

New bioclimatic models for the quaternary palaearctic based on insectivore and rodent communities.

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16	

17 Abstract

Mammal remains, preserved in archaeological and palaeontological deposits, are commonly used 18 19 to reconstruct past terrestrial climates and environments. Here we propose new species-specific 20 models for Bioclimatic Analysis, a palaeoclimatic method based on a climatic restriction index for 21 each mammal species, discriminant analysis, and multiple linear regressions. Our new models are 22 based on small mammal associations, particularly insectivores and rodents, from Quaternary paleoarctic contexts. A dataset including new localities and an updated taxonomy was constructed 23 24 in order to develop two approaches, the first using only Rodentia, the second based on associations 25 including both Rodentia and Eulipotyphla. Both approaches proved to be reliable for inferring both climate zone and quantifying temperature, precipitation, and seasonality. Rarefaction analysis 26 27 revealed these new models to be reliable even when a substantial percentage of species from the 28 original palaeocommunity was absent from the fossil site. Application of these new models to 29 small mammal associations from two sequences (Balma de l'Abeurador, France and El Mirón, 30 Spain) spanning from the Last Glacial Maximum to the Holocene are consistent with the primary 31 climatic changes recorded by regional Pyrenean proxies and showed an increase in mean annual 32 temperature of between 3 and 5°C.

33 Key-words

34 Small mammalian communities; Paleoclimatology; Paleoecology; Pleistocene; Climate
 35 reconstructions

36

37 1. Introduction

38 Understanding past climatic and environmental changes and their impact on continental 39 ecosystems remains a major challenge, particularly in terms of human biological and social 40 evolution. Currently, environmental reconstructions are built from mineralogical, geochemical, 41 floral, and faunal data recovered from palaeontological or archaeological sites (e.g. Carrasco et al., 42 2008; Frouin et al., 2013; Domingo et al., 2015; Menéndez et al., 2017; Villa et al., 2018; Britton 43 et al., 2019; Girard et al., 2019). Vertebrates are particularly sensitive to changes in climate and 44 habitat (e.g., Faunmap, 1996; Thomas et al., 2004), and it is generally assumed that these changes induce new biotic conditions, which, in turn, constrain vertebrate species distributions, biotic 45 46 interactions and, ultimately, community organization (Lyons, 2003; Moritz et al., 2008; Hernández Fernández et al., 2015; Řičánková et al., 2015; Royer et al., 2016; Blanco et al., 2018). Based on 47 this assumption, numerous methods have been developed to reconstruct environmental conditions 48 49 and quantify past climate parameters from mammalian fossil remains (for syntheses see Artemiou, 50 1984; Andrews, 1996; Lyman, 2017; Nieto & Rodríguez, 2003). Several such models are based 51 on modern species distributions and ecological niches (e.g., Atkinson et al., 1987; López García, 52 2008; Jeannet, 2010; Fagoaga et al., 2019), while others employ species richness (Montuire et al., 1997; Legendre et al., 2005; Araújo et al., 2008). Relatively few, however, utilize whole mammal 53 communities (Valverde, 1964; Legendre, 1986, 1989; Kay and Madden, 1997; García Yelo et al., 54 55 2014; Escarguel et al., 2008) or large mammal species associations (de Bonis et al., 1992; Griggo, 1996; Hernández Fernández and Vrba, 2006). Most environmental reconstructions are, in fact, 56 57 based on small mammals or uniquely rodents (Hokr, 1951; van de Weerd and Daams, 1978; Daams and van der Meulen, 1984; Chaline and Brochet, 1989; Sesé, 1991; van der Meulen and Daams, 58 1992; van Dam, 2006), as these species present several unique particularities. First, they are 59 60 frequently preserved in archaeological and palaeontological deposits in caves and rock shelters,

61 sometimes in association with human remains and artifacts and, unlike larger mammals, are less 62 susceptible to human-induced biases (e.g., Chaline, 1977; Discamps and Royer, 2017). Second, small mammals are key prey for numerous predators and inhabit a wide variety of ecological 63 64 habitats (e.g., Quéré and Le Louarn, 2011; Wilson et al., 2017, 2018). Finally, due to their small size and large taxonomic diversity, rodent communities reflect local environments and are highly 65 66 sensitive to environmental parameters and climatic changes, leading to their being considered one 67 of the most reliable proxies for inferring past environmental and climatic conditions (e.g., Hinton, 1926; Tchernov, 1975; Avery, 1987; Montuire and Desclaux, 1997; Hernández Fernández et al., 68 69 2007; García-Alix et al., 2008; Cuenca-Bescós et al., 2009; López García et al., 2010; Cuenca-70 Bescós et al., 2011; Royer et al., 2013, 2014; Rofes et al., 2015; Royer et al., 2016; Fernández-71 García et al., 2016; Laplana et al., 2016; Piñero et al., 2016; Berto et al., 2017; López García et al., 72 2017; Berto et al., 2019; Stoetzel et al., 2019; Fernández-García et al., 2020).

73 Among the many methods for reconstructing palaeonvironments, Bioclimatic Analysis 74 (Hernández Fernández, 2001) produces both qualitative and quantitative climatic reconstructions 75 built from the structure of mammal communities, including multiple taxonomic orders, and their 76 global distribution along broad climatic gradients and biozones. One of its main advantages is that 77 entire mammal communities are taken into consideration, giving equal value to all species as 78 climatic indicators related to their ecological tolerance. Here we further refine this approach by 79 using Eulipotyphla (moles, shrews and hedgehogs, commonly referred to simply as insectivores) 80 combined with Rodentia (rodents), or uniquely Rodentia recovered from Pleistocene to Late 81 Holocene Palaearctic faunal assemblages. Although recent research has begun to combine rodents 82 and insectivores in analyzes, these small mammals had almost always been previously considered 83 separately by specialists. Fossil material from palaeontological and archaeological sites commonly

84 result from the same accumulation agent and are recovered using similar excavation methods. 85 Moreover, Eulipotyphia are regularly preserved in European fossil sites and are highly sensitive to several climatic parameters, such as rainfall, (e.g., Reumer, 1995; Hernández Fernández, 2001; 86 87 Héran, 2006; van Dam, 2006; Furio et al., 2011), making them considerably more robust proxies for inferring past climates. Furthermore, integrating these taxa is equally essential for generating 88 a broader, more holistic vision of the past ecology of small mammal communities. Quaternary 89 90 sites in the Palaearctic realm frequently yield both human and mammalian fossil remains (e.g., 91 Puzachenko and Markova, 2019) and provide essential evidence for reconstructing the influence 92 of climate changes on human biological and cultural evolution. Developing new palaeoclimatic modelling techniques focused on this realms's rich fossil record potentially offers a means to 93 94 increase both the precision and spatiotemporal resolution of palaeoclimatic inferences. To test 95 these new models, we mobilized data from Balma de l'Abeurador (Languedoc-Roussillon, France) 96 and El Mirón (Cantabria, Spain), two archaeological sequences that cover the period from the Last 97 Glacial Maximum to the beginning of the Holocene.

98

99 2. Material and methods

The Bioclimatic Analysis technique originally described by Hernández Fernández (2001) and Hernández Fernández and Peláez-Campomanes (2003, 2005) consists of two sets of analyses. The first uses linear discriminant functions deduced from the bioclimatic structure of modern mammalian communities in a specific climate zone (defined following Walter's typology, 1970), which are subsequently used to classify additional observations (extinct communities in our case) in each climatic zones. The second is built from transfer functions by means of multiple linear regression analyses of climatic parameters and modern bioclimatic mammalian community
structures. These models are ultimately used to infer climatic variables for additional data (i.e.
extinct communities). These two sets of analyses use a single global dataset of localities for which
the composition of mammalian communities and climatic variables are known.

110

111 *2.1 - Localities, faunal lists, and climate variables*

112 While multiple global bioclimatic classification systems (e.g. Rivas-Martinez, 2008) are 113 available, we employed the climate typology defined by Walter (1970), which is based in the 114 interaction between monthly precipitation and average temperatures. This typology was originally 115 used in the original qualitative Bioclimatic Analysis (Hernández Fernández, 2001), due to its 116 simplicity and correlations with major terrestrial biomes. We collated faunal data from 49 modern 117 localities from across the Palearctic (Table 1, Fig. 1), seven for each of the climate zones present 118 in this biogeographic realm (see Hernández Fernández, 2001 for more details). Localities were 119 selected to be representative of the average climatic conditions of each climate zone and to be as 120 widely distributed as possible (see Table 1 for details). Our sample includes 18 localities from 121 Hernández Fernández (2001), 13 from García Yelo et al. (2014), to which we added an additional 122 18 new localities (Table 1), for a total of 49 data points. Localities in climate zones I (equatorial 123 climate), II (tropical climate with summer rains), and II/III (transitional tropical to semi-arid 124 climate) are not present in the Palaearctic realm and, therefore, contrary to the methodology used 125 by Hernandez Fernandez (2001), were excluded in the development of the new models.

In order to compare results from the two approaches, we built two faunal lists from
published data (Rutilevskiy, 1979; Hernández Fernández, 2001; Héran, 2006; García Yelo et al.,

2014) for each locality: one comprising only rodent species, the other including both rodents and
insectivores. Species taxonomies were revised for each list, integrating the most recent species
distribution data (Wilson et al., 2017; Wilson and Mittermeier, 2018; IUCN red list, 2019).

Nine climatic variables were collected for each locality (Table 1): mean annual temperature
(MAT); annual positive temperature (TP), calculated as the sum of monthly mean temperature for
months with average temperatures above 0 °C; mean temperature of the warmest month (Tmax);
mean temperature of the coldest month (Tmin); mean annual thermal amplitude (MTA), calculated
as the difference between Tmax and Tmin; thermicity index (It), which reflects the intensity of
winter cold (eq .1);

137 It =
$$10*(MAT + 2xTmin)$$
 (eq.1)

138 compensated thermicity index (Itc), which was designed to balance the harsh winter in the
139 continental climates and the extremely mild winter in the oceanic regions in order to make this
140 index comparable worldwide (see Rivas Martínez, 1994); annual total precipitation (P); drought
141 length (D), which is the period when P values are lower than 2*MAT.

142

143 2.2-Calculation of bioclimatic spectra

A climate zone *vs* species presence matrix was constructed for each locality. When a species is absent from a given climate zone, it is encoded as 0. When the species is present in that climate zone, a value of 1/n (referred to as the Climatic Restriction Index, CRI; Hernández Fernández, 2001), where *n* is the number of climates in which the species is present (the division by *n* ensures that the sum of CRI values for a given species equals 1). The more climatically 149 restricted a species is, the higher its CRI values it has. As species with higher CRI imply more 150 specific environmental requirements, they offer more reliable indications of climatic conditions. 151 CRI values for Rodentia and Eulipotyphla species were reexamined (Supplementary material 1) 152 and modified following updated taxonomic and geographical range information (Wilson et al., 153 2017; Wilson and Mittermeier, 2018). Although localities from the three tropical climatic zones 154 were not included, the large range distributions of some small mammal species extend into these 155 zones. Consequently, all ten global climate zones were used to calculate the different Bioclimatic 156 Components (BC) per locality (see below).

For a given locality, the Bioclimatic Component (BC) is the representation of each of the 10 existing climates. All 10 BC values for each locality were calculated according to the formula (eq. 2):

160
$$BC_i = (100 \sum_{j=1}^k CRI_{ij})/S$$
 (eq. 2)

where *i* stands for the *i*th climate zone, *j* for the *j*th species present in the locality, with *S* being the number of species in the locality. The 10 BC values for a given locality constitute its bioclimatic spectrum. The bioclimatic spectrum of each locality was calculated using both datasets: one comprised uniquely of rodents (Supplementary material 2), the other including both Rodentia and Eulipotyphla species (Supplementary material 3).

166

167 *2.3-Palaeoclimatic inferences*

All statistical analyses were performed using the R software package (R core team, 2019).
Linear Discriminant Analysis (LDA - Sneath and Sokal, 1973; Rencher, 2002; Legendre and

170 Legendre, 2012) with the LDA function from the R package MASS (Venables and Ripley, 2002) 171 was used to classify localities into different climate zones. We subsequently used a Leave-One-Out Cross Validation (LOOCV – Efron, 1982; Chernick, 1999) to evaluate classification quality. 172 173 Predictive equations for each of climatic variable were obtained from multiple linear regressions 174 (Sokal and Rohlf, 1995; Zar, 1999; Rencher, 2002; Legendre and Legendre, 2012) using the lm() 175 function, with the ten bioclimatic components as independent variables and the climatic variable 176 as the dependent ones. We generated a script to easily calculate the bioclimatic spectra and inferred 177 climatic zone, which is available as supplementary material (Supplementary material 4 and 178 available at https://github.com/AurelienRoyer/PalBER).

179

180 2.4-Testing sampling biases on palaeoclimatic inference through randomization

181 Small mammal associations from fossil sites potentially present biased representations of 182 past community structure due the nature of the accumulation and subsequent taphonomic 183 alterations (Clark, Beerbower and Kietzke, 1967; Denys, 1985; Andrews, 1990; Kowalski, 1990; 184 Stahl, 1996; Reed, 2007), which may, in turn, impact climatic inferences drawn from Bioclimatic 185 Analysis. For instance, many fossil assemblages of small mammals are accumulated by predators, 186 especially nocturnal raptors, who are relatively selective in their choice of prey (e.g., Andrews, 187 1990). Bioclimatic Analysis has the advantage of dampening this potential bias by taking into 188 account the general representativeness of the taxonomic composition of micromammal death 189 assemblages (i.e. the presence or absence of particular taxa) rather than their relative abundance. 190 This method does not differentially weight taxa, and the occurrence of a single fossil of a unique 191 species could theoretically alter a locality's bioclimatic spectrum and, therefore, its climate

192 reconstruction. However, although high levels of ecological redundancy in mammalian 193 communities (Gómez Cano et al., 2006) somewhat minimize this problem, it can be better 194 overcome with adequate sample size. Although the fragile nature of small mammal fossils means 195 they are unlikely to be redistributed (Álvarez-Sierra et al., 1990, Fernández-López, 2000), potential 196 species loss associated with different processes of accumulation, fossilization, and sampling bias 197 cannot be ruled out (Andrews, 1990 Behrensmeyer, 1991; Lyman, 1994; Andrews, 2006; Gómez 198 Cano et al., 2006; Turvey and Cooper, 2009; Fernández-Jalvo, Scott and Andrews, 2011), raising 199 questions as to how these potential biases affect Bioclimatic Analysis (Gómez Cano et al., 2006).

200 To investigate the potential impact of species loss, we randomly removed species in the 201 modern localities in order to replicate impoverished faunal associations. Multiple LDA and 202 multiple linear regressions were performed for each locality by randomly removing k taxa (k 203 varying from 1 to *n*-2, with *n* being number of species in the locality). For a given locality and a 204 given k, 100 impoverished samples were generated. A total of 69,000 randomized samples were 205 produced, for which simulated climatic inferences were calculated using the new rodents and 206 rodents plus insectivores models. Three locations in the arctic climate zone (IX) were left 207 unchanged, as they contain less than three species. We used logistic regressions (Sokal and Rohlf, 208 1995; Legendre and Legendre, 2012) to explore the coherence of simulated climatic inferences for 209 each climate zone. Quantitative inferences of climatic variables were evaluated using standard 210 deviations of predicted values calculated for a given locality and a given k taxa removed. 211 ANCOVAs (Zar, 1999; Sokal and Rohlf, 1995) were performed for each climate parameter using 212 log-transformed standard deviations of predicted values, with the percentage of taxon removed 213 and climate zones used as independent variables.

214 2.5-Application of the new models to palaeoclimatic reconstruction

The two new approaches for Bioclimatic Analysis allow us to analyze small mammal associations from the Palaearctic Quaternary and reconstruct climatic conditions. We used small mammal associations from La Balma de l'Abeurador (Languedoc-Roussillon, Hérault, France) and El Mirón (Cantabria, Spain), two sites located in typical temperate climate areas and that yielded sequences spanning from the end of the Late Pleistocene to the beginning of the Holocene (Fig. 2). All radiocarbon dates were calibrated at 2σ (95%) using Oxcal v4.3 (Bronk Ramsey, 2017) and the IntCal13 calibration curve (Reimer et al., 2013).

222 The site of Balma de l'Abeurador is located in southern France and currently has a mean 223 annual temperature of 13.6°C and annual precipitation around 700 mm. The central sector of the 224 cave produced six multi-level deposits (Heinz et al., 1992; Vaquer et al., 1986; Vaquer and Ruas, 225 2009), whose chronology was determined by the nature of the lithic assemblage, palynology, and 226 radiocarbon dates (Vaquer and Ruas, 2009). Small mammals from Balma de l'Abeurador were 227 first studied by Marquet (1987) and then by Mistrot (2001), who identified at least 19 taxa in an 228 assemblage comprising at least 2000 individuals (Table 2). We used Mistrot's (2001) data, as it 229 benefits from a taphonomical analysis demonstrating no bias in the representativity of the small 230 mammal assemblage. This detailed taphonomical approach, together with the high number of 231 remains (more than 100 first and second upper and lower rodent molars for each level) and an 232 adequate minimum fossil sample size of fossils, allows for reliable palaeoclimatic inferences (van 233 de Weerd and Daams, 1978; Daams et al., 1999). Several differences do exist between Mistrot's 234 (2001) and Marquet's (1987) analyses in terms of species identification, with the latter employing 235 extreme caution when identifying fossils to species. For Apodemus, Arvicola, and Microtus 236 (Terricola), Mistrot (2001) made no specific identifications, whereas Marquet (1987) and Brunet-237 Lecomte (1988) identified almost exclusively Ap. sylvaticus, Ar. Amphibius, and M. (T.)

duodecimcostatus throughout the sequence. For the insectivore genus Sorex, Neomys, and
 Crocidura, no distinction was made between Sorex araneus and Sorex coronatus, between Neomys
 anomalus and Neomys fodiens, or between Crocidura leucodon and Crocidura russula.

241

Table 2: Faunal lists of identified Rodentia and Eulipotyphla per level at Balma de l'Abeurador.
Crosses indicate species present, circles represent absence species. Data are from Mistrot (2001).

244

245 The site of El Mirón is located in northern Spain, and enjoys relatively similar climate 246 conditions to Balma de l'Abeurador, but with slightly higher annual precipitation (reaching 1,000 247 mm) and smaller seasonal differences in rainfall (Fig. 2). The site yielded a long stratigraphic 248 sequence (Strauss and Morales, 2019) and numerous radiocarbon dates (e.g., Strauss and Morales, 249 2003, 2007, 2012, 2019). Twenty-one insectivore and rodent taxa were identified from throughout 250 the 44 studied levels of the stratigraphy (Cuenca-Bescós et al., 2008, 2009). Although over 3,000 251 individuals were identified, they are highly unevenly distributed among the different levels. To 252 ensure the reliability of palaeoclimatic inferences, we selected only deposits with more than 50 253 individuals, or grouped together several successive levels (i.e. the Solutrean levels 120 to 126 from 254 the Last Glacial Maximum). For Apodemus and Sorex, no distinctions were made between Ap. sylvaticus and Ap. flavicollis or between S. araneus and S. coronatus. 255

When reliable criteria for confidently distinguishing between species are unavailable, two approaches can be taken: either building bioclimatic spectra from all the potentially present species and calculating a palaeoclimate inferences for each; or using an average species, sometimes referred to as "quimeric" species. When a taxon is only identified to genus, the biome occupation of all the potential species can be used. For instance, if one species inhabits biomes VI and VII, and the other inhabits IV and VI; the genus can be considered to inhabit IV, VI, and VII. Even if relatively little uncertainty is introduced, "quimeric" species have a lower discriminatory power than any of the potential original species while still providing useful information for community structure. Consequently, both approaches were used and their results compared.

Extinct species potentially present an additional limitation, as no past climate data is available for their geographic distribution. Following Hernández Fernández et al. (2007), we assigned extinct taxa to different BCs according to the distribution of their extant relatives. As "quimeric" species, they provide reduced discriminatory power. The El Mirón sequence has the particularity of yielding remains of the now-extinct rodent *Pliomys coronensis* (*=Pliomys lenki*), for which we used the bioclimate characterization proposed by Hernández Fernández et al. (2007).

271

272 **3-Results and discussion**

273 *3.1-Qualitative climatic classification and quantitative climatic inferences*

The number of species in modern localities from the different climate zones ranges on average between 17 and 24, except for bioclimate zones III and IX (Supplementary material 6). Localities in these two climate zones yielded a maximum of 15 species, with some containing less than 5 species. Eulipotyphla produced around 25 % of all faunal lists on average, but faunal lists from bioclimate zones IV, VI and VIII included 30 % or more of Eulipotyphla.

279 The two linear discriminant analyses based on the bioclimatic spectra correctly classified 280 nearly all localities, inducing a very low rate of error. When only Rodentia was used, incorrectly 281 classified localities rose to 6 % (Table 3), a similar figure to the one calculated by Hernández 282 Fernández (2001). When the combined Rodentia and Eulipotyphia dataset is used, only 283 Blagoveshchensk was incorrectly classified in climate zone VIII instead of VI. However, the cool-284 temperate and boreal forests throughout the region of Blagoveshchensk (Kolbek, Srutek and Box, 285 2013) are home to numerous species typical of colder climates, including Lemmus amurensis, 286 Clethrionomys rufocanus, Myopus schisticolor, Pteromys volans, or Sorex tundrensis), which 287 likely goes some way in explaining the incorrect climate zone attribution.

Table 3: Apparent error rate, prediction error, and prediction quality (proportion of observations
with a posterior probability > 0.9) for each approach (Rodentia, Rodentia plus Eulipotyphla) for
LDA. The prediction error is based on LOOCV.

291

The Leave-One-Out Cross Validation (LOOCV) indicated very low prediction error; the rodent-only dataset produced approximately 12 % misclassifications, while the dataset including both sets of mammals had both a lower level of error (Table 3) and predictions associated with a high posterior probability (Table 3). This would indicate climate inferences based on the combination of rodents and insectivores to be slightly more reliable. The high predictive power of both models reveals a close relationship between small mammal communities and climate, which makes them suitable to infer past climate conditions (Hernández Fernández, 2001).

299 Most predictive equations generated by the multiple linear regressions for each climatic 300 factor, as a function of the bioclimatic components of either Rodentia or Rodentia and

301 Eulipotyphla, produced highly significant determination coefficients (Table 4). The different 302 transfer functions for each climatic variable had similar coefficients of determination for both datasets, with minor trends towards slightly higher values for the combined Rodentia and 303 304 Eulipotyphla dataset. Values were also relatively similar to those obtained by Hernández 305 Fernández and Peláez-Campomanes (2005). Only MTA showed very low coefficients of 306 determination (<0.6). Precipitation shows a slightly better coefficient of determination than the 307 one obtained by Hernández Fernández and Peláez-Campomanes (2005), although it is still lower 308 than 0.8 (Table 4), which resulted in calculated prediction with large intervals.

309

310 *3.2-Loss of species and bias in climate inferences*

311 Overall, few incorrect qualitative inferences were generated when species were removed 312 from the faunal lists (usually below 15 % per climate zone – fig. 3, supplementary material 7). The 313 S-shaped curves of logistic regressions are skewed upwards (Fig. 3; Supplementary material 7), 314 meaning there is no "turning point" when climatic inferences become erroneous and that reliable 315 classifications can nevertheless be obtained using relatively few species, although they may have 316 a high probability of error (Fig. 4). The two approaches behaved differently according to climate 317 zone when species are randomly removed; inferences drawn from rodent communities are more 318 robust for climate zones III, V, and IX, while the second approach (Rodentia plus Eulipotyphla) is 319 more robust for climate zones IV, VI, and VIII. Incorrect inferences for communities in the polar 320 climate zone (IX) are low with the rodent-only dataset (4.2%) and much higher with the Rodentia 321 plus Eulipotyphla, with incorrect inferences mainly classified into the boreal climate zone (VIII). 322 The percentage of incorrect classifications increased for all climate zones when more taxa are

323 removed (Figs. 3 and 4, supplementary material 7). Although classification error generally 324 increased after the removal of 30-40 % of taxa from the localities (Fig. 4), most reached the 25 % 325 incorrect classifications threshold when more than 50 % of taxa were removed. Nevertheless, the 326 warm temperate (climate zone V; except for the Batumi locality) and arid temperate climate (VII) 327 zones appeared to be much more resistant to species loss independent of the dataset used (Fig. 3, 328 supplementary material 7). Substantial variability is evident between localities in the same climate 329 zone, demonstrating faunal communities from the same climate zone to be highly variable. This 330 variability reflects varying responses of different taxa inhabiting the same climate zone.

331 Quantification of climatic parameters follows similar trends. Differences between 332 theoretical and calculated values in impoverished communities for each climatic parameter are highly dependent on the percentage of remaining taxa (Supplementary material 8). These 333 334 differences vary as a function of the climate zone considered and the dataset used. For instance, in 335 the case of precipitation estimates from climate zones III and VIII (subtropical arid and boreal), 336 less impoverished communities produced larger differences when the rodent-only dataset was 337 used. Figures 5 and 6 show standard deviations of predicted values obtained for impoverished 338 faunal associations computed for each climatic parameters. These standard deviations follow 339 exponential variations, which increase with the percentage of taxa removed. Overall, standard 340 deviations of predicted values become important when half of the faunal list is removed (Fig. 5 341 and 6). ANCOVAs for each climate parameter show no significant difference between climate 342 zones (F = (6, 676), p > 0.7). The approach based on rodent communities generated steeper 343 regression curves than those based on Rodentia plus Eulipotyphia, which suggests the former to 344 be more sensitive to species loss. This is probably related to the fact that faunal lists comprised 345 exclusively of rodents contain less species than those combining both orders of small mammals.

When confronted with species loss, the combined dataset therefore appears better adapted toproducing robust climate quantifications.

As expected, our results indicate that palaeoclimatic inferences obtained by Bioclimatic Analysis depend significantly on the integrity of the past faunal communities used. Satisfactory levels of confidence can be obtained only when species loss is below 50 %, which is similar to the initial estimates produced by Gómez Cano et al. (2006). When quantifying climatic parameters, the approach combining Rodentia and Eulipotyphla proved less sensitive to faunal impoverishment, and should be preferred when species loss is suspected.

354

355 4-Palaeoclimatic inferences from past small mammal communities

356 Climate zone classifications and mean annual temperature predictions generated by the two 357 approaches were applied to multiple faunal lists from El Mirón and Abeurador (Figure 7). All 358 possibilities related to uncertainties in species identifications (see above) were tested. Climate 359 prediction remained the same for both datasets, except for the case of Abeurador level 5b, for 360 which two different climate zones (IV and VI) were predicted depending on the Eulipotyphla 361 species used. When uncertainties concern only two species, variations in mean annual temperature 362 values remain minimal (Fig. 7, Supplementary material 5). For instance, temperature variation was 363 less than 0.5°C between communities in which Apodemus specimens were identified as A. 364 sylvaticus instead of A. flavicollis, and vice versa. In the case of N. anomalus and N. fodiens, 365 variation reached almost 2°C. Observed variations can reach up to 4°C (e.g. Abeurador level 8) 366 when all possibilities are run using either of the two approaches. The highest degree of variation 367 primarily concerns uncertain species inhabiting only a few climate zone that are different between 368 them, such as *S. araneus* and *S. coronatus*. As a consequence, faunal lists employed in bioclimatic 369 modelling should only include reliably identified species. Finally, small variations could also be 370 observed according to the approach used, the one based on rodents tending to estimate mean annual 371 temperature two degrees lower.

372 Figure 8 depicts the mean annual temperature obtained from small mammal lists of 373 Abeurador and El Mirón. For levels with uncertainties in species classification, we used 374 "quimeric" species for climatic inferences. Overall, the two approaches show similar trends in both 375 sequences; an attribution to climate zone IV during the Holocene and a progressive warming with the onset of the Bølling interstadial. While the same trend can be observed for predictions based 376 377 on Rodentia and Eulipotyphla and only Rodentia, the latter tended to produce lower mean annual 378 temperature estimates. Mean annual temperatures obtained for the Abeurador sequence using the 379 rodent dataset ranges from 3.8°C to 9.2°C, whereas the approach based on Rodentia and 380 Eulipotyphla ranges from 6.3 to 11.6°C. Similarly, mean annual temperature varied throughout the 381 El Mirón sequence, averaging around 1.2°C for both approaches, which is included in the 382 confidence intervals (and so in the prediction intervals). The lowest mean annual estimations were 383 for the LGM levels 112-114. Temperatures rose around 3.0°C between Heinrich stadial 1 and the 384 beginning of the Holocene at El Mirón, which is slightly lower than was calculated for Abeurador. 385 No major differences are observed between the two approaches for precipitation and thermal index 386 conditions (Fig. 9, Supplementary material 5). Throughout the Abeurador sequence, the 387 Eulipotyphla approach showed an increase of precipitation between Heinrich Stadial 1 and the 388 Boreal interstadial, suggesting wetter climate conditions to have prevailed during the Holocene at 389 Abeurador. Nevertheless, no differences were inferred with the rodent approach, suggesting that 390 if changes in precipitation did occur, they were not important enough to have a significant impact

391 on rodent communities. A progressive increase in the Thermicity index can be observed 392 throughout the sequence, confirming milder conditions associated with the Holocene deposits. 393 Apart from a slight decrease at the end of the sequence, no abrupt changes in precipitation were 394 recorded anywhere in the El Mirón record, and no differences are evident between predictions for 395 both approaches. Several variations in the Thermicity index are evident across the sequence, with 396 several notable decreases in levels 112-114 and 107-106, whereas a slight increasing trends 397 emerges at the end of the Late Glacial. These variations were recorded for a level where low mean annual temperature were predicted, suggesting these changes were associated with modifications 398 399 in seasonality and the intensity of winter colds. Climate parameters from levels 110 and 111 of El 400 Mirón were previously explored using the Mutual Climatic Range of the rodents (Bañuls-Cardona 401 et al., 2014). Mean annual temperatures were estimated to be 3 to 4 °C higher, averaging around 402 10.6°C, whereas the mean temperature of the coldest months diverge by 6°C. The Bioclimatic 403 approach suggests colder climatic conditions for the LGM levels of El Mirón.

404 Our analysis of the Balma de l'Abeurador and El Mirón sequences primarily focused on 405 three climate variables (mean annual temperature, precipitation, and thermicity index) considered 406 susceptible to changes during this time period. These two terrestrial sequences are in relative 407 agreement with evidence from regional climate evidence recorded in speleothems as well as lake 408 and marine core records (Jalut and Turu, 2009; Moreno et al., 2012). The LGM appears as a cold, 409 relatively dry period (but not as dry as Heinrich Stadial 1), inducing a hiatus in speleothem and 410 lake records. Bioclimatic models built from LGM deposits tend to estimate lower mean annual 411 temperature and a lower Thermicity index, as is the case in El Mirón level 112.114. With that said, 412 our models revealed no major change in precipitation coincident with the LGM, with all variations 413 falling within the large confidence intervals. Speleothems from northern Spain and southern 414 France (Chauvet Cave) have suggested an increase in temperature and humidity just before the 415 Bølling interstadial (15.4 kyr), which reached their maximum during the Allerød (Genty et al., 416 2006; Moreno et al., 2010; Martínez Pillado et al., 2014). Mean annual temperature based on small 417 mammals closely mirror the mean annual temperature model built from speleothems. The 418 Younger-Dryas at El Mirón is represented by two levels with radiocarbon dates compatible with 419 this period. Overall, speleothem and pollen records suggest the Younger-Dryas to be characterized 420 by an abrupt cold event associated with a reduction in arboreal communities (Fletcher and Sánchez 421 Goñi, 2008; Morellón et al., 2018). There is currently little evidence supporting this event to 422 comprise two phases; an initial dry phase followed by a progressive increase in humidity and 423 warmer conditions (e.g. Muñoz-Sobrino et al., 2013; Bartolomé et al., 2015; Morellón et al., 2018). 424 Two hypotheses could explain the pattern observed for the Younger-Dryas at El Mirón; low 425 archaeological resolution or the relatively insignificant impact of this event on small mammal 426 communities.

427 Despite being located on different sides of the Pyrenees, these two sequences are relatively
428 close geographically and today lie in typical temperate climates. Our results suggest that warming
429 during the Holocene differed slightly of either side of this natural barrier, being less marked in
430 northern Spain compared to southern France.

431

432 5. Conclusions

By focusing on Palaearctic small mammal communities, our work refines the Bioclimatic
Analysis first developed by Hernández Fernández two decades ago and extends its use to rich
fossil record of the Eurasian Quaternary. Although rodent communities alone generate informative

436 qualitative and quantitative climatic reconstructions, combining rodents and insectivores increases 437 precision for several climatic parameters, particularly precipitation. As previously indicated by 438 Hernández Fernández and Peláez-Campomanes (2005), due to the statistical uncertainty present 439 in every climate transference function, the inferences drawn should not be taken as absolutes but 440 as indicators of general trends. Although it is robust against sampling bias, this method 441 nevertheless requires data from secure contexts with proper sampling efforts and adequate 442 taphonomical control in order to ensure accurate taxonomic identification and the representativity 443 of faunal assemblages. Although imprecisions in taxonomic identification do not impact general 444 trends, they could engender variations in climatic estimations, which highlights the importance of 445 confidence intervals. The R script developed for this analysis (and available as supplementary 446 material 4) provides a simple means for using Bioclimatic Analysis for past climate 447 reconstructions.

The application of these new models to the Quaternary sequences of Balma de l'Abeurador and El Mirón to reconstruct paleoclimate revealed relatively similar changes with those built from other proxies, such as speleothems. The bioclimatic inferences from these two archaeological sequences suggest that, depending on the side of Pyrenees, mean annual temperatures warmed from 3 to 5°C between the Late Glacial and the Holocene.

453

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- 811
- 812 Figure captions:

813 Figure 1: Maps of the 10 climate zones and 49 selected localities. See Table 1 for the list of814 localities.

Figure 2: A) location of Abeurador and El Mirón, B) Walter and Lieth climate diagrams
(calculated between 1985 and 2015) and, C) available radiocarbon dates for levels from both
sequence with associated cultural attributions. Radiocarbon dates calibrated using OxCal v4.3 and
IntCal13.

819 Figure 3: Fitted curves (red) for logistic regressions of the correctness of the LDA classification 820 for each climatic zone (vertical axes: above, correct predictions, below, incorrect predictions) 821 against the percentage of remaining taxon (horizontal axes). This percentage was calculated from 822 the randomization procedure of 49 rodent and Eulipotyphla faunal associations (seven for each 823 climate zone), which were impoverished by randomly removing species. Histograms illustrate 824 frequencies of classification as a function of the percentage of species remaining in the association 825 (number of impoverished localities is shown on the right axis). Boxplots illustrate median, quartile, 826 and range of incorrect inferences for all the randomly impoverished communities in each climate 827 zone. A detailed indication of the percentage for each climate zone estimated by incorrect 828 classifications is provided in the middle of each graph (numbers as in Table 1).

Figure 4: Evolution (colored curves) of the percentage of incorrectly classified localities (vertical axes) depending on the percentages of remaining taxa (horizontal axes) obtained with randomly impoverished communities for each locality (numbers as in Table 1) and per climate zone.
Numbers of localities are reported in table 1. Blagoveshchensk was excluded from climate zone IV. Three localities (44, 45 and 46) were not randomly impoverished for climate zone IX due to very low species numbers.

Figure 5: Fitted curves (black) and associated confidence intervals (blue) for the exponential regressions of the standard deviations of the predicted values against the percentage of remaining taxon from impoverished faunal list for the approach based on Rodentia. Point color codes for the climate zones are the same as in Fig. 1

Figure 6: Fitted curves (black) for each climatic parameter and associated confidence intervals (in
blue) for the exponential regressions of the standard deviations of predicted values against the
percentage of remaining taxon from the impoverished faunal list for Rodentia and Rodentia plus
Eulipotyphla. Point colors codes for the climate zones are the same as in Fig. 1.

Figure 7: Influence of the species identification uncertainty on Mean Annual Temperature
estimates for deposits from El Mirón and Abeurador (squares indicated rodents and circles and
rodents plus Eulipotyphla). The symbol colors represent inferred climate zone. Confidence
intervals for the estimated MAT values shown as vertical lines.

Figure 8: Evolution of the estimated Mean Annual Temperature along the El Mirón (bottom) and
Abeurador (top) chronozone by means of multiple regression. Squares represent estimated MAT
values obtained from the Rodentia approach, circles for the Rodentia plus Eulipotyphla approach.
Symbol colors indicate inferred climatic zone. Vertical lines represent confidence intervals for
estimated MAT values. Horizontal lines delimit levels associated with chronozones.

Figure 9: Evolution of estimated precipitation and thermicity index (TI) along the El Mirón (bottom) and Abeurador (top) chronozone by means of multiple regression. Squares represent estimated MAT values from the Rodentia approach, circles for the Rodentia plus Eulipotyphla approach. Symbol colors indicate inferred climatic zone. Vertical lines represent confidence intervals for estimated MAT values. Horizontal lines delimit levels associated with chronozones. 857

858 Table captions:

Table 1: Coordinates, climate zone, and climatic parameters values for the 49 localities. New localities added for this study are indicated with asterisks. Abbreviations: MAT = mean annual temperature; TP = annual positive temperature; Tmax = mean temperature of the warmest month; Tmin = mean temperature of the coldest month; MTA = mean annual thermal amplitude; It = thermicity index; Itc = compensated thermicity index; P = annual total precipitation; D= drought length.

Table 2: Faunal lists of identified Rodentia and Eulipotyphla per level at Balma de l'Abeurador.Crosses stand for species presence, circles for species absence. Data are from Mistrot (2001).

867 Table 3: Apparent error rate, prediction error, and prediction quality (proportion of observations
868 with a posterior probability > 0.9) for each approach (Rodentia, Rodentia plus Eulipotyphla) for
869 LDA. The prediction error is based on LOOCV.

870 Table 4: Coefficient of determination values obtained for multiple linear regressions for each871 climatic variable according to the two approaches (Rodentia, Rodentia plus Eulipotyphla).

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875 Supplementary captions:

Supplementary material 1: Climatic restriction index of Rodentia and Eulipotyphla. The last
column provides details for each locality (numbers as in Table 1), from which these taxa were
found.

879 Supplementary material 2: Bioclimatic spectra of Rodentia for each locality.

880 Supplementary material 3: Bioclimatic spectra of Rodentia and Eulipotyphia for each locality.

881 Supplementary material 4: Palber.R Script applied to bioclimatic method and bioclimatic spectra882 files (.csv) necessary to run it.

883 Supplementary material 5: Predictions calculated for each level of Abeurador and El Mirón,884 obtained from the two approaches.

885 Supplementary material 6: Number of rodents and Eulipotyphla per modern locality.

886 Supplementary material 7: For each climatic zone, S-shape curves (red) for logistic regressions of 887 the correctness of the LDA classification for each climate zone (vertical axes: above, correct 888 predictions, below, wrong predictions) against the percentage of the remaining taxon (horizontal 889 axes). This percentage was calculated from the randomization procedure of 49 rodent faunal 890 associations (seven for each climate zone), which were impoverished by randomly removing 891 species. Histograms illustrate frequencies of classification as a function of the percentage of 892 species remaining in the association (number of impoverished localities is shown on the right 893 axes). Boxplots illustrate median, quartile, and range of incorrect inferences for all the randomly 894 impoverished communities in each climate zone. A detailed of incorrect classifications (%) are 895 provided in the middle of each graph (numbers as in Table 1), with the average details of the 896 climate zones into which they are classified.

Supplementary material 8: Evolution of differences between estimated climate values by means of
multiple regression for each climate variable and both approaches (Rodentia or Rodentia +
Eulipotyphla) according to the percentage of remaining taxa and values calculated when the faunal
list is complete. Each figure corresponds to a combination of a given climate variable and a given
approach.