Europe, and from this moment on, it becomes very common in the late Early Pleistocene and through the Middle Pleistocene; see, among others, [30,41,126,161].

Besides its wide diffusion, the true phylogenetic relationship of *C. mosbachensis* with other modern and fossil species is still highly debated. This wolf-like, medium-sized species for a long time has been considered to be derived from *C. etruscus* and to have then originated the modern wolf (among others [125,145,151,162–168]). Nevertheless, Soergel [169] argued the possibility of a close relationship between *C. mosbachensis* and *C. arnensis*, rather than *C. etruscus*. Furthermore, some authors [151,154] suggested the presence of a different lineage of wolf-like dogs in Southern Europe, parallel to the one of Central Europe of *C. mosbachensis*. Bartolini-Lucenti et al. [170] reveals that, around 1 Ma, the medium-sized canid present in the Iberian Peninsula was very close to that of the coeval German site of Untermassfeld [126] more than a more closely derived form of *C. arnensis* (Figure 3G–N). In either case, the overall dimensions of *C. mosbachensis* are smaller than those of *C. lupus*, and it probably exhibited a more hypocarnivorous diet as compared with its extant counterpart.

3.3.3. Vulpes alopecoides

The genus *Vulpes*, which includes fossil foxes, has been present since the Late Miocene in North America and the Old World. However, the fossil record of these foxes is extremely sparse and fragmented, often attributed to various species with uncertain relationships. The Early and Middle Pleistocene records of *Vulpes* in Europe are particularly confusing and have been the subject of much debate. Four species have been described from this period: *Vulpes alopecoides* from II Tasso (Upper Valdarno, Italy), *Vulpes praeglacialis* and *Vulpes praecorsac* from Villany (Hungary), and *Vulpes angustidens* from Hundsheim (Germany). The limited fossil evidence used to describe these species has led to considerable debate among scholars regarding their relationships to both extant foxes, such as *Vulpes vulpes* and *Vulpes lagopus*, and to each other.

Vulpes alopecoides has been reported from several Early Pleistocene sites across Europe, including Dafnero-1 in Greece [171], Dmanisi in Georgia [28,172], Fonelas-P1 in Spain [157], Kastritsi in Greece [171], La Puebla de Valverde in Spain [100], Makinia in Greece [171], Pirro Nord in Italy [30], Sesklon in Greece [171], St. Vallier in France [65], Upper Valdarno in Italy [34,150], Villarroya in Spain [173,174], and Volax in Greece [171]. In contrast, *Vulpes praeglacialis* has been found in later Early Pleistocene localities, such as Apollonia-1 in Greece [32], Atapuerca Trinchera Dolina 6 TD6 in Spain [127], Barranco Leon-5 in Spain [23], Cal Guardiola in Spain [23], Gombaszög/Gombasek in Slovakia [175], Caune de l'Arago in France [176], Deutsch Altenburg 2C in Austria [177], El Chaparral in Spain [23], Fuente

Nueva 3 in Spain [23], L'Escale in France [178], Le Vallonnet in France [20], Püspökfurdö-Betfia 2 in Romania [179], Venta Micena in Spain [23], and Villany 3–8 in Hungary [180].

The classification of these species has been further complicated by differing opinions among researchers. Kormos [181] suggested that *Vulpes praeglacialis* is closely related to the modern arctic fox, assigning it to the genus *Alopex* (an earlier name for *Vulpes lagopus*). Rabeder [177], studying material from Deutsch-Altenburg 2C, proposed that *Vulpes alopecoides, Vulpes praeglacialis,* and *Vulpes angustidens* from Hundsheim form a phyletic line leading to the contemporary *Vulpes vulpes.* This lineage is characterized by an increase in size and a slight reduction in molar size [129]. Rabeder also argued that *Vulpes praecorsac* is part of the lineage leading to *Vulpes corsac* but does not represent a direct ancestor–descendant relationship. He suggested that the arctic fox diverged from the *Vulpes* lineage during the Pliocene, predating the European fossil record and earlier than the interpretation by Wang et al. [182] regarding *Vulpes qiuzhudingi.*

Bartolini-Lucenti and Madurell-Malapeira [34] conducted a comparative study of the known variability in Pleistocene fox forms and their extant counterparts. They concluded that all Early Pleistocene fox remains should be classified under *Vulpes alopecoides*, as the observed tooth morphology variations among European Pleistocene taxa were less significant than those within each extant species.

In the Vallparadís Section, the remains of Early Pleistocene foxes are so scarce and limited to the post-Jaramillo layers CGRD7 and EVT7, where a partial hemi-mandible and a P4 were recovered, respectively, and attributed to *V. alopecoides* [34,69].

3.3.4. Vulpes vulpes

The red fox (*Vulpes vulpes*) first appeared in the Middle Pleistocene of Eurasia [115]. Pei [121,183] identified a few dental specimens from Locality 1 and several cranial fragments from Locality 3 of Zhoukoudian (ZKD) as 'V. cf. *vulgaris*'. However, age calibration of these ZKD localities remains problematic. According to Li et al. [184], the *V. vulpes* remains from layers 8 to 11 of Locality 1, dated around 700–780 ka, are the oldest in the sequence [137]. Locality 3 is considered younger than Locality 1, dating to more than 400 ka [137,138]. Although Pei's [121] attribution should be revisited, the Locality 1 specimens might represent the earliest record of *V. vulpes* in the Old World.

More certain records of *V. vulpes* come from the French sites of Lunel-Viel and L'Escale, with the oldest European occurrence dated to MIS12 [178]. A questionable Middle Pleistocene occurrence of *V. vulpes* is reported from Thomas Quarry 1, level 'Grotte des Hominides' in Morocco, correlated to approximately 600 thousand years ago [185]. However, Geraads [185] noted that the weathered mandible with worn premolars and carnassial from this site is of doubtful age and significance, possibly even sub-fossil. Late Pleistocene sites such as Doukkala and 'Grotte de Gazelles' in Morocco provide more reliable evidence of the red fox's presence in North Africa [185].

Despite the taxonomic confusion, *V. alopecoides* has generally been considered the ancestor of the modern red fox, *Vulpes vulpes* [34,177]. The European Middle Pleistocene record of *V. vulpes* is extremely sparse, first appearing in the French localities of l'Escale (ca. 0.6 Ma) [178] and Caune de l'Arago (ca. 0.6–0.45 Ma) [176]. In the Iberian Peninsula, red foxes have been identified from Sima de los Huesos (MIS12, ca. 0.45 Ma) and Galería (ca. 0.3 Ma) in Atapuerca [77]. In Italy, *V. vulpes* likely appeared slightly later, around the MIS12-11 transition (ca. 0.43 Ma), at sites such as Malagrotta and Torre in Pietra [186–188].

The precise chronology of *V. vulpes* appearance and its phylogenetic relationship with the earlier late Early Pleistocene *V. alopecoides* have not been thoroughly investigated. In the Iberian Peninsula, the earliest records of *V. vulpes* are from the Middle Pleistocene sites of Trinchera Galería (ca. 0.3 Ma) and Sima de los Huesos (MIS12, ca. 0.45 Ma) within the Atapuerca complex [77]. Although the Atapuerca fox remains have not been described in detail, García [77] compared the Galería specimens metrically with living and fossil foxes, noting close similarities with the extant species.

At the Middle Pleistocene layers of the Vallparadís Section, *Vulpes vulpes* remains are rarely recorded but were previously described by Madurell-Malapeira et al. [189] (Figure 3O–Q). The *Vulpes* specimens from EVT3 (ca. 0.6 Ma) closely resemble those of the modern *V. vulpes*, though slightly larger, and can be differentiated from *V. alopecoides* based on dental morphology. Therefore, the EVT3 sample, dated to 0.6 Ma, likely represents one of the earliest records of *V. vulpes* in Western Europe, pending a more detailed revision of the similarly aged sample from the French site of L'Escale.

3.4. Ursids

Ursus deningeri

The cave bear's lineage has been recorded in Europe since the beginning of the Pleistocene (ca. 2.6 Ma), likely related to the intensification of the Northern Hemisphere glacial processes and the establishment of the 40 ka obliquity-forced cyclicity. The first species of this lineage, *Ursus etruscus*, is known from European localities such as Saint Vallier, Upper Valdarno, or Kuruksay [54,65]. Here, we support the traditional viewpoint of Kurtén [82], who first hypothesized a phylogenetic line for cave bears starting with the Early Pleistocene *U. etruscus*, followed by the Middle Pleistocene *Ursus deningeri* and finally in the Late Pleistocene for *Ursus spelaeus* s.l. Calabrian cave bears inhabited mixed environments of woodlands and/or wooded grasslands, where they fed on a broad variety of food items, with a substantial intake of meat and/or fish [54]. The trend in the derivation of morphological characters had an abrupt change since 1.2 Ma, when the first specimens of *Ursus deningeri* were recorded from Vallonnet cave as displaying clear speleoid characters and a large size [20]. This tendency towards herbivory is also documented in Untermassfeld and Gran Dolina (MIS31 and MIS21-19, respectively) [77].

In the Vallparadís Section, cave bears were abundantly recorded from MIS30 to MIS21 in layers EVT12, CGRD7, and EVT7. These specimens show significant intraspecific variability and sexual dimorphism, favoring our previous idea that the *Early–Middle Pleistocene Transition* associated with the climatic shifts was the driving force behind the final changes that conducted the cave bear's lineage to hyperherbivory behavior [21–23,54] (Figures 4 and 5).

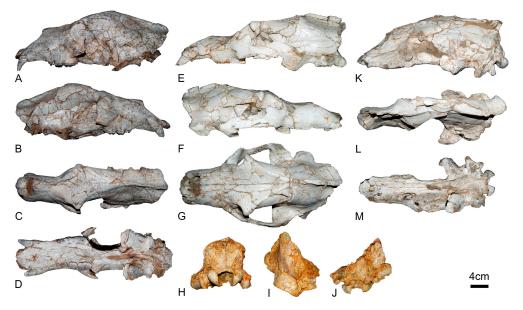


Figure 4. Crania of *Ursus deningeri* from the Vallparadís Section. Cranium with left and right C1, P4–M2 (EVT24876) from layer EVT12 in (**A**) left lateral, (**B**) right lateral, (**C**) dorsal, and (**D**) ventral views; cranium with left and right C1 and P4–M2 (EVT15872) from layer EVT7 in (**E**) left lateral, (**F**) right lateral, and (**G**) dorsal views; cranium with right m2 (EVT7116) from layer EVT7 in (**K**) left lateral, (**L**) dorsal, and (**M**) ventral views; basicranial fragment (IPS14951) from layer CGRD7 in (**H**) occipital, (**I**) dorsal, and (**J**) ventral views.

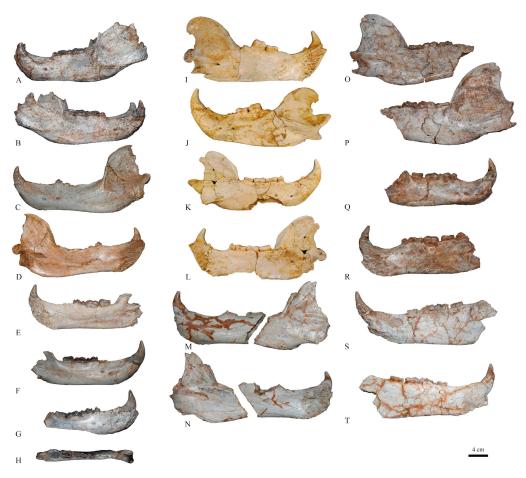


Figure 5. Mandibular remains of *Ursus deningeri* from the Vallparadís Section. Mandible with left and right c1–m3 (EVT24876) from layer EVT12 in (**A**) left buccal and (**B**) right buccal views; left hemimandible (EVT25680) from layer EVT12 in (**C**) buccal and (**D**) lingual views; right hemi-mandible with c1 and m1–m3 (EVT20710) from layer EVT12 in (**E**) lingual and (**F**) buccal views; partial right hemi-mandible with c1 and p4–m3 (EVT24876b) from layer EVT12 in (**G**) buccal and (**H**) occlusal views; mandible with left c1, p3, and m1–m2 and right c1 and p1–m3 (IPS14950) from layer CGRD7 in (**I**) left buccal, (**K**) right buccal, and (**L**) right lingual views; left hemi-mandible with c1 (EVT20234) from layer EVT7 in (**M**) buccal and (**N**) lingual views; right hemi-mandible with c1–m3 (EVT842) from layer EVT6 in (**O**) lingual and (**P**) buccal views; right hemi-mandible with c1–m3 (EVT920) from layer EVT6 in (**Q**) buccal and (**R**) lingual views.

In a recent study on the microwear and isotopic samples of the former specimens, Vizcaíno-Varo [190] found that the Vallparadís ursids are at a midpoint between the generalist omnivore behavior of *U. etruscus* and the hyperherbivory behavior of the Late Pleistocene *U. speleaus*. Additionally, the isotopic evidence shows that, probably around MIS21, these bears started hibernating and, consequently, habiting in karstic environments during winters, coinciding with the harshening of climatic conditions [190].

3.5. Mustelids

Meles meles

Badgers of the genus *Meles* have been reported from Eurasian localities since the Late Pliocene (3.5–3 Ma); their fossil record is, however, very scarce. This caused confusion among the taxonomy and prevented any good phylogenetic reconstruction of the genus. The first species to appear, almost simultaneously, are *Meles thorali* in Europe and *Meles chiai* in China [191–193]. Even if this synchronous appearance makes it difficult to hypothesize where the genus originated, the timeframe coincides with the progressive transition of

northern hemisphere biocoenosis from a subtropical, predominantly wooded habitat to more open environments and mixed forests with a progressive increase in seasonality [8]. Until recently, the earliest known evidence of *Meles meles* was found at Fuente Nueva 3, dating back to around 1.2 million years ago. However, Marciszak et al. [194] reported several *M. meles* remains from Żabia Cave in Poland, a site estimated to be between 1.7 and 1.5 Ma. If this finding is verified, it will bridge the temporal gap between the latest records of *M. thorali* at Apollonia-1 (around 1.5 Ma) and the earliest European badgers. The presence of *M. meles* at Żabia Cave could indicate that extant European badgers likely evolved from an *M. thorali*-like ancestor in Asia and then migrated to Europe, where they gradually replaced *M. thorali*.

The European badger (*M. meles*) is the only mustelid recorded in the Vallparadís Section. Abundant cranial remains of this species were recovered from the Lower and Middle Units (layers EVT10, EVT11, and EVT12; and layers EVT6, EVT7, and EVT8; [23] and references therein). The *M. meles* specimens from Vallparadís display a large intraspecific variability, characteristic of the genus, often leading to great taxonomic confusion. This appears particularly evident when comparing the two most complete skulls of Vallparadís, as they have very different sizes. However, the robust skull with a large anteriorly placed infraorbital foramen and the distolingually placed metaconule of M1 are characteristic of *M. meles* and differentiate it from *M. thorali* (Figure 6).

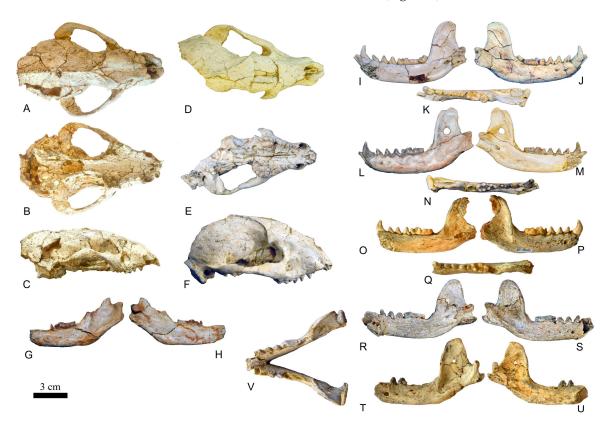


Figure 6. *Meles meles* remains recovered from the Vallparadís Section. Cranium with left and right p4–m1 (EVT25417) from layer EVT12 in (**A**) dorsal, (**B**) ventral, and (**C**) right lateral views; partial cranium with left and right p2–m1 (IPS94288) from layer EVT12 in (**D**) dorsal, (**E**) ventral, and (**F**) right lateral views; left partial hemi-mandible with m1 (IPS122011) from layer EVT7 in (**G**) buccal and (**H**) lingual views; right hemi-mandible with i1–m2 (IPS122008) from layer EVT12 in (**I**) lingual, (**J**) buccal, and (**K**) occlusal views; left hemi-mandible with i3–m1 (IPS122009) from layer EVT12 in (**L**) buccal, (**M**) lingual and (**N**) occlusal views; right hemi-mandible with c1–m1 (IPS122010) from layer EVT12 in (**O**) lingual, (**P**) buccal, and (**Q**) occlusal views; mandible with left p2 + p3 and right p1–m1 (IPS94288) from layer EVT12 in (**R**) right lingual, (**S**) right buccal, (**T**) left buccal, (**U**) left lingual, and (**V**) occlusal views.

It is unclear whether the emergence of the modern badger species reflects a general trend of progressive omnivorous adaptations within the lineage itself or is due to specific environmental conditions. Unfortunately, the few post-cranial remains of *Meles* recovered from Vallparadís (a radius and three phalanges) do not allow for inferences to be made on the evolution of the dietary behavior of this genus.

3.6. Vallparadís Guild Structure: Composition and Ecological Comparison with Other Early Pleistocene Guilds

The results of the BCA are reported in Figure 7A. The analysis clearly separates two large groups: a group of African sites (node 1) and one of all the Eurasian localities (node 3). The first group that branches out includes the East and North African guilds. Within this, the Shungura Formation of Omo, levels G–K, stems at the base of the cluster. The major difference with the other sites includes the presence of Helogale gr. hirtula and Megantereon whitei. Within the African cluster, two subgroupings branch from node 2 (percentage *p*-value = 70%): The first one (node 7) clusters the guilds of Koobi Fora Okote Fm. and that of Tighennif. The two localities have similar age and share several species (e.g., Crocuta crocuta, Hyaena hyaena, and Panthera leo). The percentage p-value (=71%) of this node is a testament of their own peculiarities. The second subcluster is that of node 9 and is well supported (percentage *p*-value = 94%). The subcluster includes East African localities with an age between ca. 1.9 and 1.2 Ma. All three localities share numerous taxa (Panthera pardus, Pseudocivetta ingens, Crocuta crocuta, and Hyaena hyaena), but the Olduvai Bed 1 equally differs from Koobi Fora KBS Fm. and Olduvai Bed 2 from a more diverse guild for exclusive carnivorans (e.g., Xenocyon africanus or Otocyon recki) and for occurrences shared only with one of the sites clustered together at node 14 (e.g., the Lutra in both Olduvai and the Lupulella in the coeval Bed 1 and KBS Fm.). The larger cluster composed of Eurasian localities (node 3) is further subdivided into numerous subgroups. Node 6 identifies the separation of Early-Middle Pleistocene Northeastern Chinese sites of Jinyuan Cave Upper levels and Zhoukoudian Locality 1. Their grouping is justified by some shared elements, e.g., Canis variabilis and Vulpes chikushanensis, and also for the peculiarity in comparison to other Asian sites. Nevertheless, the percentage *p*-value is not among the highest, although it is fairly high (p-value = 78%). The second group is that of the node 13 (percentage p-value = 98%) group, together with the Jinyuan Cave Lower levels and Nihewan Classic Fauna. As in the case of the cluster of node 6, chronological and geographic proximity might be the reason for the clusterization of these sites. They indeed share a large number of taxa (for a total of eleven species; see Supplementary Table S1). A similar reason might lie at the base of the grouping of node 11 (percentage *p*-value = 97%), that of *Gigantopithecus* Cave and of Gongwangling, two sites of Central–Southern China. The grouping is supported by the presence of taxa exclusive that are to these sites, e.g., Ailuropoda melanoleuca and Panthera pardus, or rarer taxa, like Arctonyx and Felis sylvestris. Node 8 denotes a larger set of European and Western Asian localities of Dmanisi and 'Ubeidiya. Trinchera Dolina TD6 is the first site stemming out from node 8: this position is due to the presence of clear European species (e.g., Lynx pardinus, Canis mosbachensis, and Vulpes alopecoides) but also new and rare occurrences for Early Pleistocene, like Crocuta crocuta. The grouping of Poggio Rosso and Casa Frata is well supported (percentage p-value = 94%) and justified by the close chronological and geographical position and their taxonomical composition (with six shared taxa, e.g., Canis arnensis, Lynx issiodorensis, and Ursus etruscus). The cluster of node 11 has 'Ubeidiya at its base. The Israeli site is characterized by the compresence of Eurasian and out-of-Africa taxa that are clearly affine with late Calabrian European sites (e.g., Canis mosbachensis, Xenocyon lycaonoides, and Panthera gombaszoegensis). The other localities are organized in two groups: one of earlier localities from Georgia and Southern Europe (node 16) and one of Epivillafranchian sites (node 18). In the first one, Dmanisi stems from the group made of Pirro Nord DE, Venta Micena, and Apollonia-1. At the level of carnivore guild, there is very little difference in terms of composition between the four sites: they share nine carnivores (Supplementary Table S1). This is particularly true especially for

Venta Micena and Apollonia-1, which share all the taxa (see Supplementary Table S1). A similar situation is that of the four Epivillafranchian sites (among them Vallparadís ones). These sites have six taxa in common (*Meles meles, Lynx pardinus, Pachycrocuta brevirostris, Canis mosbachensis, Xenocyon lycaonoides,* and *Ursus deningeri*), and two more are shared between EVT7/CGRD7, Vallonnet, and Untermassfeld (reaching the number of eight, respectively, nearly the 73%, the 62%, and the 73% of their record).

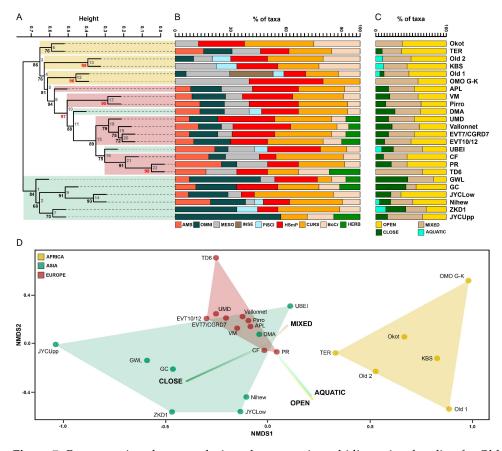


Figure 7. Bootstrapping cluster analysis and non-metric multidimensional scaling for Old World Early Pleistocene sites. (A) Dendrogram resulting from the bootstrapping cluster analysis on presence/absence of taxa in selected sites. (B) Histograms of the relative proportions of the ecological preference of the carnivorans in the analyzed guilds of the Old World, arranged according to a BCA based on ecological matrix of the sites. (C) Histograms of the relative proportions of the ecological preference of the carnivorans in the analyzed guilds of the Old World, arranged according to a BCA based on environmental matrix of the sites. (D) Non-metric multidimensional scaling plot on the environmental parameters of Old World Early Pleistocene sites. Abbreviations: AMB, hypercarnivorous ambush predators; BoCr, hypercarnivorous species with bone-cracking adaptations; CURS, hypercarnivorous cursorial predators; HERB, hypocarnivorous adaptations towards herbivory; HSmP, small-sized hypercarnivorous species; INSE, insectivorous species; MESO, mesocarnivores; OMNI, omnivorous, hypocarnivorous, and opportunistic feeders; PISC, hypercarnivores adapted to a piscivorous diet; APL, Apollonia-1 (Greece); CF, Casa Frata (Italy); DMA, Dmanisi (Georgia); EVT10/12, Vallparadís Estacio layers 10-12 (Spain); EVT7/CGRD7, Vallparadís Estacio layer 7/Cal Guardiola 7 (Spain); GWC, Gongwangling (China); JYCLow, Jinyuan Cave Lower fauna (China); JYCUpp, Jinyuan Cave Upper Fauna (China); KBS, KBS Member, Koobi Fora Formation (Kenya); Nihew, Xiashagou, Classic Nihewan (China); Okot, Okote Member, Koobi Fora Formation (Kenya); Old 1, Olduvai Bed I (Tanzania); Old 2, Olduvai Bed II (Tanzania); OMO G-K, Members GeK, Shungura Formation (Ethiopia); Pirro, Pirro Nord (Italy); PR, Poggio Rosso (Italy); TD6, Trincher Dolina 6 (Spain); UBEI, 'Ubeidiya (Israel); UMD, Untermassfeld (Germany); Vallonnet, Vallonnet (France); VM, Venta Micena (Spain); ZKD1, Zhoukoudian Locality 1 (China).

Figure 7C reports the result of NMDS on the occurrences reported in each site, taking into consideration the environmental preferences of the taxa. In the plot, the distribution of the localities reflects the similarity between them (i.e., the closer the sites, the more similar) and generally confirms the results of the BCA, as described above. In the graph, European localities are greatly associated with one another, despite their chronological differences. Among these, particularly the sites of Poggio Rosso and Casa Frata have carnivore guilds similar between them but fairly distinct from the group composed of late Early Pleistocene of Venta Micena, Pirro Nord, Vallonnet, and Vallparadís layers. The exception among the European sites is Trinchera Dolina TD6, which lies greatly separated along the NMDS Axis 2. The limited number of carnivorans and the occurrence of peculiar elements, chiefly Crocuta crocuta and Cuon alpinus, might explain this distance from the other European guilds. Asian localities occupy a wide space in the NMDS. For instance, the (nearly) coeval Gigantopithecus Cave, Nihewan Classic Fauna, and Jinyuan Cave Lower Fauna are separated in two positions, with the latter two much closer to each other than to the Southern Chinese locality of Gigantopithecus Cave. In turn, this site is closer to the Late Calabrian site of Gongwanling. Regarding temporal distinctions, Jinyuan Cave Upper Fauna is more distant from the Lower Fauna than from any other East Asian locality. This is possibly due to the arrival of new taxa like Canis variabilis and Ursus deningeri. The two sites of Dmanisi and 'Ubeidiya are located considerably far from the Eastern Asian ones and much closer to European sites. Dmanisi lies close to Apollonia-1 and Venta Micena, as their carnivoran guilds share numerous taxa (e.g., Pachycrocuta brevirostris, Homotherium crenatidens, Panthera gombaszoegensis, Xenocyon lycaonoides, and Ursus etruscus). 'Ubeidiya differs slightly from the European sites because of the mixture of African/Eurasian elements, such as Crocuta crocuta, Vormela peregusna, Canis mosbachensis, and Ursus etruscus. Far on the positive end of the NMDS 1 Axis, the African sites make up a separate and distinguishable group. Among them, we see that there is some kind of difference, as there is no evident clustering based on age or in terms of geographic or stratigraphic proximity. Tighennif is the most similar to European localities, given the number of European taxa recorded in the Algerian site. The high NMDS 2 Axis values of Shungura Fm. of Omo, levels G-K, might be explained by the presence of unique taxa (e.g., *Helogale*) not recorded in the other selected sites and the limited number of occurrences.

4. Discussion

4.1. The Vallparadís Section Carnivore Guild: Ecological and Taxonomical Comparison

The results of the BCA and the NMDS (Figure 7) show the different grouping of the carnivore guilds of Africa and Eurasia. The various guilds are correctly clustered into groups which testify the similarities but also the peculiarities between one another, for instance, the cluster of African guilds. These sites are separated by the other ones both taxonomically and ecologically (Figure 7), confirming previous results [28]. It is worth mentioning the differences in environmental preferences among African guilds: the large dominance of open habitat taxa in all guilds with different proportions in terms of aquatic and mixed habitat species. Another well-characterized grouping is that of Gigantopithecus Fauna and Gongwangling, as opposed to that of Nihewan Classic Fauna–Jinyuan Cave Lower Fauna and of Jinyuan Cave Upper Fauna-Zhoukoudian Loc. 1. This is evident in both BCA and in NDMS. Furthermore, the latter enhances the differences between clusters, adding the environmental parameters. The pattern of Asian localities might be the signal of the complete pattern of temporal and spatial turnover/latitudinal distribution and segregation that characterized and still characterize Asia today. Following the preferences of the carnivorans from the Asian sites supports a certain distinction in terms of habitats, with Gigantopithecus Fauna, Gongwangling, and Jinyuan Cave Upper Fauna dominated by a closed environment, in comparison to Nihewan Classic Fauna, Jinyuan Cave Lower Fauna, and Zhoukoudian Loc. 1, in which taxa of a more open environment dominate. The European localities are well clusterized together, with a clear affinity between Dmanisi and 'Ubeidiya, as reported and discussed in other previous works [27,28,33,41]. This

clusterization and the position in the NMDS space suggest that a majority of the European localities were characterized by relatively mixed environments, with no dominance between open- or closed-environment taxa. Among the European locality, only the Upper Valdarno localities of Poggio Rosso and Casa Frata are marked by proportionally more abundant open habitat taxa, in agreement with current understanding of the environments of Upper Valdarno Basin around 1.9–1.8 Ma [31,40]. In comparison to other considered localities from Asia, Africa, and Europe, the carnivore guild from the Vallparadís Section is, unsurprisingly, close to other Western European Epivillafranchian localities, particularly Untermassfeld and Vallonnet. This is consistently supported by the taxonomic composition of their guilds, equally composed by common elements like *Meles meles, Lynx pardinus*, and *Ursus deningeri*. This is confirmed by both the BCA and the NMDS (Figure 7).

4.2. The Vallparadís Section Carnivorans in the Iberian Context

The latest Early Pleistocene (Epivillafranchian) is not abundantly recorded in the Iberian Peninsula, with few sites exhibiting low diversity of recorded carnivorans, precluding accurate comparisons with the Vallparadís Section guild. On the Northeastern Iberia, the sites of the Incarcal complex basically have a good record of two carnivoran species (*H. crenatidens* and *P. brevirostris*) correlated with MIS21 [23]. However, at the close Bòvila Ordis site, no carnivore was recorded in the layers with an approximate age of 1.2 Ma [23]. Further south, in the Francolí valley, the recently discovered site of Barranc de la Boella, with an approximate age of 0.9 Ma, records *P. gombaszoegensis*, Hyaenidae indet., *U. deningeri*, and *Canis* sp. Unfortunately, there are only a few poorly preserved specimens [25].

In the Southern Iberian Peninsula, the site of Húescar-1, with a debated chronology around 0.8 Ma, records very few specimens of *H. crenatidens*, *P. gombaszoegensis*, Hyaenidae indet., and *C. mosbachensis* [195]. In Cadiz, in the Chaparral site [106], well-preserved but scarce specimens of *V. pardoides*, *C. mosbachensis*, and *V. alopecoides* have been recovered. A similar low diversity of carnivorans was documented in the Quibas site [18] (MIS30), with scarce records of *Lynx pardinus* and few indeterminate canid fragments.

Particularly interesting in this discussion is the site of Cueva Negra del Estrecho del Rio Quípar, with an estimated chronology of 990–772 ka [196]. The former authors describe carnivoran remains attributable to *Ursus* sp., Mustelidae indet., *Crocuta* sp., and *Lynx* sp. Probably the most interesting in this assemblage is the putative presence of a spotted hyena, coeval with the earliest records of this genus outside Africa in Trinchera Dolina [77]. However, a quick examination of the remains attributed to *Crocuta* open serious doubts on former authors' attribution. *Crocuta* remains are not described in detail and only discussed based on a biometrical analysis of the lower p4 of a partial and poorly preserved right hemi-mandible [196]. The dimensions of the p4 fit perfectly with the specimens of the Vallparadís Section, securely attributed to *Pachycrocuta*. Additionally, the presence of mesial and distal accessory cuspulids on the p4 and the massive and high mandibular symphysis clearly favors its attribution to *P. brevirostris* [196].

A similar scenario is observed in the Early Pleistocene layers of the the Atapuerca site complex. Specifically, the layer TE9c of the Sima del Elefante site, with an estimated chronology of 1.2 Ma [197], evidences the presence of *P. gombaszoegensis, Lynx pardinus,* cf. *Pannonictis,* cf. *Baranogale, Mustela* sp., *C. mosbachensis, V. alopecoides,* and *U. deningeri* [197]. In the Trinchera Dolina TD6 layer [77], dated approximately 0.9 Ma, the recorded species includes *U. deningeri, C. crocuta, M. palerminea, L. pardinus, C. mosbachensis,* and *V. alopecoides* [77]. Other layers in the complex also include *H. crenatidens* and various carnivorans never described in detail. Despite the abundance of small mustelid species, which are extremely rare in the Iberian record, the most significant highlight of the Atapuerca site is the confirmed presence of the genus *Crocuta* in Early Pleistocene layers [77].

Finally, the most diverse carnivore guild on Iberian Early Pleistocene is found at the Cueva Victoria site [23], with an estimated chronology of 1.0–0.8 Ma; however, several doubts about the stratigraphical provenance of some specimens exist. The record includes

H. crenatidens, M. adroveri, P. gombaszoegensis, V. pardoides, L. pardinus, P. brevirostris, U. deningeri, X. lycaonoides, C. mosbachensis, and *V. alopecoides.* Additionally, an unpublished third metatarsal is clearly attributable to *P. s. fossilis.* This guild is the same as the one recorded in the Vallparadís Section; however the uncertainties about their stratigraphical scheme preclude detailed discussion on their implications.

To summarize, the carnivoran guild of the latest Iberian Early Pleistocene shows no significant changes compared to the previous Late Villafranchian guild recorded at sites like Venta Micena [23], at least until MIS30 (ca. 1.0 Ma). Since MIS30, African-origin carnivorans such as *P. s. fossilis* and *C. crocuta* started to being recorded in Iberia but only in two sites or complexes, probably due to the scarce and fragmentary record of the Iberian Epivillafranchian record. Despite no secure records of *P. pardus* during this period, the main causes of the disappearance of several species may include the competition with newcomer species and the environmental changes associated with the onset of the *Early–Middle Pleistocene Transition* (i.e., increase in seasonality, aridity, and dissymmetrical climatic cycles prompted by an increase in wood cover as compared with previous Late Villafranchian).

4.3. The Vallparadís Section's Carnivorans in the Eurasian Context

The scenario for the latest Early Pleistocene sites in Europe is similar to the Iberian one, characterized by several sites, predominantly in Mediterranean Europe, with poorly recorded carnivoran guilds. The European sites from the 1.2–0.8 Ma interval include Collecurti, Cava Rediccioli, Slivia, and Frantoio in Italy [198,199]; Apollonia-1 in Greece [32]; Blassac-la-Girondie, Bois-de-Riquet, Cagnes-sur-Mer, Ceyssaguet, Chagny, Rosières, Vallonnet, Soleilhac, Saint-Prest, Sartanette, Tour-de-Grimaldi, Trois Pigeons, and Durfort in France [39,200]; Untermassfeld in Germany [201]; Happisburgh in England [202]; Trilica in Montenegro [203]; Somssich Hill 2 in Hungary [204]; or Akhalkalaki in Georgia [205].

Among the former list, only Untermassfeld, Vallonnet, and Ceyssaguet exhibit a comparable or even most diverse carnivoran guild, which includes *H. crenatidens*, *M. adroveri*, *P. gombaszoegensis*, *A. pardinensis*, *V. pardoides*, *L. pardinus*, *F. sylvetris*, *P. brevirostris*, *U. deningeri*, *X. lycaonoides*, *C. mosbachensis*, *V. alopecoides*, and *M. meles*. Additionally, at Le Vallonnet, two lower molars of *P. pardus* were personally studied by the authors, supporting the taxonomic attribution made by [20].

Despite the different taxonomical attributions provided by different scholars, the Epivillafranchian carnivore guild of Europe displays homogeneity, except for the scarcely recorded African immigrants (lion, leopard, and spotted hyaena). However, several minor issues need to be addressed.

Firstly, since the beginning of the Pleistocene, U. etruscus was the only ursid recorded in European assemblages. This omnivorous specie, adapted to moderately wooded environments, was recorded at numerous sites. However, starting around 1.2 Ma and probably influenced by the climatic instability associated with the Early-Middle Pleistocene Transition, these forms progressively become stouter, larger, and more sexually dimorphic, starting their transition to a purely herbivorous diet [190]. Ursus deningeri was first recorded at the Vallonnet cave but also at the Vallparadís Section and Ceyssaguet. Alternatively, in Trinchera Dolina, Frantoio, and Untermassfeld, the potential same derived form has been cited under the attribution of *U. dolinensis* [77]. Secondly, in previous papers, one of us (J. M.-M.) hypothesized an Iberian origin for the Mediterranean lynx, first recorded at the Avenc Marcel cave (1.6 Ma) [113,114]. Recent research studies demonstrated that, by the end of Late Villafranchian and Epivillafranchian, L. pardinus was already distributed throughout all of Mediterranean Europe. This species was present at sites such as Pirro Nord, Vallonnet, Ceyssaguet, the Vallparadís Section, Cueva Victoria, or Untermassfeld, making it difficult to determine the precise geographical origin of the species. A more parsimonious hypothesis suggests a gradual transition from L. issiodorensis-like forms to smaller L. pardinus-like ones. Lastly, the evolutive history of the wild cat (*Felis sylvestris*) remains poorly understood. The presence of a mandible of this species in the approximately 2.0 Ma layers of the Italian Upper Valdarno complex is known. However, until the late Middle Pleistocene, it seems that this species did not exist in

Europe. Despite the exceedingly limited record, often consisting of one or two fragments, this evidence from the Epivillafranchian European sites pointed to a continuous presence of the wild cat in Europe.

In summary, the Calabrian carnivore guild remained stable throughout the Late Villafranchian but began to decline with the disappearance of several Villafranchian-character species during the arid phase associated with MIS30. This period is precisely recorded at the Vallparadís Section, which also marks the first appearance of lions and the last occurrence of dirk-toothed cats (layers EVT10-12). The various layers of the Vallparadís Section have become a key reference for studying these changes in the carnivoran guild. Future stable isotope and microwear analyses will provide further insights into the impact of the Early–Middle Pleistocene Transition on European taphocenosis [190].

5. Conclusions

The *Early–Middle Pleistocene Transition* in the Northern Hemisphere marked a shift from the predominantly obliquity-forced cyclicity to the strongly asymmetric and severely cold precession-forced cyclicity characteristic of the Middle and Late Pleistocene. These changes, along with increased aridity and seasonality, profoundly affected the European late Early Pleistocene taphocenosis. The Vallparadís Section, with its multiple layers comprising a 12 m thick sequence, is currently the only European site where the impact on large mammal assemblages during this period can be studied in detail. This period roughly coincides with the transition from Oldowan to Acheulian culture in Western Europe.

In this study, we detail the carnivoran guild composition of the Vallparadís Section and its changes over time, comparing it with other Iberian and European sites. The results show the stability of most of the guild during the Calabrian Early Pleistocene, with the arrival of several newcomers of African origin (e.g., steppe lions). These long-surviving Calabrian species are recorded until the interglacial stage MIS31 in layers EVT7 and CGRD7 of the section (e.g., *Megantereon* or *Xenocyon*).

Supplementary Materials: The following supporting information can be downloaded at https:// www.mdpi.com/article/10.3390/quat7030040/s1, Figure S1: Geographical location of the Vallparadís Section within the Iberian Peninsula and the city of Terrassa. Additionally, composite stratigraphic section with the layer of precedence of the studied specimens; Figure S2: Map of the Old World showing the considered localities in the present study. Number: 1 = Vallparadís Section (Iberia); 2 = Apollonia-1 (Greece); 3 = Casa Frata (Italy); 4 = Dmanisi (Georgia); 5 = Gongwangling (China); 6 = Jinyuan Cave Lower Fauna (China); 7 = Jinyuan Cave Upper Fauna (China); 8 = KBS Member, Koobi Fora Formation (Kenya); 9 = Xiashagou, Classic Nihewan (China); 10 = Okote Member, Koobi Fora Formation (Kenya); 11 = Olduvai Bed I (Tanzania); 12 = Olduvai Bed II (Tanzania); 13 = Members G-K, Shungura Formation (Ethiopia); 14 = Pirro Nord (Italy); 15 = Poggio Rosso (Italy); 16 = Trinchera Dolina 6 (Spain); 17 = Ubeidiya (Israel); 18 = Untermassfeld (Germany); 19 = Vallonnet (France); 20 = Venta Micena (Spain); 21 = Zhoukoudian Locality 1 (China); Table S1: Database of species occurrences per selected sites used in the analysis. Abbreviations: APL, Apollonia-1 (Greece); CF, Casa Frata (Italy); DMA, Dmanisi (Georgia); EVT10/12, Vallparadís Estacio layers 10-12 (Spain); EVT7/CGRD7, Vallparadís Estacio layer 7/Cal Guardiola 7 (Spain); GWC, Gongwangling (China); JYCLow, Jinyuan Cave Lower Fauna (China); JYCUpp, Jinyuan Cave Upper Fauna (China); KBS, KBS Member, Koobi Fora Formation (Kenya); Nihew, Xiashagou, Classic Nihewan (China); Okot, Okote Member, Koobi Fora Formation (Kenya); Old 1, Olduvai Bed I (Tanzania); Old 2, Olduvai Bed II (Tanzania); OMO G-K, Members GeK, Shungura Formation (Ethiopia); Pirro, Pirro Nord (Italy); PR, Poggio Rosso (Italy); TD6, Trinchera Dolina 6 (Spain); UBEI, 'Ubeidiya (Israel); UMD, Untermassfeld (Germany); Vallonnet, Vallonnet (France); VM, Venta Micena (Spain); ZKD1,. Zhoukoudian Locality 1 (China). References: [9,21,23,28-44].

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