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3	Pan-Mediterranean Holocene vegetation
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26 ABSTRACT

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28 Aim

29 The Mediterranean is characterised by diverse and spatially-heterogeneous mosaic 30 landscapes. Within this study a cluster analysis-based method is developed for the 31 classification of Mediterranean vegetation types based on modern and fossil pollen datasets. 32 The application of this approach to multiple pollen records spanning the Mediterranean 33 region has allowed temporal variations in vegetation dynamics to be explored throughout the 34 Holocene. We ask how far back stable baseline vegetation communities can be identified in the pollen record, and whether those types considered to be characteristic of the 35 36 Mediterranean landscapes have been present in the past as well as at the present. 37 Location 38 The research location includes the land areas surrounding the Mediterranean Sea. The pollen 39 sites are principally located in mainland Spain, southern France, Greece and Turkey, Italy, North Africa, the Levant, and some Mediterranean islands. 40 41 Methods 42 5641 samples from 158 fossil pollen records (cores) and 1799 modern pollen surface samples 43 were harmonised taxonomically and pollen count data summed into 200-year time windows 44 on a common timescale from 11,000 BP to the present-day. Cluster analysis and community 45 classification was used to identify major vegetation types along with other approaches to explore patterns in ecological datasets, such as Simpson's diversity index and non-metric 46 47 multidimensional scaling. 48 **Results**

The pollen datasets were classified into eleven closed forest/woodland and five open or
scrubland vegetation types. Closed vegetation clusters declined from the mid-Holocene with

a marked increase in open or human-modified vegetation types since 3500 BP and with an
increasing rate of vegetation change and habitat diversity through time.

53 Main conclusions

54 The Mediterranean has been a dynamic landscape throughout the Holocene with frequent

55 changes in land cover identified in the pollen datasets. The pollen-inferred clusters reveal a

56 wider range of Mediterranean vegetation types than identified in previous studies; for

57 example, including both beech and alder woods. Evergreen Oleaceae-dominated shrubland is

58 much better represented in modern than in fossil samples while mesic forest was abundant in

59 the past but is uncommon today.

60

61 **Keywords:** Disturbance; Holocene; Human impacts; Land-cover; Landscape structure;

62 Mediterranean; Pollen; Vegetation

63

65 INTRODUCTION

66

67 Pollen analysis offers a valuable tool for exploring long-term changes in vegetation and land cover (Jackson & Blois 2014; Edwards et al., 2017). Numerous studies have mapped and 68 69 classified European vegetation using fossil pollen for discrete time intervals through the 70 Holocene in relation to climate and major environmental gradients (e.g. Huntley, 1990; 71 Prentice et al., 1996; Peyron et al., 1998). This has led to the description of plant functional 72 types, the identification of 'biomes' and their spatial and temporal distribution (e.g. Marinova 73 et al., 2017), climate reconstruction based on pollen and vegetation modelling (e.g. Guiot & Kaniewski, 2015) and geographical mapping of individual taxa (Brewer et al., 2017). 74 75 Understanding patterns of vegetation dynamics and biogeographic change allows improved 76 knowledge of the habitats within ecoregions and their responses to disturbance and 77 environmental change, which can provide frameworks for biodiversity conservation policies 78 and ecological modelling (Blasi et al., 2014). Previous studies have often focused on 79 continental scales. For example the 'BIOME 6000' mapping project (Prentice et al., 1996; 80 Prentice & Webb, 1998) provided a pollen-based landscape reconstruction for Europe for two 81 time periods (6000 and 0 BP) based on modern and fossil data, notably for climate-biosphere modelling. However, continental-scale descriptions of vegetation types can be less 82 83 informative about ecologically diverse sub-regions, such as the Mediterranean. There is 84 potential for more detailed classification and reconstruction of Mediterranean ecoregions (e.g. 85 Collins et al., 2012), which can allow exploration of how vegetation types developed in 86 relation to human land modification. 87

The present vegetation of the Mediterranean region is the product of a distinctive climate and
natural environment transformed by human activities over many millennia (Jalut et al., 2009).

90 Improved understanding of the unique vegetation assemblages that characterise the 91 Mediterranean landscape and how these have changed over millennia has potential value in 92 identifying vegetation types at risk from future land use change. Much of the modern 93 Mediterranean vegetation has been altered by deforestation, over-grazing, erosion and 94 burning (Vogiatzakis et al., 2006). In recent decades these factors have led to a shift from 95 highly heterogeneous landscapes maintained by multiple different land uses, to more 96 homogeneous landscapes, for example, following land abandonment on some European shores of the Mediterranean (Pausas, 1999). Wildfires are also an important feature of 97 98 Mediterranean ecosystems and their associated disturbance regimes (Rundel et al., 1998). 99 Animal grazing has been a constant disturbance factor throughout the Holocene, and has been 100 important in maintaining open and mosaic landscapes (Blondel, 2006). Plant distribution is 101 also restricted by environmental factors such as species' climatic tolerances and interspecific 102 interactions (e.g. García-Valdés et al., 2015). Climate is an important driver of vegetation 103 turnover through previous interglacial periods (e.g. Langgut et al., 2011) and some studies 104 employ pollen analysis as a tool to reconstruct past climate (e.g. Mauri et al., 2015). Since the 105 mid-Holocene, the increasing impact of human land use has been demonstrated in numerous 106 studies (e.g Colombaroli & Tinner, 2013). Consequently, disentangling the impacts of 107 climate and human land use on vegetation shifts can be complex, notably for recent 108 millennia.

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The aims of this study are to: 1) develop a data-led method for the classification of vegetation types based on modern and fossil pollen datasets across the Mediterranean region; 2) identify temporal and spatial variations in the abundance of different vegetation types and explore whether stable baseline vegetation communities can be identified in the pollen record; and 3) compare these results with previous studies that have classified and mapped vegetation from pollen data and modern vegetation associations. These aims will be achieved via use of
cluster analysis to derive a numerical classification of vegetation types from pollen taxa
associations. This approach has the potential to be applied to multiple sequences covering
broad geographic regions and is flexible enough to accommodate additional datasets in the
future. Databases of modern pollen samples (Davis et al., 2013) and fossil pollen (e.g. Fyfe et
al., 2009) have been used for this macro-ecological study.

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122 METHODS

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124 Modern and fossil pollen datasets

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126 Pollen count data have been obtained from the European Modern Pollen Database (Davis et 127 al., 2013) and the European Pollen Database for fossil records (version: May 2016) (Leydet, 128 2007-2017). The analyses are based on 1799 modern surface samples and 158 fossil pollen 129 records (cores) from 143 sites (Supplementary Information 1). Only pollen sequences with reliable chronologies (sensu Giesecke et al. 2013) were used. These provide more 130 independent age control for the mapping of vegetation change than has been possible in 131 previous studies. The pollen count data from each site have been summed into 200-year time 132 133 windows from 11000 BP to present. These allow vegetation changes of sub-millennial 134 duration to be identified, avoid 'smoothing' that might result from using longer time windows (Fyfe et al., 2015) and enable direct temporal comparison among sites. The modern (i.e. 135 surface) pollen and most recent (200 BP - present) fossil samples are separated in the 136 137 analyses. The majority of the modern pollen samples derive from moss polsters or soils and 138 there is no duplication of core top samples between the modern and fossil datasets. This has resulted in 3852 fossil samples distributed across 55 time windows, and a total of 5641 139

modern/fossil pollen samples. Pollen sites located in areas with characteristic Mediterranean
climate and vegetation have been selected for analysis (Fig. 1), as defined using the
vegetation zones illustrated by Izdebski et al. (2015) and Ozenda & Borela (2000). Pollen
sites from the Rhone Valley in southern France were also included in the initial cluster
analysis to identify land-cover types at higher elevation, but these sites were subsequently
removed for analysis of patterns in the Mediterranean, leading to 105 fossil records and 1610
modern samples.

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148 The taxonomy of the modern and fossil datasets has been harmonised and simplified using 149 the EPD nomenclature (Leydet, 2007-2017). Different levels of taxonomic resolution were 150 chosen for specific purposes. For example, key indicator species; i.e. taxa that are associated 151 with certain land use types, such as arable weeds, were grouped at the species level (e.g. 152 ribwort plantain (*Plantago lanceolata*)), whereas other taxa were combined at a higher 153 taxonomic level, such as all evergreen oak species and varieties (e.g. Quercus ilex and 154 *Quercus suber*). There are challenges for pollen analysts in subdividing the Quercus group 155 based on the morphology of the pollen grains. For example, *Quercus cerris/suber* types, 156 which represent both evergreen and deciduous types, are often combined by analysts due to difficulties in distinguishing them using a light microscope. Certain assumptions are required 157 158 when utilising large databases that have been compiled from the work of numerous different 159 analysts, for example, 'Quercus undifferentiated' could represent evergreen or deciduous 160 *Quercus* when not identified to species level. Within the dataset presented here, deciduous Quercus appears in 150 sequences and evergreen Quercus has been identified in 125 161 162 sequences, therefore in the majority of cases analysts have separated different oak types. Only taxa with >1% occurrence that occur in >50 samples were included, resulting in 260 163 164 pollen taxa. The geographical coverage of this dataset is good for most of the European

Mediterranean but poorer for the African side. This needs to be taken into account when interpreting results, as all phytogeographical conditions may not be represented. The terms 'vegetation clusters' and 'vegetation/pollen assemblages' are used interchangeably and 'cluster group' is used to describe the resulting collection of pollen samples that fall into a particular 'cluster' number.

170

171 Data analysis

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173 Ward's hierarchical agglomerative clustering method (Ward, 1963; Murtagh & Legendre, 174 2014) has been applied to the amalgamated modern and fossil pollen dataset using the 'Rioja' 175 R package (Juggins, 2015). This unsupervised data-driven approach was used to assign pollen 176 samples to cluster groups based on the similarity of their taxa assemblages without excluding 177 any land pollen taxa from the dataset (other than those <1% with fewer than 50 occurrences). 178 Whilst many approaches exist for classification of multivariate data, these generally produce 179 similar results in pollen datasets (Felde et al., 2014; Felde et al., 2016). Cluster analysis 180 allows hierarchical aggregation at different classification levels (García-Madrid et al., 2014). 181 Detailed comparison of the pollen samples and assemblages at different clustering levels has allowed exploration of the cluster groups that separate at a lower clustering level (e.g. six 182 183 clusters) and are represented by more distinct vegetation groups (Table 1). Unsupervised 184 cluster analysis assigns pollen samples into a predetermined number of groups. The ratio of change between cluster groups through time, which may represent a measure of community 185 turnover, was calculated by identifying the number of pollen samples that switch to a new 186 187 cluster or remain within the same cluster between successive time windows and transforming these values into a ratio of change to non-change. 188

190 After assigning pollen samples to clusters statistically, a phytosociological classification 191 approach was used to identify the frequent and abundant taxa within each group based on the 192 number of occurrences of the taxon, the average percentage, median and interquartile range 193 (IQR). The resulting community tables were used to facilitate the selection of interpretive 194 name descriptors for each cluster based on the most frequent/abundant taxa along with 195 comparisons with other classification systems (Küchler & Zonneveld, 1998; Bossard et al., 196 2000; Allen, 2009). The pollen cluster groups have also been compared against land cover 197 and land use types defined by the Corine (COoRdination of INformation on the Environment) 198 remotely-sensed land cover maps (European Environment Agency (EEA), 2016) and the 199 results of previous studies (Huntley, 1990; Prentice et al., 1996; Guliot et al., 1996; Peyron et 200 al., 1998). An individual taxon's frequency is determined by calculating its number of 201 occurrences divided by the number of samples in the cluster and assigning one of five 202 frequency classes based on cut-off values between each group, which follows the method 203 used by Perez et al. (2015). If a taxon appears in 81-100% of all samples in the cluster group 204 it is assigned the highest frequency class. The relative proportions of samples in each cluster 205 group have been calculated for each fossil pollen 200-year time window and the modern 206 pollen surface samples.

207

Non-metric multidimensional scaling (nMDS) was applied to the data using the R 'vegan' package (Oksanen et al., 2016) as a complementary method to summarise major variation in the dataset. Kernel density plots of nMDS scores have been produced to show the overlap or distinctiveness of cluster groups in ordination space with contours to highlight greater density of samples. Simpson's diversity index has been calculated for each pollen sample using raw pollen percentage data. This diversity index was selected as it takes both species richness and evenness into account (Simpson, 1949) and is frequently used to explore diversity change in

215 pollen datasets (e.g. Morris et al., 2014). Although species diversity indices provide only 216 limited information about community ecology and species abundance (Chiarucci et al., 2011) 217 this index provides a complementary method and has allowed diversity change in the pollen 218 datasets to be explored independently of the cluster analysis approach. The differential pollen 219 productivity of different plant types is not accounted for in the methods presented here; 220 therefore, some high pollen producers, such as *Pinus*, may be over-represented in some 221 clusters. Lower pollen producers are recognised within the method, as they may appear 222 consistently represented in certain clusters as frequent taxa in spite of their lower abundance 223 values. 224 225 **RESULTS** 226 227 **Defining pollen cluster groups** 228 229 After analysing the community classification of various groups of sites at different hierarchical levels (Table 1) within the dendrogram, 16 clusters have been chosen as a 230 231 suitable number to reflect the range of land cover types represented in the pollen dataset 232 (Table 2). These are based on a hierarchical classification that aggregates the 16 clusters into 233 eight major groups. Preston et al. (2013) suggested that decisions about the number of 234 clusters should be "based on the purpose of the study because there is no single 'correct' 235 classification". Choosing 16 clusters allows us to address the aims of this study and identify changes in key vegetation assemblages through time. Some of the open land cover clusters 236 237 are composed of complex mixtures of relatively low abundance, but frequently occurring taxa, whereas forest (closed vegetation) clusters are usually distinguished by one or two 238 239 dominant arboreal taxa (Fig. 2). The open vegetation clusters include both 'anthromes'

240 (anthropogenic biomes; Ellis, 2015), such as grassland, with indicators of arable and 241 disturbed land, and naturally occurring open land cover types, such as steppe. Some clusters 242 were composed entirely of either fossil or modern samples, i.e. indicating the possible 243 existence of non-analogue vegetation assemblages. Clusters are presented according to their corresponding 'super group', i.e. a broader group that they are statistically assigned to at a 244 245 lower clustering level (Table 1). Many of the pollen records do not cover the entire Holocene; 246 therefore some periods of time are represented by fewer sites than others. The number of sites 247 does not drop below 30 for any time interval, which is a suitable number for calculating 248 averages. However, site numbers do drop below 30 when split by elevation group, and 249 interpretations need to be made with caution when site numbers are very low for a particular 250 time window.

251

252 Some cluster groups comprise pollen samples that are similar to one another while other 253 cluster groups are more diverse when mapped by nMDS scores (Fig. 3). For example, all of 254 the open vegetation clusters occupy a larger area in nMDS space while the forested clusters occupy smaller areas on the plots. The nMDS stress value was 0.224, which indicates that the 255 256 analysis cannot adequately collapse the variability within the 260 taxa across the samples into two dimensions. This explains the overlap of clusters within the nMDS ordination space and 257 258 highlights the value of the cluster analysis approach in providing a more detailed and useful 259 way to identify major vegetation types.

260

261 Defining Mediterranean vegetation types

262

263 *Open vegetation assemblages*

264 The pollen taxa assemblages derived for clusters 1.1 (sclerophyllous parkland), 1.2 265 (evergreen oak shrubland: Oleaceae), 1.3 (sclerophyllous steppe/parkland) and 1.4 266 (parkland/grassland) are typical of open landscapes, which are often recognised as being 267 human modified (Table 1; see Fyfe et al., 2018). Cluster 1.1 is a very mixed group with 268 median values below 8% for all frequently-occurring taxa, while cluster 1.2 has high values 269 for Oleaceae, which includes both cultivated olive groves and wild trees. The most frequent 270 and abundant taxon in cluster 1.3 is goosefoot family (Chenopodiaceae), a typical steppe-land 271 type, but this cluster also includes frequent occurrence and relatively high abundance of pine 272 (Pinaceae), grass (Poaceae) and mugwort/wormwood (Artemisia). Cluster 1.4 273 (parkland/grassland) is a very large group that includes 583 pollen samples. The cluster 274 includes high values for grass, with frequent occurrence of pine, deciduous/evergreen oak 275 (Quercus), arable and grassland indicators. Many of these taxa have broad IQR values 276 indicating a mixed and varied group. Within cluster 2.0 (evergreen shrubland/woodland: 277 Quercus), certain taxa have higher IQR than others, for example, evergreen oak has a fairly 278 narrow IQR, while grass in contrast has a large IQR, indicating that evergreen oak is the 279 distinctive taxon for this cluster.

280

281 The removal of aquatic pollen taxa and spores (such as *Myriophyllum, Menyanthes,*

282 *Sphagnum, Potamogeton*) from the dataset means that identifying wetlands is restricted.

283 However, sedges (Cyperaceae) include both aquatic and dryland plants, so this taxon has

been included in the dataset. Cluster 3.0 (pasture/wetland) is distinguished by high values of

sedge and also includes frequent occurrence of pine and grass. It is common at sites above

286 500 masl, where it appears to represent pastureland, as well as some lowland sites where it is

287 more likely to be indicative of wetlands.

290 Clusters 4.0 (pine forest), 5.1 (pine woods), 5.2 (pine steppe) and 7.0 (fir forest) are 291 dominated by pine (Table 1), which produces high amounts of pollen that can travel great 292 distances (Broström et al., 2008) and can be over-represented in pollen records. There are 293 many Mediterranean pine species with contrasting ecologies, ranging from *Pinus sylvestris*, 294 which is typical of cooler montane environments, to *Pinus halepensis* that grows in coastal 295 regions. These different species of pine may not always be consistently or easily 296 distinguished palynologically, and this limits the environmental significance of the combined 297 Pinus clusters; however, the associated taxa (the constant, frequent and common types) can assist in understanding the nature of the vegetation communities represented in these 298 299 samples. Cluster 5.2 (pine steppe) is dominated by a combination of pine, grass and sedge 300 with similar values for each, whereas cluster 7.0 is a distinctive fir cluster group, which also 301 includes a mixture of other deciduous and coniferous woodland types. Abies and Picea have 302 been separated from *Pinus* in the dataset and Pinaceae is mainly composed of *Pinus*, but as 303 other genera such as *Cedrus* are in the same family, the name used here to describe this group 304 is Pinaceae.

305

306 Deciduous and mixed forests/woodlands

307 This "super group" includes clusters 6.1 (deciduous oak woods), 6.2 (deciduous oak

parkland), 8.1 (alder woods), 8.2 (coniferous forest), 8.3 (beech woods) and 8.4 (mesic forest)
(see Table 1 for a list of the dominant taxa in these groups). Cluster 6.1 is a distinct deciduous
oak class whereas grass and pine are also key taxa in cluster 6.2. Cluster 8.1 includes samples
with a high, but wide, IQR for alder, with frequent but lower values of a selection of other
deciduous woodland types, pine, sedge and grass. Cluster 8.3 has high average values for
beech (*Fagus*) and other taxa that occur frequently in the dataset, and cluster 8.2 represents a

314 combination of spruce, pine and fir. Cluster 8.4 is distinct from the other clusters in this broad
315 group and described as 'mesic forest' with high average values for elm, hazel and oak,
316 although, pine, grass, birch, alder and sedge also frequently occur in samples in this group
317 and all taxa have fairly wide IQR, indicating that this is a varied woodland group.

318

319 Deciphering patterns of vegetation change

320

321 The overall patterns of Holocene vegetation change (Fig. 4) reveal that pine forest (4.0), pine 322 woods (5.1) and deciduous oak woods (6.1) dominated the early Holocene period prior to 8000 BP. These were then partly replaced by other cluster groups, especially wooded 323 324 parkland (e.g. cluster 6.2: deciduous oak parkland and 1.1: sclerophyllous parkland) into the 325 mid-Holocene (i.e. 8000-4500 BP). Some vegetation clusters increased in abundance during 326 the latter part of the record after ~4500 BP (e.g. 3.0 (pasture/wetland) and 1.4 327 (parkland/grassland). A number of clusters also only appear in the second half of the 328 Holocene, such as 8.3 (beech woods), 8.1 (alder woods) and 1.2 (evergreen shrubland: 329 Oleaceae). The declining nMDS scores and varied ratio of change in the early Holocene 330 appears to reflect shifts in vegetation patterns as a result of natural processes, notably 331 adaptation to the onset of interglacial climatic conditions. This was followed a period of 332 greater vegetation stability between ~9000 and 6000 BP with more consistent nMDS scores 333 and steadily increasing diversity. Since the mid to late Holocene there has been an increasing ratio of change in the pollen-inferred vegetation clusters (since ~5000 BP) and nMDS scores 334 (since ~2000 BP), while the Simpson's diversity index steadily increased throughout the 335 336 entire Holocene. The nMDS patterns (Fig. 3) indicate that the open vegetation types that have prevailed during the later Holocene are more diverse taxonomically. Overall, the ratio of 337 change and analysis of ecosystem trajectories (Fyfe et al., 2018) indicate continual turnover 338

342	Grouping sites according to elevation has revealed that needle-leaf forests (4.0, 5.1 and 5.2)
343	and deciduous oak woods/parkland (6.1 and 6.2) were abundant in the early Holocene at the
344	lowest elevation (below 500 m) (Fig. 5). Parkland/grassland (1.4), sclerophyllous
345	steppe/parkland (1.3) and pasture/wetland (3.0) then become more abundant from the mid-
346	Holocene, and in the most recent 5000 years evergreen shrubland (Oleaceae) (1.2) and alder
347	woods (8.1) increase. Evergreen shrubland (1.2) is most abundant in this lowest elevation
348	group. At mid-elevation levels (between 500-1200 m) in addition to pine forest types, mesic
349	forest (8.4) has high values in the early Holocene, while beech woods (8.3) become
350	significant after ~3500 BP. Conifer trees are most common at the highest elevation level
351	(above 1200 m), with pine forest/woods types (4, 5.1 and 5.2) having very high values in the
352	early Holocene (greater than 80% when pine clusters are combined), which are then partly
353	replaced by fir, conifer and pine steppe (7.0, 8.2 and 5.2).
354	
355	There are cluster groups in the fossil record that are not represented in the modern dataset; for
356	example, mesic forest (8.4), which may represent a non-analogue community, as it is also
357	absent in the most recent fossil pollen time window (200 BP - present). However, there are no
358	groups that are present in the modern dataset that are not also present in the fossil dataset,
359	although some clusters appear rarely, for example, 2.0 (evergreen shrubland: Quercus) and
360	1.2 (evergreen shrubland: Oleaceae). The modern surface samples cover a much larger
361	geographical area, as there are around ten times more site locations than in the fossil dataset,
362	and greater diversity of landscape types may be therefore expected in the modern dataset. On

the other hand, the fossil dataset is much larger than the surface pollen dataset in terms of the
number of samples included and encompasses landscape types that have changed over time.

366 **DISCUSSION**

367

368 The pollen-based results indicate that modern/recent vegetation in the Mediterranean basin is 369 spatially diverse, in accord with vegetation mapping using remote sensing and ground-level 370 field survey, as reflected by the Corine remote sensed land cover map (Fig. 6). The similarity 371 between pollen clusters and the Corine maps is best in areas with homogeneous land cover, whereas more heterogeneous landscapes lead to poorer matches (Woodbridge et al., 2014). 372 373 The Corine map and pollen-inferred vegetation clusters show good overall similarities, for 374 example, parkland/grassland (1.4) and sclerophyllous parkland (1.1) are similarly reflected by 375 both datasets in north central Spain while the evergreen shrubland (Oleaceae) (1.2) cluster is 376 located in areas of olive groves in south central Spain shown on Corine maps. However, there 377 are also dissimilarities, which relate to the difference in these two data types in sensing vegetation patterns (Woodbridge et al., 2014). This partly results from the amalgamation of 378 379 pollen types reaching a site in a mosaic landscape and the difficulty in identifying some 380 landscape types palynologically. Differential pollen productivity and dispersal may lead to 381 over-representation of pine forests and under-representation of some herbaceous taxa and 382 thus open vegetation types.

383

Plotting pollen samples in nMDS space according to cluster group (Fig. 3) provides an
independent assessment of the major patterns within the dataset, although the full range of
variability is not captured by two dimensions. Similar vegetation types identified according to
the cluster analysis approach occupy similar areas of nMDS ordination space, such as all pine

dominated forest types (4.0, 5.1 and 5.2) while the two evergreen shrubland types (1.2 and
2.0) also occupy similar nMDS space.

390

391 Temporal dynamics in vegetation patterns

392

393 The cluster analysis-based approach and datasets presented here and in a companion paper 394 (Fyfe et al., 2018) focussing on exploring trajectories of change, permit a more detailed 395 description of Mediterranean vegetation change through the Holocene than some previous 396 studies. Prentice et al. (1996), for example, identified only four key vegetation types in the 397 Mediterranean: xerophytic woods/scrub, temperate deciduous forest, broad-leaved 398 evergreen/warm mixed forest, and steppe. These vegetation types are reflected by the limited 399 array of pollen clusters shown in Fig. 6. Within the 'BIOME 6000' mapping project (Prentice 400 et al., 1996), the Mediterranean was dominated by "temperate deciduous broad-leaf forest" at 6000 BP and by "xerophytic woods/scrub" and "warm-temperate evergreen broad-leaf and 401 402 mixed forest" at 0 BP. The vegetation clusters presented in our new analysis reveal a much 403 wider range of forest types, which appear to shift northwards between 6000 BP and the 404 present day. For example, they include 6.2 (deciduous oak parkland), 4.0 (pine forest) and 5.1 405 (pine woods) in northern Spain at 6000 BP, while 1.1 and 1.3 (sclerophyllous clusters) were 406 restricted to southern Spain. The forest types in northern Spain were then replaced by clusters 407 1.1 and 1.3 and other open vegetation types (e.g. 1.4 (parkland/grassland) and 3.0 (pasture/wetland) while some forests persisted, primarily in upland regions, with the 408 appearance of others such as 8.3 (beech woods). This matches the BIOME 6000 analysis of 409 410 vegetation change in Iberia and Greece between the mid-Holocene and modern presented in 411 Roberts et al (2004; plate 7).

413 In a cluster analysis-based approach, Huntley (1990) identified key taxa associations using percentage data for 44 pollen taxa, which were placed on a six-point smoothed scale, and 414 415 mapped these across Europe for specific time slices. According to Huntley (1990), at 6000 416 BP the Mediterranean was represented by three vegetation types (described in Table 2). The 417 pollen cluster results presented for the current study provide a more detailed reconstruction of 418 vegetation assemblages and greater representation of open landscapes (Fig. 6). For example, cluster 1.4 (parkland/grassland) and 1.1 (sclerophyllous parkland) are the main open 419 420 landscapes in Italy, while 1.3 (sclerophyllous steppe/parkland) and 1.2 (evergreen shrubland: 421 Oleaceae) are more widely represented in Spain.

422

423 Differences between the cluster results for the modern and most recent fossil datasets (Fig. 4) 424 may be partly influenced by differences in the spatial distribution of sites. Coniferous, fir and 425 alder woods (clusters 8.2, 7.0 and 8.1) are poorly represented in the surface pollen dataset, 426 but are present in the fossil data, including the most recent time window (200 BP - present). 427 The modern samples are not always co-located with the fossil datasets and thus sample 428 different parts of the landscape. Differences between modern and most recent fossil pollen 429 vegetation types are also due to real changes in land cover that have occurred in recent 430 decades, and are linked to both direct and indirect anthropogenic intervention. This has been 431 demonstrated by Colombaroli et al. (2007) with the disappearance of Abies alba at low 432 altitudes in the Mediterranean since the Neolithic. It is clearly reflected in the cluster results (Fig. 5) when sites are separated by elevation groups, as fir forest (7.0) is well represented 433 above 500 m, but does not appear in the dataset below this altitude. 434

435

436 Various factors lead to the existence of non-analogue vegetation assemblages (Williams &
437 Jackson, 2007), such as rates of postglacial migration (Médail & Diadema, 2009), climate

438 change, human disturbance (Jalut et al., 2009) and altitudinal changes; for example, Geri et 439 al. (2010) found that loss of Mediterranean forest has mostly been confined to low altitude 440 areas. The altitudinal differences in the dataset reflect the ecotone boundaries of vegetation 441 types and land use differences. For example, the only open vegetation type present >1200 m 442 is 1.4 (parkland/grassland) and the only forest types represented at low elevation are 6.1 and 443 6.2 (deciduous oak parkland/woods). In a comparison of Holocene vegetation diversity across 444 a transect from central Europe to the Mediterranean, Colombaroli & Tinner (2013) identified how diverse open-land ecosystems increase when human disturbance intensifies while forest 445 446 becomes less diverse, highlighting the important role of anthropogenic fire and other disturbance related activities. Pollen taxa diversity appears to increase with the development 447 448 of cultural landscapes and not just as a by-product of deforestation (Colombaroli & Tinner, 449 2013).

450

451 The patterns of change in different plant species and communities may sometimes be missed 452 in the broad scale synthesis presented here. For example, evergreen *Quercus ilex* communities, which are recognised as more fire resistant than broad-leaf oak forests, 453 454 developed before the emergence of Neolithic farming and were heavily impacted by land use strategies (e.g. Calò et al., 2012), but this pattern is not clearly shown in the changing 455 456 abundance of the evergreen shrubland (dominated by Quercus) cluster (2.0). This suggests 457 that regional patterns may become hidden within the broad scale synthesis, and that smaller spatial scale analyses would be required to investigate sub-regional patterns of vegetation 458 459 change in more detail.

460

461 **Potential and actual natural vegetation**

463 There has been recent debate about potential natural vegetation (PNV) and the disconnect 464 between PNV, actual vegetation and the role that palaeoecological evidence of past vegetation might play in determining or assessing PNV (e.g. Carrión & Fernandez, 2009; 465 466 Chiarucci et al., 2010; Loidi et al., 2010; Jackson, 2013). Jackson (2013) sees PNV as a 467 'useful fiction', but one that might be better informed through use of data such as those 468 presented here, which can be used to develop understanding of the processes that controlled 469 historical vegetation character and change. PNV, when applied to the Iberian Peninsula, has 470 modelled vegetation with a single dominant tree taxon at broad spatial scales (Carrión & 471 Fernandez, 2009). This is not the case for the clusters generated in this analysis, which indicates frequent co-dominant species, even taking into account differential pollen 472 473 productivity between taxa (Broström et al., 2008). Whilst it is recognised that the PNV 474 concept encapsulates vegetation that would develop in the absence (or cessation of) human 475 activity (Loidi et al., 2010), it is clear from palynological records that vegetation turnover 476 rather than stability has been the norm in the Mediterranean, even in the absence of human 477 impacts associated with agriculture and pastoralism.

478

479 Several long, continuous pollen records from the Mediterranean (e.g. Sadori et al., 2013) 480 have demonstrated continuous vegetation change during past interglacial periods. These 481 suggest that the character of 'natural' vegetation succession in the Mediterranean has been 482 predictable and comparable during different interglacials (Tzedakis, 2007). Open woodland develops first, followed by expansion of deciduous oak, sclerophyll scrub, elm, hazel and 483 lime, then by fir, beech and spruce, followed by revertance to open woodland. The fact that 484 485 the early Holocene vegetation clusters presented here show similar patterns implies that predominantly natural processes were driving vegetation succession at least until ~8000 BP. 486 487 Magri (1995) suggested that the Holocene interglacial may currently be experiencing a later 488 phase that matches previous interglacials during which natural openings in forests and the 489 development of grasslands produced environments suitable for agriculture and pastoralism in 490 the current interglacial. However, the complexity of land cover types and frequency of 491 vegetation cluster change during the second half of the Holocene is likely to have been 492 amplified by intensifying human impacts, a pattern also identified by Carrión and Fernandez 493 (2009). In the pollen-based synthesis presented here, for example, total non-arboreal pollen across the Mediterranean increased from ~35% to ~47% between 3500 BP and recent times 494 495 (Fig. 4). Looking to the future, dynamic vegetation models driven by future climate change 496 scenarios predict a shift from mesic forest (the current PNV) to more open shrubland and 497 eventually steppe as a result of reduced water availability (Hickler et al., 2012). 498 Transformation of the vegetation by human agency has already begun this process over the 499 past millennia. 500 501 **CONCLUSIONS**

502

503 We have demonstrated a method that allows exploration of vegetation dynamics in time and 504 space and potentially the detection of non-analogue communities. A high degree of spatio-505 temporal heterogeneity is recognised in Mediterranean vegetation patterns (Shoshany, 2000) 506 and this is evident within the results presented here. In comparison with some previous 507 classification studies, such as the BIOME mapping vegetation classification (Prentice et al., 1996; Roberts et al., 2004), which often focussed on discrete time windows, our results show 508 509 almost continuous turnover of most plant communities during the last 11,000 years, as 510 demonstrated by the rate of vegetation cluster change (Fig. 4). This indicates that stable 511 baseline vegetation communities cannot be easily identified, at least not at a pan-512 Mediterranean scale. Our transformation and synthesis of multiple modern and fossil pollen

records has revealed key vegetation types that existed in the past and are rare or absent in the
modern landscape (e.g. mesic forest) and vice versa (e.g. evergreen shrub dominated by
Oleaceae). The pollen cluster results highlight the potential of this approach to explore
complex vegetation patterns and provide detailed information about vegetation assemblages
not previously identified, which is open to future development with the availability of
additional datasets.

519

520 While it can be difficult to establish precise causes for changes in vegetation, stochastic 521 characteristics, associated with a range of disturbance factors, such as natural fire, climate fluctuations and human impacts, will have resulted in what are often considered to represent 522 523 'subclimax' vegetation communities. There is long-standing evidence of the complex 524 interaction between plant assemblages, disturbance and climate (Colombaroli et al., 2007; 525 Guiot & Kaniewski). If stable natural vegetation communities never existed when viewed on 526 a Holocene timescale, then attempts to use them as targets for environmental restoration may 527 be misplaced. Understanding land cover types in an area of the world with a long-standing 528 (multi-millennial) history of human transformation has potential value in informing 529 palaeoecologists, conservationists, and climate modellers, among other groups concerned 530 with vegetation change and has potential to help identify biogeographic regions that are more 531 resilient or susceptible to future disturbance.

532

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534

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542	helpful comments.
543	
544	DATA AVAILABILITY STATEMENT
545	
546	The datasets underpinning this study are publically available from the European Pollen
547	Database (http://www.europeanpollendatabase.net/). The results of the analyses carried out
548	are available from the authors upon request.
549	
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719	41 , 2080-2092.
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721	BIOSKETCHES
722	
723	Jessie Woodbridge is a Post-Doctoral Research Fellow at Plymouth University (UK) and is
724	conducting research that aims to reconstruct changes in European land cover throughout the
725	Holocene using pollen data. Her research background is based on palaeoenvironmental
726	reconstruction, specifically in relation to understanding human impacts and past climate
727	trends.
728	
729	Ralph Fyfe is Professor of Geospatial Information at Plymouth University. His research
730	focuses on developing innovative methodological approaches to reconstruct patterns of
731	vegetation cover at local to European scales, and deciphering the processes that led to these
732	patterns with an emphasis on human agency.
733	
734	Neil Roberts is Professor of Geography at Plymouth University. His research emphasises past
735	climatic and environmental change since the last glacial maximum, specifically using lake-

736 sediment archives from eastern Africa, the Mediterranean and West Asia. Professor Roberts 737 is a fellow of the Royal Geographical Society, and has served on national and international 738 committees concerning past global changes. 739 740 **TABLES** 741 742 Table 1 Comparison of pollen-inferred vegetation groups within the modern and fossil 743 datasets using Ward's hierarchical clustering method. Towards the left of the table, the 744 groups combine at lower levels of clustering and to the right cluster groups separate as different pollen taxa assemblages are detected at a higher clustering level. The inter-quartile 745 746 range of the dominant taxa (appearing in 81-100% of samples in the cluster) are provided. 747 748 Table 2 Pollen-inferred vegetation clusters compared with Corine remote sensed land cover 749 maps (EEA, 2016) and vegetation types defined by previous studies (Huntley, 1990; Prentice 750 et al., 1996; Peyron et al., 1998). 751 752 **FIGURES** 753 754 Fig 1 Modern and fossil pollen site locations colour coded according to elevation group. The 755 white area represents the Mediterranean region for which results are presented in this study. 756 Pollen samples in the Rhone Valley of southern France are also shown on the map as these 757 sites contributed to the generation of vegetation cluster groups. 758 759 Fig 2 Box plots for pollen taxa within each vegetation cluster group for the combined fossil

and modern pollen datasets. The middle vertical bar for each taxon represents the median, the

761 lower and upper boundaries of the boxes show the interquartile range, and the dashed 762 horizontal lines show the minimum and maximum occurrence of the taxon in the cluster. The 763 bars representing taxa that are frequent or abundant in a specific cluster group are coloured 764 black and those that are not frequent or abundant are grey (a version of this diagrams has 765 been published in Fyfe et al., 2018). 766 767 Fig 3 Modern and fossil pollen samples: non-metric multidimensional scaling (nMDS) kernel 768 density plots for 16 cluster analysis derived vegetation groups. Darker colours indicate 769 greater density of samples and lighter colours indicate fewer samples. 770 771 Fig 4 Percentage of pollen samples statistically assigned to each cluster group throughout the 772 Holocene for the entire Mediterranean combined fossil and modern dataset with ratio of 773 vegetation cluster group change, average non-metric multidimensional scaling (nMDS) 774 scores, average Simpson's diversity index and average non-arboreal pollen sum (% NAP) 775 (scale: 28-48%) (11,000 BP to modern). 776 777 Fig 5 Percentage of pollen samples statistically assigned to each cluster group throughout the Holocene separated by elevation group: a) <500 m (415 modern and 47 fossil pollen sites), b) 778 779 500-1200 m (643 modern and 28 fossil pollen sites), and c) >1200 m (498 modern and 30 780 fossil pollen sites) (11,000 BP to modern). 781 782 Fig 6 Pollen-inferred vegetation clusters (6200-6000 BP and modern surface samples) 783 compared with the results of BIOME 6000 vegetation maps (Prentice et al., 1996) for modern 784 pollen (0 BP) and 6000 BP and the Corine remotely-sensed (EEA, 2006) land cover map. 785

786	Fig 7 Frequency of vegetation types according to the pollen cluster results and the BIOME
787	6000 vegetation maps for 0 BP and 6000 BP maps (Prentice et al., 1996). Similar vegetation
788	types are shown alongside one another.
789	
790	Supplementary Information 1: EPD fossil pollen site metadata for the records used in these
791	analyses (for details of the modern pollen sites see Davis et al., 2013).
792	
792 793	Supplementary Information 2: Pollen-inferred vegetation classification tables: land-cover
	Supplementary Information 2: Pollen-inferred vegetation classification tables: land-cover types (clusters) and pollen taxa percentage (inter-quartile range) in the combined modern and
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793 794	types (clusters) and pollen taxa percentage (inter-quartile range) in the combined modern and

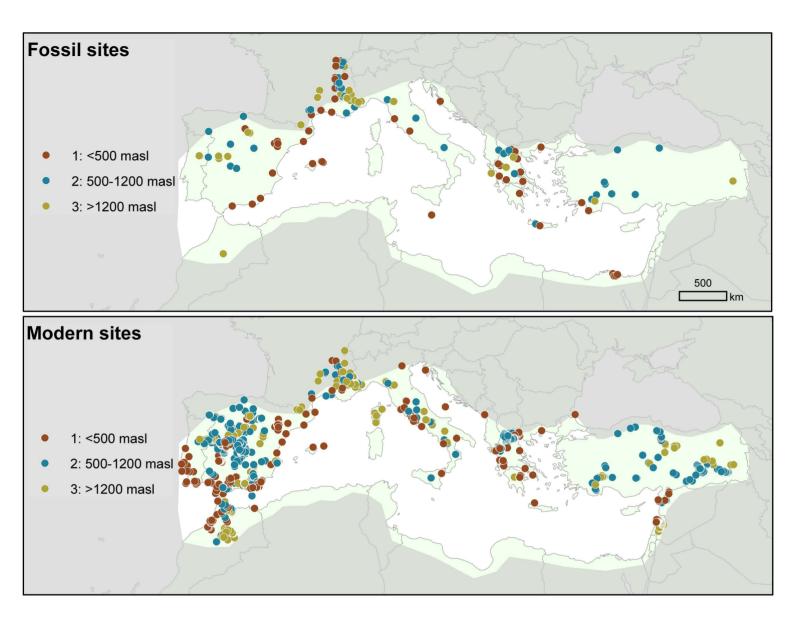
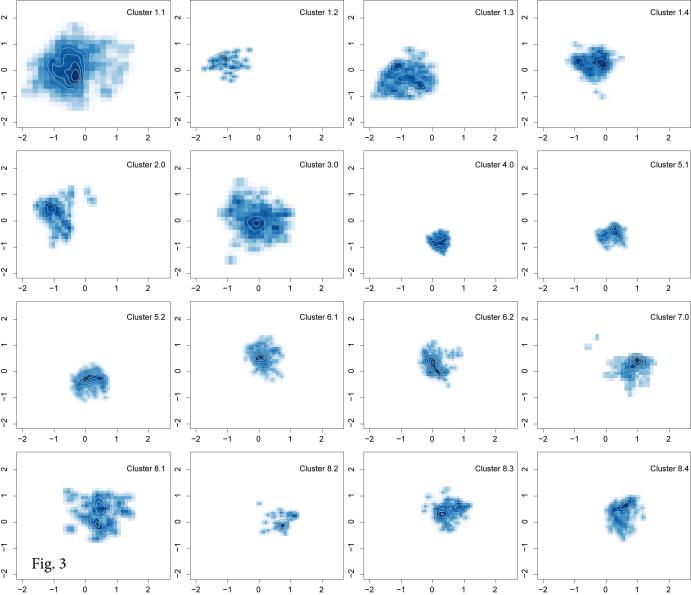
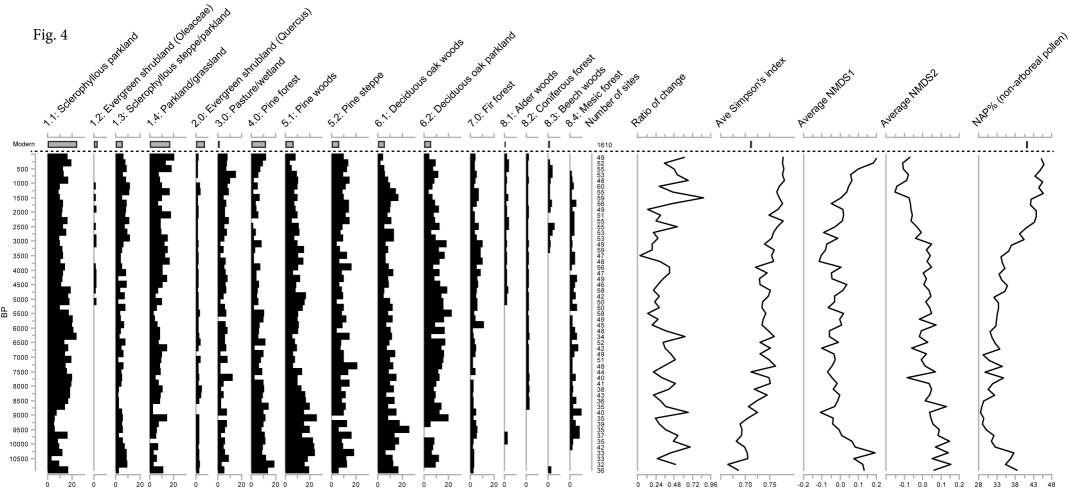
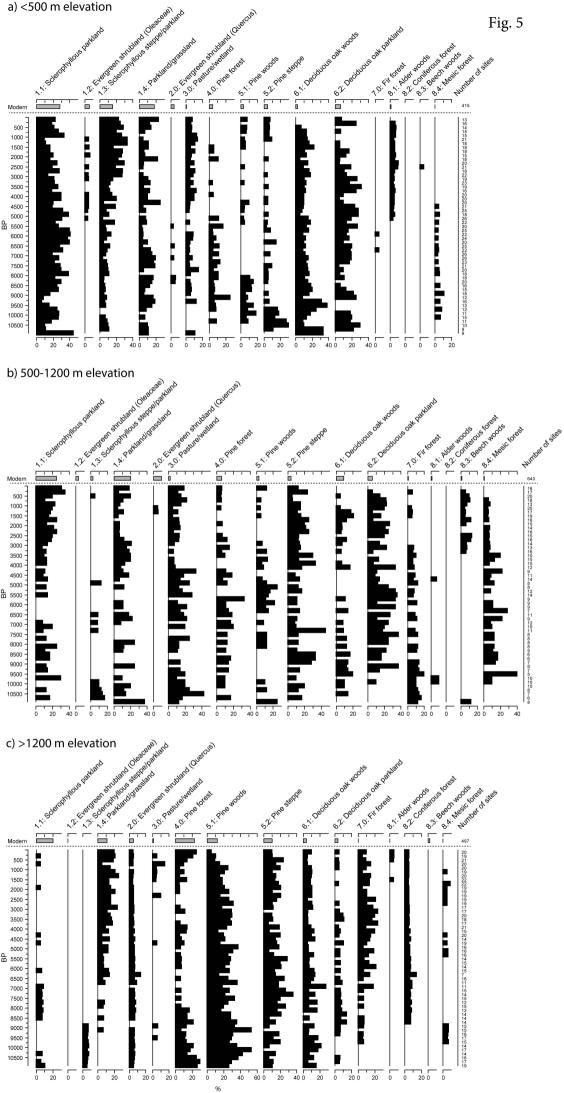
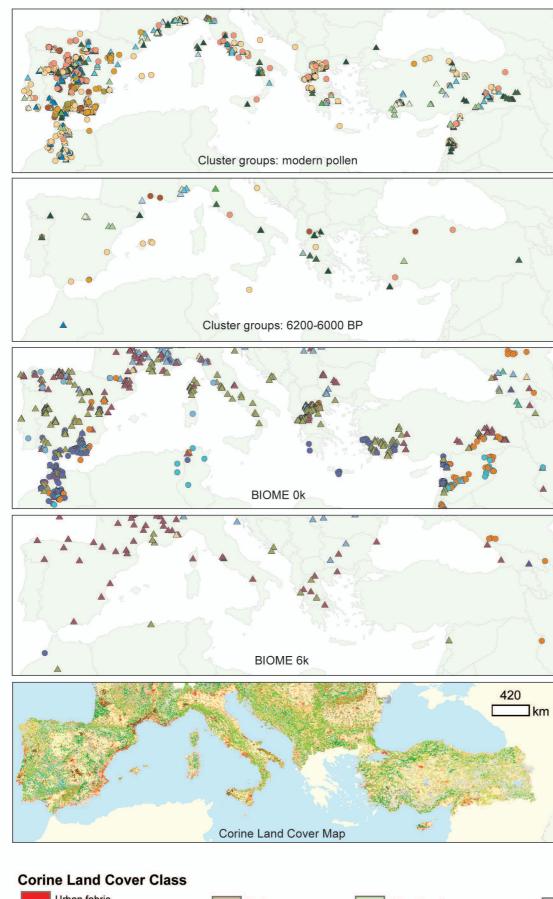


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Asteraceae (Lactucoides)-	1	I	₽ +	[+								H	e i	4	E I	
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Ranunculaceae-			4	4	4		-	4	} I	H-I	F H	ŀ	P (4	H	-1
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Pollen cluster group

- 1.1: Sclerophyllous parkland
- 1.2: Evergreen shrubland (Oleaceae)
- 1.3: Sclerophyllous steppe/parkland
- 1.4: Parkland/grassland
- ▲ 2.0: Evergreen shrubland (Quercus)
- 3.0: Pasture/wetland
- ▲ 4.0: Pine forest
- △ 5.1: Pine woods
- ▲ 5.2: Pine steppe
- ▲ 6.1: Deciduous oak woods
- ▲ 6.2: Deciduous oak parkland
- ▲ 7.0: Fir forest
- ▲ 8.1: Alder woods
- ▲ 8.2: Coniferous forest
- ▲ 8.3: Beech woods
- △ 8.4: Mesic forest

BIOME vegetation types

- Steppe
- Xerophytic woods/scrub
- Tundra
- Warm-temperate evergreen broadleaf and mixed forest
- Cool evergreen needleleaf forest
- Cool mixed forest
 Cool-temperate evergreen needleleaf
- and mixed forest
- Cold evergreen needleleaf forest
 Temperate deciduous broadleaf forest
- Desert



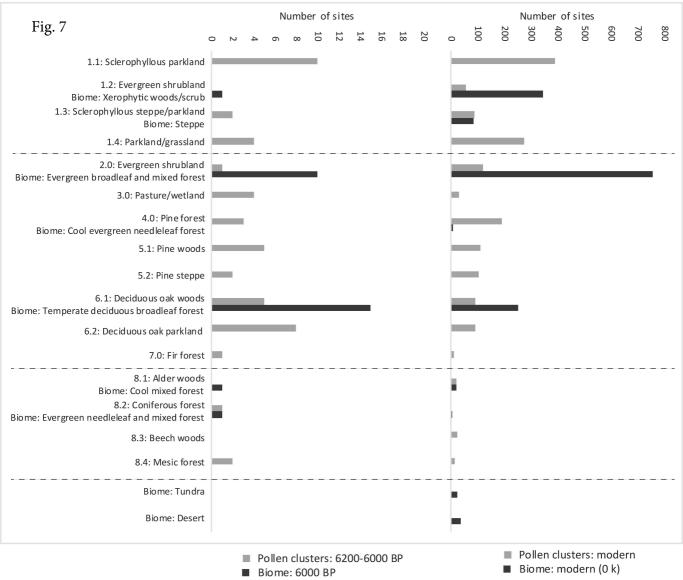


Table 1 Comparison of pollen-inferred vegetation groups within the modern and fossil datasets using Ward's hierarchical clustering method. Towards the left of the table, the groups combine at lower levels of clustering and to the right cluster groups separate as different pollen taxa assemblages are detected at a higher clustering level. The inter-quartile range of the dominant taxa (appearing in 81-100% of samples in the cluster) are provided.

3 clusters	8 clusters	12 clusters	13 clusters	15 clusters	16 clusters	Dominant taxa (most frequently occurring taxa within the cluster)
		1	1	1	1.1 : Sclerophyllous parkland	Poaceae (4-12.5), Pinaceae (2.8-13.3), <i>Quercus</i> deciduous (1.4-8.6), <i>Asteraceae</i> : <i>Asteroideae</i> (0.9-6.4), <i>Quercus</i> evergreen (11.2-14.9), Oleaceae (10.7-8.6), <i>Asteraceae</i> : <i>Lactuceae</i> (10.8-6.2), Ericaceae (10.5-6.2)
	1: Open vegetation/			9	1.2 : Evergreen shrubland (Oleaceae)	Oleaceae (39-61.8), <i>Quercus</i> evergreen (5.4-17.8), Poaceae (6-12.9), <i>Asteraceae:</i> <i>Asteroideae</i> (1.1-5.2), Chenopodiaceae (0.7-4.4), Cyperaceae (10.8-6.1)
1: open	shrubland	3	3	3	1.3 : Sclerophyllous steppe/parkland	Chenopodiaceae (29.1-56.7), Pinaceae (2.4-18), Poaceae (2.9-13.4), Artemisia (1.1- 10.9), <i>Quercus</i> evergreen (1.5-8.3), Oleaceae (0.7-7.3)
vegetation		7	8	8	1.4: Parkland/grassland	Poaceae (26.2-43), <i>Quercus</i> deciduous (2.3-11.3), Pinaceae (2.4-10.3), Asteraceae: Lactuceae (0.5-4.4), Asteraceae: Asteroideae (0.6-3.3), Cyperaceae (I1.5-11.3)
	2: Evergreen shrubland	5	5	7	2.0: Evergreen shrubland (Quercus)	Quercus evergreen (50.6-69.4), Poaceae (4.2-23.7), Brassicaceae (0.5-2), Chenopodiaceae (0.4-1), Pinaceae (I2-10)
	3: Pasture/wetland	11	12	14	3.0: Pasture/wetland	Cyperaceae (34.6-52.2), Pinaceae (5.2-18.8), Poaceae (4.4-15.2), <i>Quercus</i> deciduous (2.1-10.1)
	4: Pine forest	2	2	2	4.0: Pine forest	Pinaceae (67-82.1), Poaceae (1.5-4.8), <i>Quercus</i> deciduous (0.6-4.2), Betula (10.4-5.7)
2: Pine	5: Pine woods/steppe		4	4	5.1: Pine woods	Pinaceae (43.3-54.5), Poaceae (3.5-8.9), <i>Quercus</i> deciduous (1.1-11.2)
forests		4	6	6	5.2: Pine steppe	Pinaceae (26.9-40.5), Poaceae (3.4-16.9), Cyperaceae (2-18.3), <i>Quercus</i> deciduous (1.9-9.7), Artemisia (0.4-1.7)
	6: Deciduous (broad-			7	6.1 : Deciduous oak woods	Quercus deciduous (40.8-53.5), Poaceae (4.2-13.8), Pinaceae (2.3-8.9)
	leaved)	6	7	10	6.2: Deciduous oak parkland	Quercus deciduous (23.1-35.4), Poaceae (7.1-18.3), Pinaceae (3.7-21.5)
3: Deciduous	7: Fir forest	8	9	11	7.0 : Fir forest	Abies (31.4-52.3), Quercus deciduous (5.7-11.6), Corylus (3.2-10.5), Pinaceae (2-10.2)
and mixed forests		9	10	12	8.1: Alder woods 8.2: Coniferous forest	Alnus (25.8-46.7), Quercus deciduous (4.2-9.4), Cyperaceae (1.8-9.6)
	8: Mixed woods	10	11	13	8.3: Beech woods	Picea (22.6-32.7), Pinaceae (3.6-18.2), Abies (4.3-17.2)
		12	13	15	8.4: Mesic forest	Fagus (18.7-36.3), <i>Quercus</i> deciduous (9.2-16), Poaceae (3.7-14.1), Corylus (3.5-8.4)

Table 2 Pollen-inferred vegetation clusters compared with Corine remote sensed land cover maps (EEA, 2016) and vegetation types defined by previous studies (Huntley, 1990; Prentice et al., 1996; Peyron et al., 1998).

Cluster name descriptor	Corine Land Cover Classes	Comparison with previous studies					
Cluster name descriptor	Conne Land Cover Classes	Prentice et al. (1996) Peyron et al. (1998)	Huntley (1990)				
1.1: Sclerophyllous parkland (Poaceae- <i>Quercus</i> -Oleaceae-herbaceous and sclerophyll taxa)	Sclerophyllous vegetation	Grass / heath	Quercus-Pinus-sclerophyll taxa Quercus-Chenopodiaceae- Pistacia				
1.2: Evergreen shrubland (Oleaceae-Quercus evergreen-Poaceae- herbaceous and sclerophyllous taxa)	Transitional woodland-shrub / broad-leaved forest / sclerophyllous	Warm-temperate sclerophyll trees/shrub	<i>Quercus</i> -Chenopodiaceae- Pistacia				
1.3 : Sclerophyllous steppe/parkland (Chenopodiaceae-Poaceae-Artemisia- Quercus evergreen-Oleaceae)	Sclerophyllous vegetation	Steppe forb / shrub	Chenopodiaceae <i>Quercus</i> -Pinus-Ericales- Chenopodiaceae				
 1.4: Parkland/grassland (Poaceae, <i>Quercus</i> deciduous, Asteraceae) 	Natural grasslands / pastures / agriculture	Grass	No similar group				
2.0 : Evergreen shrubland (<i>Quercus</i> evergreen, Poaceae)	Transitional woodland-shrub / broad-leaved forest / sclerophyllous	Warm temperate broad-leaved evergreen	<i>Quercus</i> -Chenopodiaceae- Pistacia				
3.0 : Pasture/wetland (Cyperaceae, Poaceae, Quercus deciduous)	Moors and heathland	Сурегасеае	Quercus-Pinus-sclerophyll taxa				
4.0: Pine forest (Pinaceae)	Coniferous forest / mixed forest	Boreal evergreen conifer	Pinus-Betula				
5.1 : Pine woods (Pinaceae-Poaceae- <i>Quercus</i> deciduous)	Coniferous forest / mixed forest	Boreal evergreen conifer	Pinus-Betula				
5.2 : Pine steppe (Pinaceae-Poaceae- <i>Quercus</i> deciduous- <i>Artemisia</i>)	Coniferous forest / mixed forest	No similar group	Pine-Betula-Chenopodiaceae- Quercus				
6.1 : Deciduous oak woods (<i>Quercus</i> -Poaceae-Pinaceae)	Broad-leaved forest	Temperate summer green	Quercus-Pinus-sclerophyll taxa				
6.2 : Deciduous oak parkland (<i>Quercus</i> -Poaceae-Pinaceae-sclerophyll taxa)	Broad-leaved forest	Temperate summer green	<i>Quercus</i> -Pinus-sclerophyll taxa				
7.0: Fir forest (<i>Abies-Quercus</i> deciduous- <i>Corylus</i> -Pinaceae)	Coniferous forest / mixed forest	Boreal evergreen conifer	No similar group				

8.1 : Alder woods (<i>Alnus-Quercus</i> deciduous-Cyperaceae)	Broad-leaved forest	Temperate summer green	Corylus-Quercus-Alnus
8.2 : Coniferous forest (<i>Picea</i> -Pinaceae- <i>Abies</i>)	Coniferous forest / mixed forest	Boreal evergreen conifer	Picea-Pinus-Betula
8.3 : Beech woods (<i>Fagus-Quercus</i> deciduous-Poaceae- <i>Corylus</i>)	Broad-leaved forest	Cool temperate summer green	Fagus-Picea-Carpinus
8.4 : Mesic forest (<i>Corylus-Quercus</i> deciduous-Poaceae-Pinaceae-Ulmaceae	Broad-leaved forest	Cool temperate summer green	Corylus-Quercus-Alnus