

EuPPollNet: A European database of plant-pollinator networks

Jose B. Lanuza ^{1,2,3} | Tiffany M. Knight ^{2,3,4} | Nerea Montes-Perez ¹ | Paola Acuña ⁵ | Matthias Albrecht ⁶ | Maddi Artamendi ^{7,8} | Isabelle Badenhauer ^{9,10,11} | Joanne M. Bennett ^{2,3} | Paolo Biella ^{12,13,14} | Ricardo Bommarco ¹⁵ | Andree Cappellari ^{16,17} | Sílvia Castro ⁵ | Yann Clough ¹⁸ | Pau Colom ¹⁹ | Joana Costa ⁵ | Christophe Dominik ⁴ | Yoko L. Dupont ²⁰ | Reinart Feldmann ²¹ | Emeline Felten ²² | Victoria Ferrero ⁵ | William Fiordaliso ²³ | Alessandro Fisogni ²⁴ | Úna Fitzpatrick ²⁵ | Marta Galloni ²⁶ | Hugo Gaspar ⁵ | Elena Gazzea ¹⁶ | Will Glenney ⁴ | Irina Goia ²⁷ | Juan Pedro González-Varo ^{1,28} | Nina Hautekèete ²⁴ | Veronica Hederström ¹⁸ | Ruben Heleno ⁵ | Sandra Hervias-Parejo ¹⁹ | Jonna Heuschele ^{2,29,30} | Bernhard Hoiss ³¹ | Andrea Holzschuh ³¹ | Sebastian Hopfenmüller ³¹ | José M. Iriondo ³² | Birgit Jauker ³³ | Frank Jauker ³³ | Jana Jersáková ³⁴ | Katharina Kallnik ³¹ | Reet Karise ³⁵ | David Kleijn ³⁶ | Stefan Klotz ⁴ | Theresia Krausl ¹⁸ | Elisabeth Kühn ⁴ | Paula Dominguez Lapido ⁷ | Carlos Lara-Romero ³² | Michelle Larkin ³⁷ | Emilien Laurent ²² | Yicong Liu ⁴ | Sara Lopes ⁵ | Francisco López-Núñez ⁵ | João Loureiro ⁵ | Ainhoa Magrach ^{7,38} | Marika Mänd ³⁵ | Natasha de Manincor ²⁴ | Lorenzo Marini ¹⁶ | Rafel Beltran Mas ¹⁹ | François Massol ²⁴ | Corina Maurer ^{6,39} | Denis Michez ²³ | Francisco P. Molina ¹ | Javier Morente-López ³² | Sarah Mullen ⁴⁰ | Georgios Nakas ⁴¹ | Lena Neuenkamp ^{42,39} | Arkadiusz Nowak ^{43,44} | Catherine J. O'Connor ^{5,45} | Aoife O'Rourke ⁴⁰ | Erik Öckinger ¹⁵ | Jens M. Olesen ²⁰ | Øystein H. Opedal ⁴⁶ | Theodora Petanidou ⁴¹ | Yves Piquot ²⁴ | Simon G. Potts ⁴⁷ | Eileen Power ⁴⁰ | Willem Proesmans ²² | Demetra Rakosy ^{4,48} | Sara Reverte ²³ | Stuart P. M. Roberts ⁴⁷ | Maj Rundlöf ⁴⁹ | Laura Russo ⁴⁹ | Bertrand Schatz ⁵⁰ | Jeroen Scheper ³⁶ | Oliver Schweiger ⁴ | Pau Enric Serra ¹⁹ | Catarina Siopa ⁵ | Henrik G. Smith ^{49,18} | Dara Stanley ⁴⁰ | Valentin Ștefan ^{4,2} | Jane C. Stout ⁴⁰ | Louis Sutter ⁶ | Elena Motivans Švara ^{2,4,3} | Sebastian Świerszcz ^{43,51} | Amibeth Thompson ^{3,2} | Anna Traveset ¹⁹ | Annette Trefflich ⁵² | Robert Tropek ^{13,53} | Adam J. Vanbergen ²² | Montserrat Vilà ¹ | Ante Vujić ⁵⁴ | Cian White ⁴⁰ | Jennifer B. Wickens ⁴⁷ | Victoria B. Wickens ⁴⁷ | Marie Winsa ¹⁵ | Leana Zoller ^{3,2} | Ignasi Bartomeus ¹

Corresponding author= barragansljose@gmail.com

All authors excluding the first three and last are ordered alphabetically

¹ Estación Biológica de Doñana (EBD-CSIC), Seville, Spain, ² German Centre for Integrative Biodiversity Research (iDiv) Halle-Jena-Leipzig, ³ Martin Luther University Halle-Wittenberg, Institute of Biology, Halle, Germany, ⁴ Department of Community Ecology, Helmholtz Centre for Environmental Research – UFZ, Halle, Germany, ⁵ Department of Life Sciences, Centre for Functional Ecology, University of Coimbra, Coimbra, Portugal, ⁶ Agroecology and Environment, Agroscope, Zürich, Switzerland, ⁷ Basque Centre for Climate Change-BC3, Leioa, Spain, ⁸ Universidad del País Vasco, Euskal Herriko Unibertsitatea (UPV-EHU), Leioa, Spain, ⁹ Centre d'Etudes Biologiques de Chizé,

Université de La Rochelle, Villiers en Bois, France, ¹⁰ LTSER “ZA Plaine & Val de Sèvre”, CNRS, Villiers en Bois, France, ¹¹ Unité de Recherche Pluridisciplinaire Prairies Plantes Fourragères, INRA, Lusignan, France, ¹² Department of Zoology, Faculty of Science, University of South Bohemia, České Budějovice, Czech Republic, ¹³ Institute of Entomology, Biology Centre of the Academy of Sciences of the Czech Republic, České Budějovice, Czech Republic, ¹⁴ Department of Earth and Environment Sciences (sect. Landscape Ecology), University of Pavia, Pavia, Italy, ¹⁵ Department of Ecology, Swedish University of Agricultural Sciences, Uppsala, Sweden, ¹⁶ Department of Agronomy, Food, Natural Resources, Animals and Environment, University of Padua, Padua, Italy, ¹⁷ Department of Biology and Biotechnology “Charles Darwin”, Sapienza University of Rome, Rome, Italy, ¹⁸ Centre for Environmental and Climate Science, Lund University, Lund, Sweden, ¹⁹ Mediterranean Institute for Advanced Studies (IMEDEA, UIB-CSIC), Esporles, Spain, ²⁰ Department of Biological Sciences, University of Aarhus, Aarhus, Denmark, ²¹ Helmholtz Centre for Environmental Research – UFZ, Leipzig, Germany, ²² Agroécologie, INRAE, Institut Agro, Université de Bourgogne, Université de Bourgogne Franche-Comté, Dijon, France, ²³ Laboratoire de Zoologie, Université de Mons, Mons, Belgium, ²⁴ Evo-Eco-Paleo, CNRS, Université de Lille, Lille, France, ²⁵ National Biodiversity Data Centre, County Waterford, Ireland, ²⁶ Department of Evolutionary Experimental Biology (BES), University of Bologna, Bologna, Italy, ²⁷ Faculty of Biology and Geology, Babeş-Bolyai University, Cluj-Napoca, Romania, ²⁸ Conservation Science Group, Department of Zoology, University of Cambridge, Cambridge, UK, ²⁹ ESS Department, Helmholtz Centre for Environmental Research – UFZ, Leipzig, Germany, ³⁰ BZF Department, Helmholtz Centre for Environmental Research – UFZ, Halle (Saale), Germany, ³¹ Department of Animal Ecology and Tropical Biology, Biocenter, University of Würzburg, Würzburg, Germany, ³² Area of Biodiversity and Conservation, ESCET, Universidad Rey Juan Carlos, Madrid, Spain, ³³ Department of Animal Ecology, Justus Liebig University Giessen, Giessen, Germany, ³⁴ Department of Ecosystems Biology, Faculty of Science, University of South Bohemia, České Budějovice, Czech Republic, ³⁵ Institute of Agricultural and Environmental Sciences, Estonian University of Life Sciences, Tartu, Estonia, ³⁶ Nature Conservation and Plant Ecology Group, Wageningen University, Wageningen, The Netherlands, ³⁷ Botany and Plant Science, School of Natural Sciences and Ryan Institute, National University of Ireland Galway, Galway, Ireland, ³⁸ IKERBASQUE, Basque Foundation for Science, Bilbao, Spain, ³⁹ Institute of Plant Sciences, University of Bern, Bern, Switzerland, ⁴⁰ Botany Department, Trinity College Dublin, Dublin, Ireland, ⁴¹ Department of Geography, University of the Aegean, Mytilene, Greece, ⁴² Department of Botany, Institute of Ecology and Earth Sciences, University of Tartu, Tartu, Estonia, ⁴³ Center for Biological Diversity Conservation, Polish Academy of Sciences, Botanical Garden, Warsaw, Poland, ⁴⁴ Laboratory of Geobotany and Plant Conservation, Department of Biosystematics, Opole University, Opole, Poland, ⁴⁵ Cardiff School of Biosciences, Cardiff University, Cardiff, UK, ⁴⁶ Faculty of Biological and Environmental Sciences, Research Centre for Ecological Change, University of Helsinki, Helsinki, Finland, ⁴⁷ Centre for Agri-Environmental Research, School of Agriculture, Policy and Development, University of Reading, Reading, UK, ⁴⁸ Department for Integrative Zoology, Faculty of Life Sciences, University Vienna, Vienna, Austria, ⁴⁹ Department of Biology, Lund University, Lund, Sweden, ⁵⁰ CEFE, EPHE-PSL, CNRS, University of Montpellier – University of Paul Valéry, Montpellier, France, ⁵¹ Polish Academy of Sciences, The Franciszek Górski Institute of Plant Physiology, Opole, Poland, ⁵² State Institute of Agriculture and Horticulture Saxony-Anhalt, Bernburg, Germany, ⁵³ Department of Ecology, Faculty of Science, Charles University, Prague, Czech Republic, ⁵⁴ Department of Biology and Ecology, Faculty of Sciences, University of Novi Sad, Novi Sad, Serbia

Abstract

Motivation: Pollinators play a crucial role in maintaining Earth’s terrestrial biodiversity and human food production by mediating sexual reproduction for most flowering plants. Indeed, the intricate network of interactions formed by plants and pollinators constitutes the backbone of plant-pollinator community stability and functioning. However, rapid human-induced environmental changes are compromising its long-term persistence. One of the major challenges for pollinator conservation is the lack of robust generalisable data capturing how plant-pollinator communities are structured across space and time. Here, we present the EuPPollNet database, a fully open and reproducible European-level database containing harmonized taxonomic data on plant-pollinator interactions referenced in both space and time. This database offers an open workflow that allows researchers to track data-curation decisions and edit them according to their preferences, while also providing other ecological variables of interest. Furthermore, this work provides an in-depth assessment of the taxonomic and sampling coverage of the database at the European level, complemented by analyses of key structural properties in plant-pollinator networks. We hope this database can help researchers to: 1) identify taxonomic, ecological, and geographical gaps of knowledge on plant-pollinator interactions; and 2) explore the impacts of global change on plant-pollinator networks to guide future conservation planning for both plant and pollinator species.

Main Types of Variables Included: EuPPollNet contains 1,144,369 interactions between plants and pollinators from 1,147 distinct locations (i.e., plant-pollinator networks), which belong to 51 different studies distributed across 17 European countries. In addition, information about sampling methodology, habitat type, bio-climatic region, and further taxonomic rank information for both plant and pollinator species are also provided.

Spatial location and grain: The database contains 1,147 different sampling locations from natural and anthropogenic habitats that fall in 5 different bio-climatic regions. All records are geo-referenced and presented in the World Geodetic System 1984 (WGS84).

Time period and grain: Species interaction data was recorded between 2004 and 2021. All records are time-referenced and most of the studies documented interactions in a single flowering season (68.63%).

Major taxa and level of measurement: The database contains interaction data at the species level for 94.72% of the records, including a total of 1,353 plant and 2,065 pollinator species. The database covers 5.34% of the European species of flowering plants, 33.82% of bees, 26.21% of butterflies, and 33.08% of syrphids species at the European level.

Software format: The database was built with R software and is stored as “.rds” and “.csv” formats. The construction of the database is fully reproducible and can be accessed at the following PERMANENT LINKS.

KEYWORDS

plant-pollinator networks, plant-pollinator interactions, flowering plants, Angiosperms, pollinators, nestedness, connectance

1 | INTRODUCTION

The interaction between plants and pollinators is one of the most well-documented mutualisms on Earth. Plant-pollinator interactions involve a great diversity of species, largely attributed to their coevolutionary history (Ollerton 2017), and are critically important for terrestrial biodiversity and the human economy. The synergistic effects of climate change with other global change pressures are threatening worldwide biodiversity (Sala et al. 2000; Bellard et al. 2014), including plant and pollinator species as well as their interactions (Goulson et al. 2015; Settele, Bishop, and Potts 2016; Eichenberg et al. 2021). Under this scenario, the increasing availability of biodiversity data plays a major role in our ecological understanding of species and guiding conservation planning (Heberling et al. 2021). However, our knowledge of plant and pollinator species and their network of interactions still exhibits major temporal, spatial and taxonomic biases (Archer et al. 2014; Troia and McManamay 2016; Poisot et al. 2021; Marshall et al. 2024), limiting our ability to effectively protect their biodiversity.

Europe is one of the continents with a larger amount of available biodiversity data (Proença et al. 2017), yet still exhibits major gaps (Wetzel et al. 2018; Bennett et al. 2018). While species checklists need to be treated carefully, especially at a macro-ecological scale (Grenié et al. 2023), the growing number of European plant and pollinator checklists along with occurrence data, is setting a foundation for the conservation of its flora and their pollinators. However, species richness is just one component of biodiversity and documenting the interaction between plants and pollinators is essential for understanding the fate of the species within an ecosystem (Jordano 2016). Numerous works have studied plant-pollinator interactions in the last decades, originating thousands of plant-pollinator interaction networks worldwide. Several initiatives have tried to integrate plant-pollinator interaction data into databases such as *Mangal* (Poisot et al. 2016) or *GloBI* (Poelen, Simons, and Mungall 2014), resulting in numerous large scale comparative analyses that have enhanced our understanding of the ecology of plants and pollinators (e.g., European wild bee data trends; Marshall et al. 2024). Despite all these resources, Europe lacks accessible harmonized plant-pollinator interaction data that allow researchers to evaluate the state of the art of plant-pollinator interactions at European level, which will guide research efforts, conservation planning and will set a foundation for future global change research.

The interactions between different plant and pollinator species within a community form complex networks. Macro-ecological analyses of the topology of these networks have revealed common properties across them, such as truncated power-law degree distributions (Jordano 1987) or nestedness (Bascompte et al. 2003). In addition, this type of large scale analyses can help understanding landscape level processes that cannot be explored at the community level, such as ecological patterns across geographic regions (Olesen and Jordano 2002; Traveset et al. 2016) or environmental gradients (Ramos-Jiliberto et al. 2010; Rech et al. 2016; Saunders et al. 2023). Although macro-ecological approaches make significant contributions to knowledge, they tend to be rare, and are strongly influenced by the spatio-temporal availability and nature of the data (Burkle and Alarcón 2011; Trøjelsgaard and Olesen 2013). For instance,

plant-pollinator studies tend to differ in sampling effort and methodology which can impact the structure of the resulting plant-pollinator networks (Gibson et al. 2011; Jordano 2016). Further, despite little sampling effort could capture most of the relevant functional species (Hegland et al. 2010), most plant-pollinator networks have unobserved interactions because undersampling (Olesen et al. 2011; Chacoff et al. 2012). Thus, the lack of strong spatio-temporal coverage, along with the current intrinsic limitations of sampling plant-pollinator networks, highlight the need of keep gathering and integrating informative species interaction data to properly unravel the different ecological processes that shape plant-pollinator interactions at large scales.

Here, we present the EuPPollNet database, an acronym derived from European plant-pollinator networks, which contains harmonized information on interaction data of plants and pollinators at European level. The primary focus of the pollinator taxonomic groups is on the main orders of entomofauna that visit and pollinate flowering plants in Europe. These include insect species from the orders Hymenoptera, Diptera, Lepidoptera, and Coleoptera (Potts et al. 2015), which comprise almost the totality of recorded interactions in the EuPPollNet database (99.88%). To understand the scope of the database, we examine the taxonomic and sampling coverage of the different plant and pollinator species at European level with the help of the most up-to-date species checklists and rarefaction analyses. In addition, we investigate how two key structural metrics of plant-pollinator networks, such as connectance and nestedness, change across the latitudinal range of studies and bioclimatic regions. Finally, we also examine whether plant-pollinator networks fulfill the expectation of being more nested than null expectations. EuPPollNet aims to cover a wide range of taxonomic groups and habitats, while also providing other variables of interest that allow better control for the ecological context and sampling methods. In addition, EuPPollNet offers a transparent and accessible workflow of its data management and species harmonization that allows it to be reused and keep building on it over time. This database provides a large number of community-level networks with curated and harmonized data, distinguishing it from other currently available resources that contain plant-pollinator interactions. We expect that this database will help to evaluate macro-ecological processes and current gaps of plant-pollinator interactions at European level.

2 | METHODS

Data acquisition

This database is the result of one of the working packages of the European project Safeguard. The EuPPollNet database includes published and unpublished studies compiled initially by a wide number of researchers and institutions within the European continent. Data acquisition followed a non-systematic approach. First, data was directly asked to members of the Safeguard project and then, the request was extended to data owners outside of the project. These other studies were identified by direct communication with other colleagues and by directly

searching studies on Google Scholar of under-represented regions within the database. To keep high quality standards that will allow robust future ecological research, we only included studies meeting the following criteria: 1) studies that contained time- and geo-referenced records of plant-pollinator interactions; and 2) studies with phyto-centric plant-pollinator networks with quantitative visitation data.

Dataset description

The database contains 51 independent published and unpublished studies conducted during the time period 2004 - 2021 on 17 different countries (**Figure 1a** and **Figure 1b**). The different studies differ in sampling effort and methodology, although most studies took place within a single flowering season (68.63%), sampled a given location an average of 7.22 days, and documented interactions mostly by using transects as sampling method (62.75%). The database includes a total of 1,144,369 distinct interactions, considering interaction as the contact of a given pollinator to the reproductive structure of a particular plant. Most of the pollinator species in the database belong to the orders Hymenoptera and Diptera, each comprising approximately 1,000 species. However, the majority of plant-pollinator interactions are from Hymenoptera species (91.04%; **Figure 1c**). Notably, *Apis mellifera* represents 70.68% of the total interaction records from the database and an average of 31.00% of the total interactions per network.



Figure 1. (a) Approximate locations of the studies of the EuPPollNet database showing the total number of pollinator (i.e., orange heptagon) and plant (i.e., green circles) species per study. The sizes of these shapes are proportional to the respective species counts. For visualization purposes, we have focused only on the European region with studies and selected a single location per study. (b) Number of studies by year within the database. (c) Number of species and interactions across the 4 main pollinator orders at European level.

Data structure

The EuPPollNet database is available in both .csv and .rds formats. The .csv file contains data in long format, while the .rds file stores the data in a list structure, with networks organized within each study. The file contains a total of 30 columns that depict where and when the plant-pollinator interaction was observed (**Table 1**). To provide a standard unit of plant-pollinator interaction across studies, each row depicts a single interaction between a plant and a pollinator species. To build a plant-pollinator network matrix at a site level, authors only need to group interactions by plant and pollinator species, site and study. Information about habitat type and bioclimatic region are also provided in this file. In addition, the interaction dataset includes a column that describes the presence or absence of floral counts for each study. The flower count data is provided in a separate file given that one third of studies lack floral measurements and that the units differ greatly across studies. Finally, metadata at the study level is provided, including information about the authors, digital object identifier if available, sampling time and taxonomic coverage of the main pollinator groups for each study.

Sampling coverage

The completeness of the EuPPollNet database was evaluated by exploring the rarefied accumulation curves of plant and pollinator species and their interactions across the different sampling sites. In addition, an outstanding question in ecology is how many pollinators are required to pollinate flowering plants (Kleijn et al. 2015). To that end, we also calculated the accumulation curve of pollinator species with increasing number plant species. The rarefied and extrapolated sampling curves were obtained with the help of the **iNEXT** package (Hsieh and Chao 2016). This was complemented with 100 random accumulation curves obtained programmatically to guide the visualization of the different rarefied curves.

Taxonomic harmonization

All plant and pollinator species names were checked and standardized according to large scale taxonomic databases. To ensure reproducibility of the workflow, we have conducted this harmonization in R with **rgbif** (Chamberlain, Oldoni, and Waller 2022) as pivotal package to check for species names and retrieve further taxonomic information (i.e., phylum, order, family and genus) from the Global Biodiversity Information Facility (GBIF). The protocol for plants and pollinators is similar but slightly different given the availability of the different taxonomic resources. For transparency, we have included in the database the old species name, the new assigned name, and, if the name of the species is uncertain (e.g., species complex or species alike).

For plants: (i) we initially verified the exact matches against the GBIF species checklist; (ii) selected unmatched cases and fix orthographic errors; (iii) retrieved again taxonomic information for those unmatched cases, evaluated accuracy of fuzzing matching and programmatically fixed

Table 1. Column names and their descriptions within the EuPPollNet database.

| Variable | Description |
|-----------------------------|---|
| Study_id | Identifier of the study |
| Network_id | Identifier of a site sampled within a study |
| Sampling_method | Type of plant-pollinator sampling |
| Authors_habitat | Type of habitat as described by the authors |
| EuPPollNet_habitat | Type of habitat homogenized across studies |
| Bioregion | European biogeographical regions |
| Country | Country where the plant-pollinator interaction was observed |
| Locality | Locality where the plant-pollinator interaction was observed |
| Latitude-Longitude | Coordinates of the observed interaction in decimal degrees |
| Date | Year, month and day when the observation took place |
| Interaction | Number of interactions. By default is 1 as interactions are provided ungrouped |
| Plant_old_name | Plant species name given by the authors |
| Plant_accepted_name | Harmonized plant species name in the database |
| Plant_rank | Taxonomic rank of the observation |
| Plant_order | Order taxonomic rank of the observed plant species |
| Plant_family | Family taxonomic rank of the observed plant species |
| Plant_genus | Genus taxonomic rank of the observed plant species |
| Plant_unsure_id | Category to indicate if the plant species name is unsure (Yes) or not (No) |
| Plant_uncertainty_type | If the name is unsure, type of species uncertainty is provided |
| Pollinator_old_name | Pollinator species name given by the authors |
| Pollinator_accepted_name | Harmonized pollinator species name in the database |
| Pollinator_rank | Taxonomic rank of the observation |
| Pollinator_order | Order taxonomic rank of the observed pollinator species |
| Pollinator_family | Family taxonomic rank of the observed pollinator species |
| Pollinator_genus | Genus taxonomic rank of the observed pollinator species |
| Pollinator_unsure_id | Category to indicate if the pollinator species name is unsure (Yes) or not (No) |
| Pollinator_uncertainty_type | If the name is unsure, type of species uncertainty is provided |
| Flower_data | Floral data availability (Yes) or (No) |
| Flower_data_merger | Column to merge floral data with the interaction dataset |

records that are still not found; (iv) finally, we used the World Flora Taxonomic Backbone (Govaerts et al. 2021; WFO, July 7, 2022) as the ultimate filter for taxonomic information as we used it to calculate the plant taxonomic coverage of our database.

For pollinators: (i) we first created a checklist of species names for the most representative pollinator groups at European level by combining the recently published checklists of bees (Ghisbain et al. 2023), syrphids (Kočić et al. 2023) and butterflies (Wiemers et al. 2018); (ii) then, we compared pollinator species names against the checklist and recovered some unmatched cases with restrictive fuzzy matching by using **stringdist** package (Van der Loo et al. 2014); (iii) we programmatically fixed unmatched records when necessary and retrieved the taxonomic information for all species from GBIF; (iv) we fixed the non-found cases in the GBIF checklist and made sure that all species names from bees, syrphids and butterflies were named according to their respective species checklists.

Taxonomic coverage

To assess the completeness of plant and pollinator species in the EuPPollNet database at European level, we used the aforementioned checklists for plants and pollinators. Specifically for plants, we refined the checklist to include only European flowering plants and excluded taxonomic groups not associated with biotic pollination. We did this by first excluding the families considered to have exclusively a wind pollination mode (see Culley, Weller, and Sakai 2002), and then by filtering out the genera with wind or non-biotic pollination from families that exhibit both biotic and non-biotic pollination modes. Additionally, we manually included some exotic species and added unresolved species names that were not present in the accepted names of the checklist at the current version of usage. For pollinators, we compared only the taxonomic coverage of bees, syrphids and butterflies by using their species checklists at European level. Finally, to evaluate if the presence-absence of interaction records for bees and flowering plants follows a phylogenetic pattern within the database, we calculated its phylogenetic signal at genus and family level, respectively. The phylogenetic signal was calculated by using the *phylosig* function from the **phytools** package (Revell 2012). We extracted the phylogenetic information for bees from a genus level phylogeny (Hedtke, Patiny, and Danforth 2013) and processed it with the help of the packages **ape** (Paradis et al. 2019), **MCMCglmm** (Hadfield 2010) and **phytools**; and for plants the phylogenetic tree was obtained from a species level PLANT phylogeny (Smith and Brown 2018) with the help of **rtree** package (Li 2023).

Habitat type and bioclimatic region

The different sites per study were described with a habitat type by the authors. As these habitats are not standardized across studies, they were standardized with the additional help of land cover information and visual checks on current satellite imagery. For each georeferenced site, the land cover information was extracted from Corine Land Cover (CLC, version 2018) with the help of the **Terra** package (Hijmans et al. 2022). Based on the habitat classification

from the authors and the CLC classification, we created habitat categories that intend to summarize the diversity of habitats in the EuPPollNet database (see habitat type definition in supplementary material). These categories allow a quick comparison and understanding of the habitat types from the database. However, we advise authors to revise this classification if they intend to rely on this field for their analyses as this is a non-fully objective process. Moreover, Europe is characterized by a great variety of environmental conditions that harbor different biota. Thus, to allow authors to explore set of studies that share similar environmental conditions and species, we assigned to each site a biogeographical region. The biogeographical regions were downloaded from the European Environment Agency (version 2016) and were matched to the different sites with the help of a spatial joint from the **sf** package (Pebesma et al. 2018).

Network analyses

To get a general overview of some of the main aspects of these set of plant-pollinator networks, we evaluated how connectance and nestedness change across the latitudinal range of studies and bioclimatic regions. We implemented linear models, with the network metrics as the response variable, and both latitude and bioclimatic region as predictors in two separate models. Finally, with the help of a Tukey test from the package *emmeans*, we evaluated if nestedness and connectance differ statistically across bioclimatic regions. In addition, we explored the association between connectance and nestedness with the number of species per network (i.e., geometric mean of plants and pollinators) by conducting Kendall rank correlation coefficient. Finally, we evaluated if the different networks are more or less nested than expected by chance. Importantly, we implemented two different metrics of nestedness given the different analytical objectives. Common metrics of nestedness are highly correlated to network size, limiting our ability to compare this metric across networks. To avoid this, we implemented a normalised version of nestedness (NODFc) that corrects by connectance and the number of species (Song, Rohr, and Saavedra 2017). This metric was calculated with the help of the **maxnodf** package (Hoeppke and Simmons 2021), which employs a computational efficient approach to calculate it. However, to compare if networks are more or less nested than expected by chance, we employed the traditional z-score approach with the widely used nestedness metric (NODF) from Almeida-Neto et al. (2008), as this approach allow us to compare our results with previous published nestedness analyses in plant-pollinator networks and only compares each unique network against their randomized versions. We calculated 100 null models for each network with the help of the *vaznull* function from the package **Bipartite** (Dormann, Gruber, and Fründ 2008) that implements the null model from Vázquez et al. (2005). These null networks have the same connectance and number of plant and pollinator species as the empirical ones, but different marginal totals. In other words, interactions are reshuffled with almost equal probabilities to their relative abundances. Connectance was estimated for each network by using the function *networklevel* from **Bipartite**.

3 | RESULTS

Sampling coverage

The estimated sampling coverage of plant and pollinator species within the database across the different sampling sites is approximately 97% for both taxonomic groups. This suggests that the rarefied accumulation curves of both plant and pollinator species exhibit already a “quasi-asymptotic” growth of species richness by considering the current number of sampling sites or networks (**Figure 2a-2b**). The predicted observed species richness by doubling the sampling effort on the already sampled habitat types within the database will only increase pollinator richness by 24.13% and plant richness by 21.14%. However, the sampling coverage of interactions is 74.17%, and by doubling the sampling effort the predicted number of unique interactions recorded will have approximately a twofold increase (54.83%; **Figure 2c**). Similarly, to the plant and pollinator species accumulation curves across sampling sites, when we consider the accumulated pollinator richness across sampled plant species, this curve also shows a “quasi-asymptotic” growth with a sampling coverage value of 96.66%. The predicted recorded pollinator species by doubling the number of plants sampled is expected to increase by 21.90% (**Figure 2d**). In addition, when we explore how common or rare are plant and pollinator species across all sampling sites, we find that the majority of plant and pollinator species tend to be regionally specific, and only a minor portion of them is being shared across a broad range of sampling sites (**Figures 2e-2f**). For instance, most plants (84.70%) and pollinators (87.07%) are exclusively found in less than 1% of sampling sites; and the most common plant (*Trifolium pratense*) and pollinator (*Bombus pascuorum*) species are only found in 37.32% and 64.80% of sampling sites, respectively.

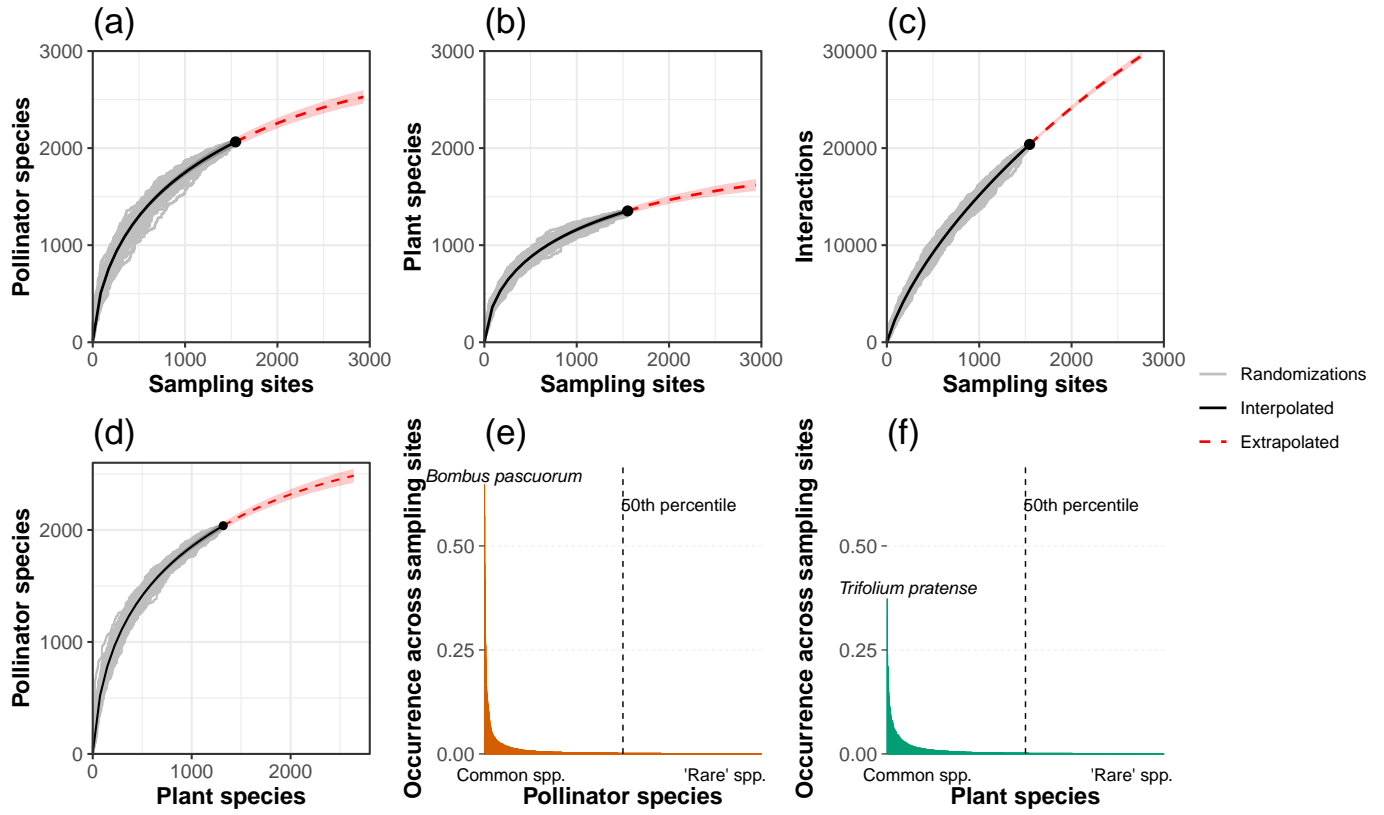


Figure 2. Graphs (a-b-c) indicate the accumulation curves for pollinators, plants, and their interactions across sampling sites. Grey solid lines represent 100 randomized accumulation curves, the black solid lines represent the interpolated curve (i.e., the mean across curves), and the red dashed lines illustrate the extrapolated curve for approximately 3000 sampling sites. The solid black points indicate the number of species and interactions contained in the database. Graph (d) shows the accumulation curve of pollinator species across an increasing number of plant species. This last graph uses the same color and shape structure as the ones in the top panel. Graphs (e-f) indicate the percentage of occurrence (i.e., incidence) of plant and pollinator species across sampling sites. Species on the left (i.e., common) are found in many sampling sites, while species on the right (i.e., rare) are found in few or only a single sampling site. Note that indeed *Apis mellifera* is the most common pollinator but was excluded from this visualization.

Taxonomic coverage

The database contains a total of 2,065 pollinator and 1,353 plant species. The coverage of the main pollinators groups occurring in Europe is 33.82% for bees, 33.08% for syrphids and 26.21% for butterflies (see **Figure S1** for coverage at the family level for bees and butterflies, and at the subfamily level for syrphids). Bees constitute 90.58% of the interactions of the database, and 78.03% of the interactions when excluding honey bees. Within the database, 83.82% of bee genera have at least one species with interaction records, and the average coverage at the bee genus level is 36.99% (**Figure 3**). In addition, the presence or absence of interactions records for bees seems not to follow any sort of phylogenetic pattern ($\lambda = 0.07$; $P = 0.65$). The database coverage of all flowering plant species occurring in Europe is 5.34% (**Figure 4**), with an average coverage of 9.04% at the plant family level. Approximately, half of the plant families have at least one species with interaction records (52.56%), and the presence or absence of interaction data for the different plant species also seems not to follow a statistically relevant phylogenetic pattern ($\lambda = 0.26$; $P = 0.07$).

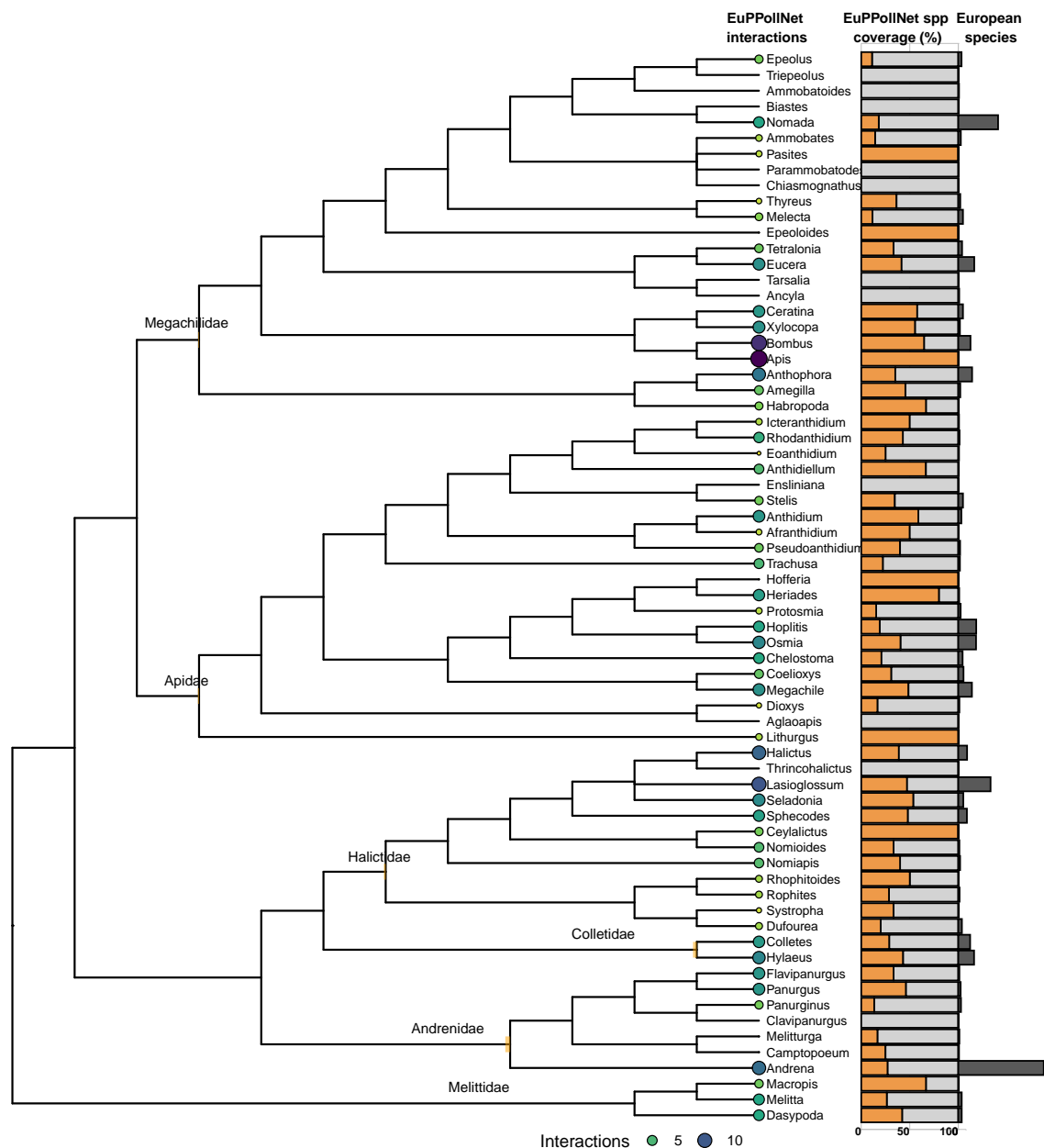


Figure 3. Phylogenetic representation of bee genera in the EuPPollNet database within the context of all European bee genera. The number of interactions recorded per genus in the database is illustrated using circles, with their sizes proportional to the number of interactions on a logarithmic scale. Additionally, a gradient of colors ranging from yellow to dark purple aids in this visualization. The coverage of species recorded in EuPPollNet per genus at the European level is depicted with orange and light grey bars, representing the percentage of species included and not included in the database, respectively, out of the total number of bee species per genus at the European level. Grey bars indicate the total number of species per genus at the European level.

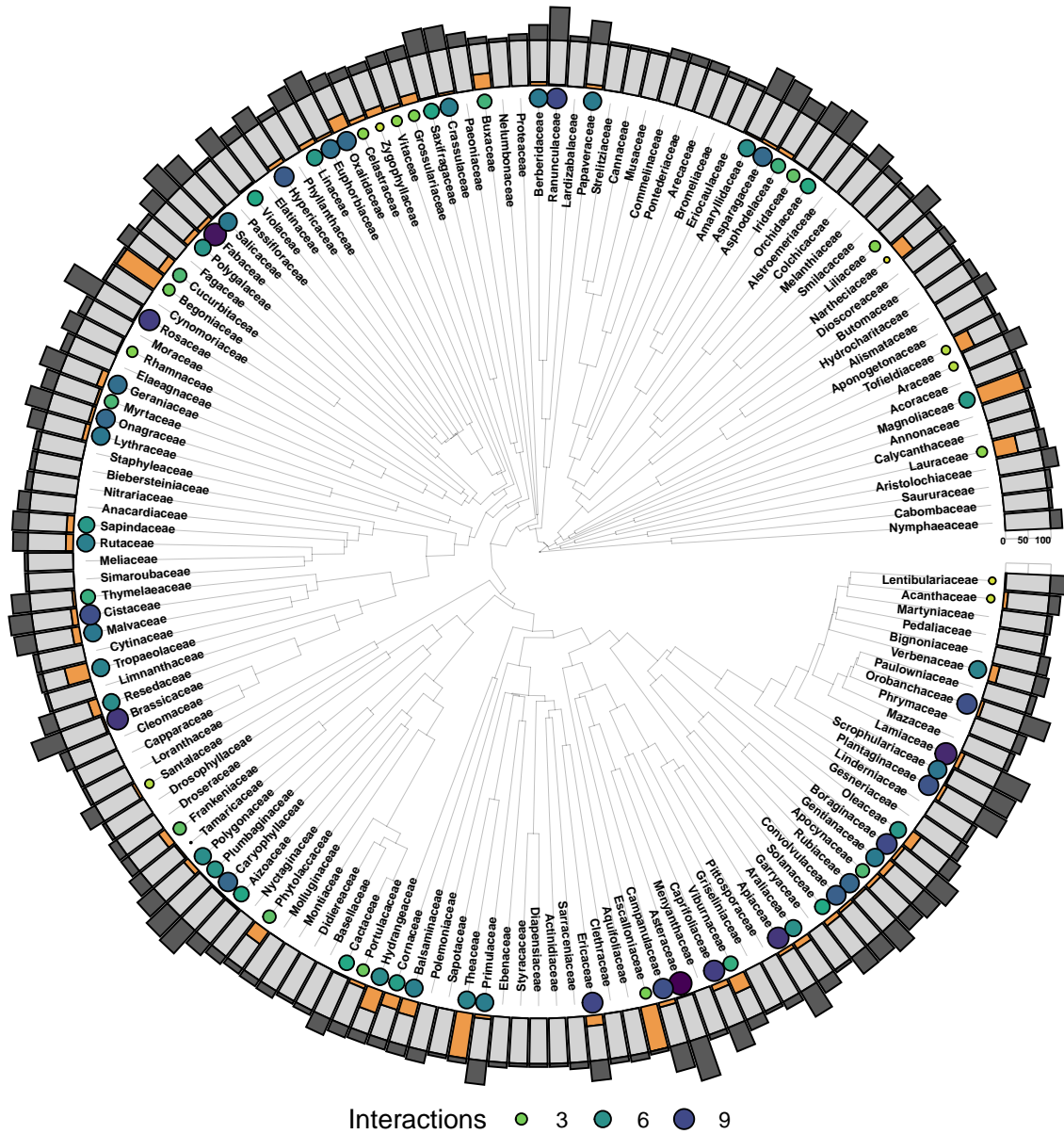


Figure 4. Phylogenetic representation of the plant families in the EuPPollNet database within the context of all flowering plant families occurring in Europe. The number of interactions recorded per genus in the database is illustrated using circles, with their sizes proportional to the number of interactions on a logarithmic scale. Additionally, a gradient of colors ranging from yellow to dark purple aids in this visualization. The coverage of species recorded in EuPPollNet per genus at the European level is depicted with orange and light grey bars, representing the percentage of species included and not included in the database, respectively, out of the total number of flowering plant species per family at the European level. Grey bars indicate the total number of species per family at the European level on logarithmic scale.

Habitat type and bioclimatic region

The proportion of the major pollinator orders within the database differed across habitats and bioclimatic regions (**Figure 5**). As expected, Hymenoptera was the main taxonomic order on the majority of habitats, exceeded only by Diptera on the habitat categories of riparian vegetation, moors and heathland, and alpine grasslands. Overall, the proportion of Lepidoptera and Coleoptera was low across all habitats but Coleoptera showed a notably increase in sclerophyllous vegetation and beaches, dunes and sands habitat categories. Similar patterns were observed when exploring the pollinator proportions by bioclimatic region. In this particular case, Hymenoptera was the predominant order in all bioclimatic regions with Diptera taking more importance in Alpine and Atlantic regions. Again, Lepidoptera shows low proportions across all bioclimatic regions and Coleoptera seems to be only relevant in Mediterranean regions at European level. Notably, the number of studies (**Figure 5**) and sampling sites (**Figure S2**) also differs across habitats and bioclimatic regions. The habitats sampled by a higher number of studies in the database are pastures (27), semi-natural grassland (11) and sclerophyllous vegetation (9). However, the habitats that contain a higher number of sampling sites are pastures (589), agricultural margins (432) and agricultural land (141). The bioclimatic regions with a higher number of studies are continental (22), atlantic (13) and mediterranean (12); and the ones that contain a higher number of sampling sites are atlantic (459), continental (454) and boreal (439).

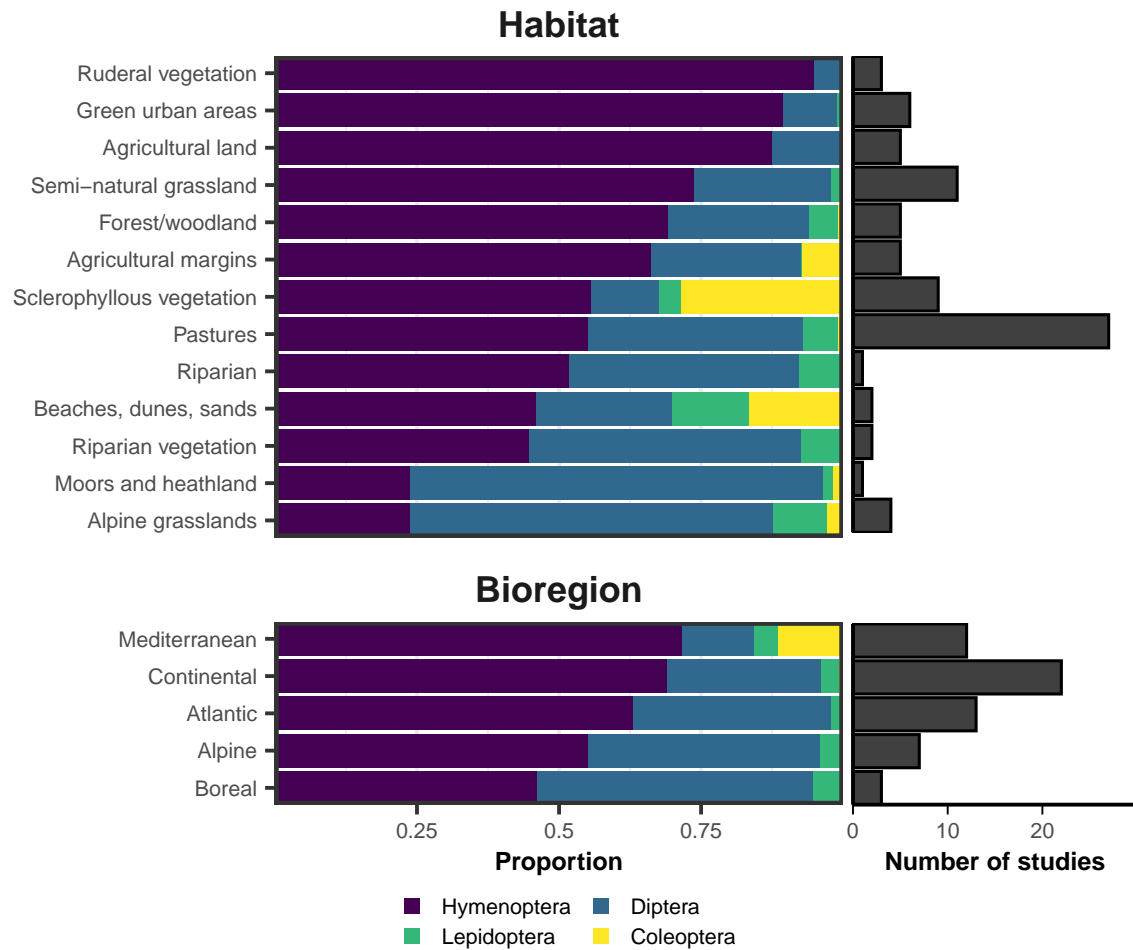


Figure 5. Proportion of the major pollinator orders by habitat types and bioclimatic regions in the EuPPollNet database. The orders, from left to right, include Hymenoptera, Diptera, Lepidoptera and Coleoptera. The horizontal barplot on the right indicates the number of studies that were conducted on each habitat type or bioclimatic region. Note that a single study can contribute to more than one habitat or bioclimatic region. Areas with a greater number of studies are more likely to depict accurate proportions of the different pollinator orders in those systems.

Network properties

Connectance and nestedness did not show major changes across the latitudinal range of studies (connectance: $R^2 = 0$, $P = 0.14$, **Figure 6a**; NODFc: $R^2 = 0.01$, $P = 0.02$, **Figure 6b**). However, differences were found for connectance but not nestedness across bioclimatic regions. Specifically, the alpine region differed statistically in connectance to the other bioclimatic regions (Tukey test $P < 0.01$). In addition, the atlantic and continental bioclimatic regions also showed statistical meaningful differences in connectance (Tukey test $P < 0.01$). Connectance values ranged between 0.03 to 0.4 ($\bar{x} = 0.14$) and as expected, followed a negative exponential relationship with the geometric mean of plant and pollinator species per network (Kendall $\tau = -0.75$, $P < 0.01$; **Figure 6c**). Nestedness values (NODFc) ranged between 1.34 to 7.94 ($\bar{x} = 2.81$), and as expected for this metric, were independent of the mean number of species (Kendall $\tau = -0.06$, $P = 0.03$; **Figure S3**). Finally, we found that only 31.62% of networks were statistically more nested than expected by chance, with 68.38% showing no statistical difference, and none being less nested than null expectations (**Figure 6d**).

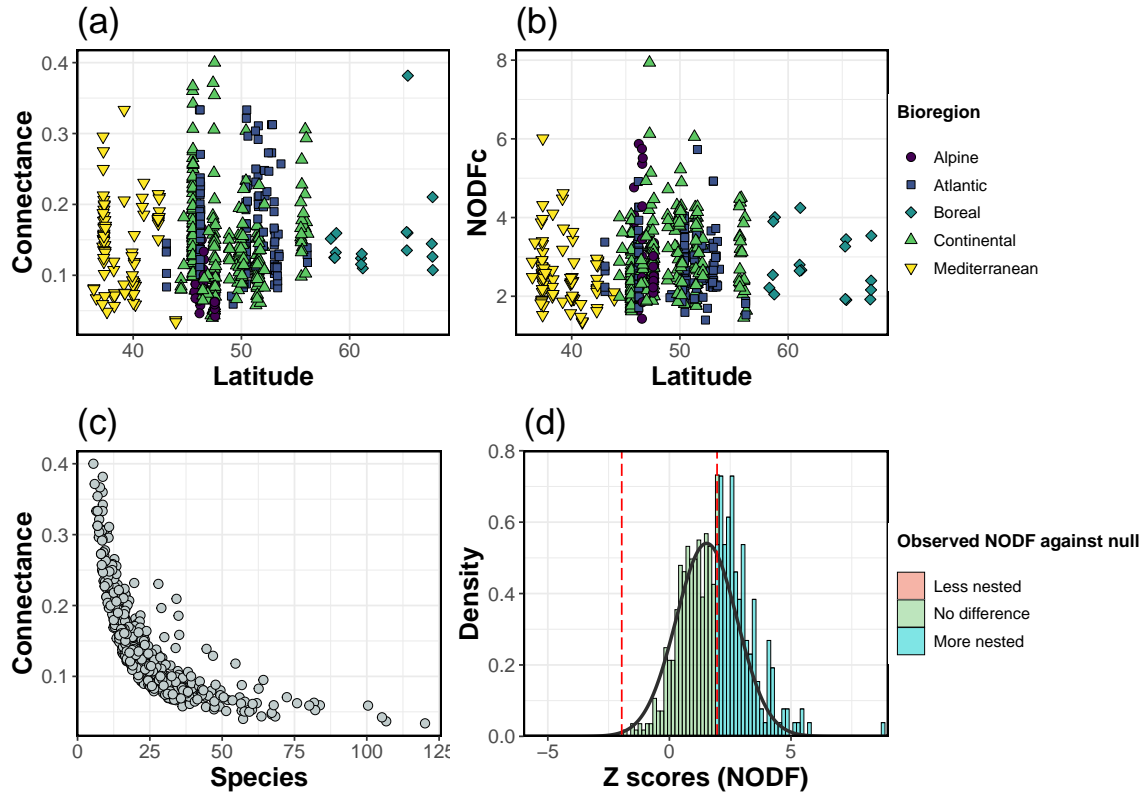


Figure 6. Graphs (a-b) indicate network connectance and nestedness across the latitudinal range of the studies from the database. In addition, the bioclimatic region where each network is located is indicated with points of different shapes and colours. Graph (c) shows the association between network connectance and the geometric mean of plant and pollinator species per network. Graph (d) shows the distribution of z-scores when comparing the nestedness from the empirical networks with their randomised counterparts (100 null models for each network). The vertical red dashed lines represent the z critical value for a two tailed test with $\alpha = 0.05$. Z-scores to the left of the first vertical red dashed line indicate that networks are less nested than expected by chance (red), those between the two dashed lines indicate no statistical difference from random expectations (green), and those to the right indicate that networks are more nested than expected by chance (blue).

3 | DISCUSSION

Things to do:

-Intro: Somewhere, but maybe later, we should mention what we add. 1) Way more networks than those on Mangal or Web of life (which you should cite). 2) curated metadata and harmonization, 3) Community level data and not only isolated interactions as in GloBI.

Although Europe...

Geographic and taxonomic biases! Supporting reference: <https://doi.org/10.1111/ecog.05926>

Networks, ref:

<https://www.biorxiv.org/content/10.1101/2020.04.02.021691v1.abstract>

Contextualise European diversity of plant and pollinator species with global biodiversity

References

- Almeida-Neto, Mário, Paulo Guimaraes, Paulo R Guimaraes Jr, Rafael D Loyola, and Werner Ulrich. 2008. "A Consistent Metric for Nestedness Analysis in Ecological Systems: Reconciling Concept and Measurement." *Oikos* 117 (8): 1227–39.
- Archer, C Ruth, Christian Walter Werner Pirk, Luísa G Carvalheiro, and Sue W Nicolson. 2014. "Economic and Ecological Implications of Geographic Bias in Pollinator Ecology in the Light of Pollinator Declines." *Oikos* 123 (4): 401–7.
- Bascompte, Jordi, Pedro Jordano, Carlos J Melián, and Jens M Olesen. 2003. "The Nested Assembly of Plant–Animal Mutualistic Networks." *Proceedings of the National Academy of Sciences* 100 (16): 9383–87.
- Bellard, Céline, Camille Leclerc, Boris Leroy, Michel Bakkenes, Samuel Veloz, Wilfried Thuiller, and Franck Courchamp. 2014. "Vulnerability of Biodiversity Hotspots to Global Change." *Global Ecology and Biogeography* 23 (12): 1376–86.
- Bennett, Joanne M, Amibeth Thompson, Irina Goia, Reinart Feldmann, Valentin Ştefan, Ana Bogdan, Demetra Rakosy, et al. 2018. "A Review of European Studies on Pollination Networks and Pollen Limitation, and a Case Study Designed to Fill in a Gap." *AoB Plants* 10 (6): ply068.
- Burkle, Laura A, and Ruben Alarcón. 2011. "The Future of Plant–Pollinator Diversity: Understanding Interaction Networks Across Time, Space, and Global Change." *American Journal of Botany* 98 (3): 528–38.
- Chacoff, Natacha P, Diego P Vázquez, Silvia B Lomáscolo, Erica L Stevani, Jimena Dorado, and Benigno Padrón. 2012. "Evaluating Sampling Completeness in a Desert Plant–Pollinator Network." *Journal of Animal Ecology* 81 (1): 190–200.
- Chamberlain, Scott, Damiano Oldoni, and John Waller. 2022. "Rgbiif: Interface to the Global Biodiversity Information Facility API."
- Culley, Theresa M, Stephen G Weller, and Ann K Sakai. 2002. "The Evolution of Wind Pollination in Angiosperms." *Trends in Ecology & Evolution* 17 (8): 361–69.
- Dormann, Carsten F, Bernd Gruber, and Jochen Fründ. 2008. "Introducing the Bipartite Package: Analysing Ecological Networks." *Interaction* 1 (0.2413793): 8–11.
- Eichenberg, David, Diana E Bowler, Aletta Bonn, Helge Bruelheide, Volker Grescho, David Harter, Ute Jandt, Rudolf May, Marten Winter, and Florian Jansen. 2021. "Widespread Decline in Central European Plant Diversity Across Six Decades." *Global Change Biology* 27 (5): 1097–1110.
- Ghisbain, Guillaume, Paolo Rosa, Petr Bogusch, Simone Flaminio, ROMAIN LE DIVELEC, Achik Dorchin, Max Kasperek, et al. 2023. "The New Annotated Checklist of the Wild Bees of Europe (Hymenoptera: Anthophila)." *Zootaxa* 5327 (1): 1–147.
- Gibson, Rachel H, Ben Knott, Tim Eberlein, and Jane Memmott. 2011. "Sampling Method Influences the Structure of Plant–Pollinator Networks." *Oikos* 120 (6): 822–31.
- Goulson, Dave, Elizabeth Nicholls, Cristina Botías, and Ellen L Rotheray. 2015. "Bee Declines Driven by Combined Stress from Parasites, Pesticides, and Lack of Flowers." *Science* 347 (6229): 1255957.
- Govaerts, Rafaël, Eimear Nic Lughadha, Nicholas Black, Robert Turner, and Alan Paton. 2021.

- “The World Checklist of Vascular Plants, a Continuously Updated Resource for Exploring Global Plant Diversity.” *Scientific Data* 8 (1): 215.
- Grenié, Matthias, Emilio Berti, Juan Carvajal-Quintero, Gala Mona Louise Dädlow, Alban Sagouis, and Marten Winter. 2023. “Harmonizing Taxon Names in Biodiversity Data: A Review of Tools, Databases and Best Practices.” *Methods in Ecology and Evolution* 14 (1): 12–25.
- Hadfield, Jarrod D. 2010. “MCMC Methods for Multi-Response Generalized Linear Mixed Models: The MCMCglmm r Package.” *Journal of Statistical Software* 33: 1–22.
- Heberling, J Mason, Joseph T Miller, Daniel Noesgaard, Scott B Weingart, and Dmitry Schigel. 2021. “Data Integration Enables Global Biodiversity Synthesis.” *Proceedings of the National Academy of Sciences* 118 (6): e2018093118.
- Hedtke, Shannon M, Sébastien Patiny, and Bryan N Danforth. 2013. “The Bee Tree of Life: A Supermatrix Approach to Apoid Phylogeny and Biogeography.” *BMC Evolutionary Biology* 13: 1–13.
- Hegland, Stein Joar, Jennifer Dunne, Anders Nielsen, and Jane Memmott. 2010. “How to Monitor Ecological Communities Cost-Efficiently: The Example of Plant–Pollinator Networks.” *Biological Conservation* 143 (9): 2092–2101.
- Hijmans, Robert J, Roger Bivand, Karl Forner, Jeroen Ooms, Edzer Pebesma, and Michael D Sumner. 2022. “Package ‘Terra’.”
- Hoeppke, Christoph, and Benno I Simmons. 2021. “Maxnodf: An r Package for Fair and Fast Comparisons of Nestedness Between Networks.” *Methods in Ecology and Evolution* 12 (4): 580–85.
- Hsieh, TC, and Anne Chao. 2016. “iNEXT: An r Package for Rarefaction and Extrapolation of Species Diversity (h Ill Numbers).” *Methods in Ecology and Evolution* 7 (12): 1451–56.
- Jordano, Pedro. 1987. “Patterns of Mutualistic Interactions in Pollination and Seed Dispersal: Connectance, Dependence Asymmetries, and Coevolution.” *The American Naturalist* 129 (5): 657–77.
- . 2016. “Sampling Networks of Ecological Interactions.” *Functional Ecology* 30 (12): 1883–93.
- Kleijn, David, Rachael Winfree, Ignasi Bartomeus, Luísa G Carvalheiro, Mickaël Henry, Rufus Isaacs, Alexandra-Maria Klein, et al. 2015. “Delivery of Crop Pollination Services Is an Insufficient Argument for Wild Pollinator Conservation.” *Nature Communications* 6 (1): 7414.
- Kočić, Anja, Ante Vujić, Tamara Tot, Marina Janoviké Milosavljevuć, and Maarten De Groot. 2023. “An Updated Checklist of the Hoverflies (Diptera: Syrphidae) of Slovenia.” *Zootaxa* 5297 (2): 189–227.
- Li, Daijiang. 2023. “Rtrees: An r Package to Assemble Phylogenetic Trees from Megatrees.” *Ecography* 2023 (7): e06643.
- Marshall, Leon, Nicolas Leclercq, Luísa G Carvalheiro, Holger H Dathe, Bernhard Jacobi, Michael Kuhlmann, Simon G Potts, Pierre Rasmont, Stuart PM Roberts, and Nicolas J Vereecken. 2024. “Understanding and Addressing Shortfalls in European Wild Bee Data.” *Biological Conservation* 290: 110455.
- Olesen, Jens M, Jordi Bascompte, Yoko L Dupont, Heidi Elberling, Claus Rasmussen, and

- Pedro Jordano. 2011. “Missing and Forbidden Links in Mutualistic Networks.” *Proceedings of the Royal Society B: Biological Sciences* 278 (1706): 725–32.
- Olesen, Jens M, and Pedro Jordano. 2002. “Geographic Patterns in Plant–Pollinator Mutualistic Networks.” *Ecology* 83 (9): 2416–24.
- Ollerton, Jeff. 2017. “Pollinator Diversity: Distribution, Ecological Function, and Conservation.” *Annual Review of Ecology, Evolution, and Systematics* 48: 353–76.
- Paradis, Emmanuel, Simon Blomberg, Ben Bolker, Joseph Brown, Julien Claude, Hoa Sien Cuong, Richard Desper, and Gilles Didier. 2019. “Package ‘Ape’.” *Analyses of Phylogenetics and Evolution, Version 2* (4): 47.
- Pebesma, Edzer J et al. 2018. “Simple Features for r: Standardized Support for Spatial Vector Data.” *R J.* 10 (1): 439.
- Poelen, Jorrit H, James D Simons, and Chris J Mungall. 2014. “Global Biotic Interactions: An Open Infrastructure to Share and Analyze Species-Interaction Datasets.” *Ecological Informatics* 24: 148–59.
- Poisot, Timothée, Benjamin Baiser, Jennifer A Dunne, Sonia Kéfi, François Massol, Nicolas Mouquet, Tamara N Romanuk, Daniel B Stouffer, Spencer A Wood, and Dominique Gravel. 2016. “Mangal–Making Ecological Network Analysis Simple.” *Ecography* 39 (4): 384–90.
- Poisot, Timothée, Gabriel Bergeron, Kevin Cazelles, Tad Dallas, Dominique Gravel, Andrew MacDonald, Benjamin Mercier, Clément Violet, and Steve Vissault. 2021. “Global Knowledge Gaps in Species Interaction Networks Data.” *Journal of Biogeography* 48 (7): 1552–63.
- Potts, S, Koos Biesmeijer, Riccardo Bommarco, T Breeze, L Carnevalheiro, Markus Franzen, Juan P González-Varo, et al. 2015. “Status and Trends of European Pollinators. Key Findings of the STEP Project.”
- Proença, Vânia, Laura Jane Martin, Henrique Miguel Pereira, Miguel Fernandez, Louise McRae, Jayne Belnap, Monika Böhm, et al. 2017. “Global Biodiversity Monitoring: From Data Sources to Essential Biodiversity Variables.” *Biological Conservation* 213: 256–63.
- Ramos-Jiliberto, Rodrigo, Daniela Domínguez, Claudia Espinoza, Gioconda Lopez, Fernanda S Valdovinos, Ramiro O Bustamante, and Rodrigo Medel. 2010. “Topological Change of Andean Plant–Pollinator Networks Along an Altitudinal Gradient.” *Ecological Complexity* 7 (1): 86–90.
- Rech, André Rodrigo, Bo Dalsgaard, Brody Sandel, Jesper Sonne, Jens-Christian Svenning, Naomi Holmes, and Jeff Ollerton. 2016. “The Macroecology of Animal Versus Wind Pollination: Ecological Factors Are More Important Than Historical Climate Stability.” *Plant Ecology & Diversity* 9 (3): 253–62.
- Revell, Liam J. 2012. “Phytools: An r Package for Phylogenetic Comparative Biology (and Other Things).” *Methods in Ecology and Evolution*, no. 2: 217–23.
- Sala, Osvaldo E, FIII Stuart Chapin, Juan J Armesto, Eric Berlow, Janine Bloomfield, Rodolfo Dirzo, Elisabeth Huber-Sanwald, et al. 2000. “Global Biodiversity Scenarios for the Year 2100.” *Science* 287 (5459): 1770–74.
- Saunders, Manu E, Liam K Kendall, Jose B Lanuza, Mark A Hall, Romina Rader, and Jamie R Staver. 2023. “Climate Mediates Roles of Pollinator Species in Plant–Pollinator Networks.” *Global Ecology and Biogeography* 32 (4): 511–18.

- Settele, Josef, Jacob Bishop, and Simon G Potts. 2016. "Climate Change Impacts on Pollination." *Nature Plants* 2 (7): 1–3.
- Smith, Stephen A, and Joseph W Brown. 2018. "Constructing a Broadly Inclusive Seed Plant Phylogeny." *American Journal of Botany* 105 (3): 302–14.
- Song, Chuliang, Rudolf P Rohr, and Serguei Saavedra. 2017. "Why Are Some Plant–Pollinator Networks More Nested Than Others?" *Journal of Animal Ecology* 86 (6): 1417–24.
- Traveset, Anna, Cristina Tur, Kristian Trøjelsgaard, Ruben Heleno, Rocío Castro-Urgal, and Jens M Olesen. 2016. "Global Patterns of Mainland and Insular Pollination Networks." *Global Ecology and Biogeography* 25 (7): 880–90.
- Troia, Matthew J, and Ryan A McManamay. 2016. "Filling in the GAPS: Evaluating Completeness and Coverage of Open-Access Biodiversity Databases in the United States." *Ecology and Evolution* 6 (14): 4654–69.
- Trøjelsgaard, Kristian, and Jens M Olesen. 2013. "Macroecology of Pollination Networks." *Global Ecology and Biogeography* 22 (2): 149–62.
- Van der Loo, Mark PJ et al. 2014. "The Stringdist Package for Approximate String Matching." *R J.* 6 (1): 111.
- Vázquez, Diego P, Robert Poulin, Boris R Krasnov, and Georgy I Shenbrot. 2005. "Species Abundance and the Distribution of Specialization in Host-Parasite Interaction Networks." *Journal of Animal Ecology*, 946–55.
- Wetzel, Florian T, Heather C Bingham, Quentin Groom, Peter Haase, Urmas Kõljalg, Michael Kuhlmann, Corinne S Martin, et al. 2018. "Unlocking Biodiversity Data: Prioritization and Filling the Gaps in Biodiversity Observation Data in Europe." *Biological Conservation* 221: 78–85.
- Wiemers, Martin, Emilio Balletto, Vlad Dincă, Zdenek Faltýnek Fric, Gerardo Lamas, Vladimir Lukhtanov, Miguel L Munguira, et al. 2018. "An Updated Checklist of the European Butterflies (Lepidoptera, Papilionoidea)." *ZooKeys*, no. 811: 9.

Acknowledgments

We thank all the taxonomist and ecologist that has made this database possible by contributing with their fieldwork data. This research was funded by the H2020 European project Safeguard (101003476).

Author contributions

Supplementary material

Supplementary text 1

Habitat definitions:

- 1) **Ruderal vegetation:** Plants growing on highly disturbed sites such as road sides or mineral extraction sites.
- 2) **Agricultural margins:** Sides of crops that can include any type of vegetation from low growing plants to trees.
- 3) **Green urban areas:** Parks, private gardens or small pastures within an urban setting. Botanical gardens are included in this category.
- 4) **Agricultural land:** Includes any type of crop and any type of vegetation growing within them.
- 5) **Forest/woodland understory:** Any plant community sampled under a wooded group of plants. The forest could be embedded in an agricultural setting or in a fully natural scenario. We have included here agro-forestry areas and open to dense forest. Note that we have excluded from this category forest that contains sclerophyllous vegetation.
- 6) **Semi-natural grassland:** Low growing plant community with relatively low disturbances but under low pressure such as seasonal mowing or extensive grazing.
- 7) **Pastures:** Any type of low growing plant community that is highly influenced by human disturbance. For instance, agriculture, mowing, moderate to high grazing or urban environments. Note that this category also includes old pastures with regrowth of woody vegetation.
- 8) **Sclerophyllous vegetation:** Any type of system with a dominant shrub community adapted to drought. Typical of the Mediterranean region. Note, that we have include in this category also woodlands (open coniferous forest) where the shrub community was the main focus of the study.
- 9) **Beaches, dunes, sands:** Plant communities growing on sandy soil.
- 10) **Riparian vegetation:** Plant communities growing on river margins.
- 11) **Natural grasslands:** Low growing plant communities with little or none human disturbance. Often located in high elevation areas within Europe.
- 12) **Moors and heathland:** Low growing woody vegetation characteristic from low fertile soils near the coast or in alpine areas.

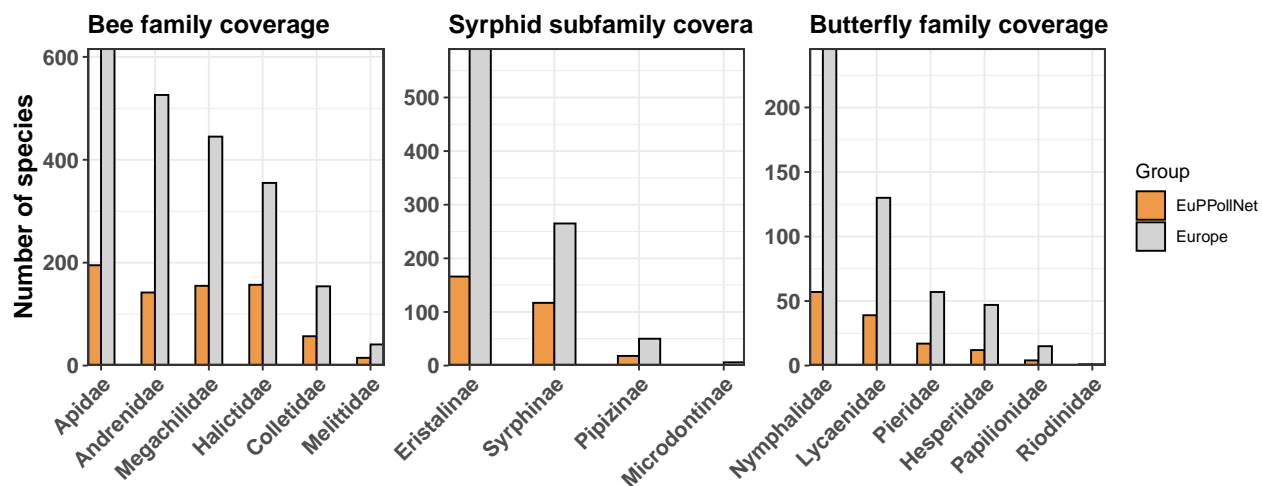


Figure S1. Coverage of the EuPPollNet species for bees (family level), syrphids (subfamily level) and butterflies (family level) in relation to the total number of European species within these taxonomic groups.

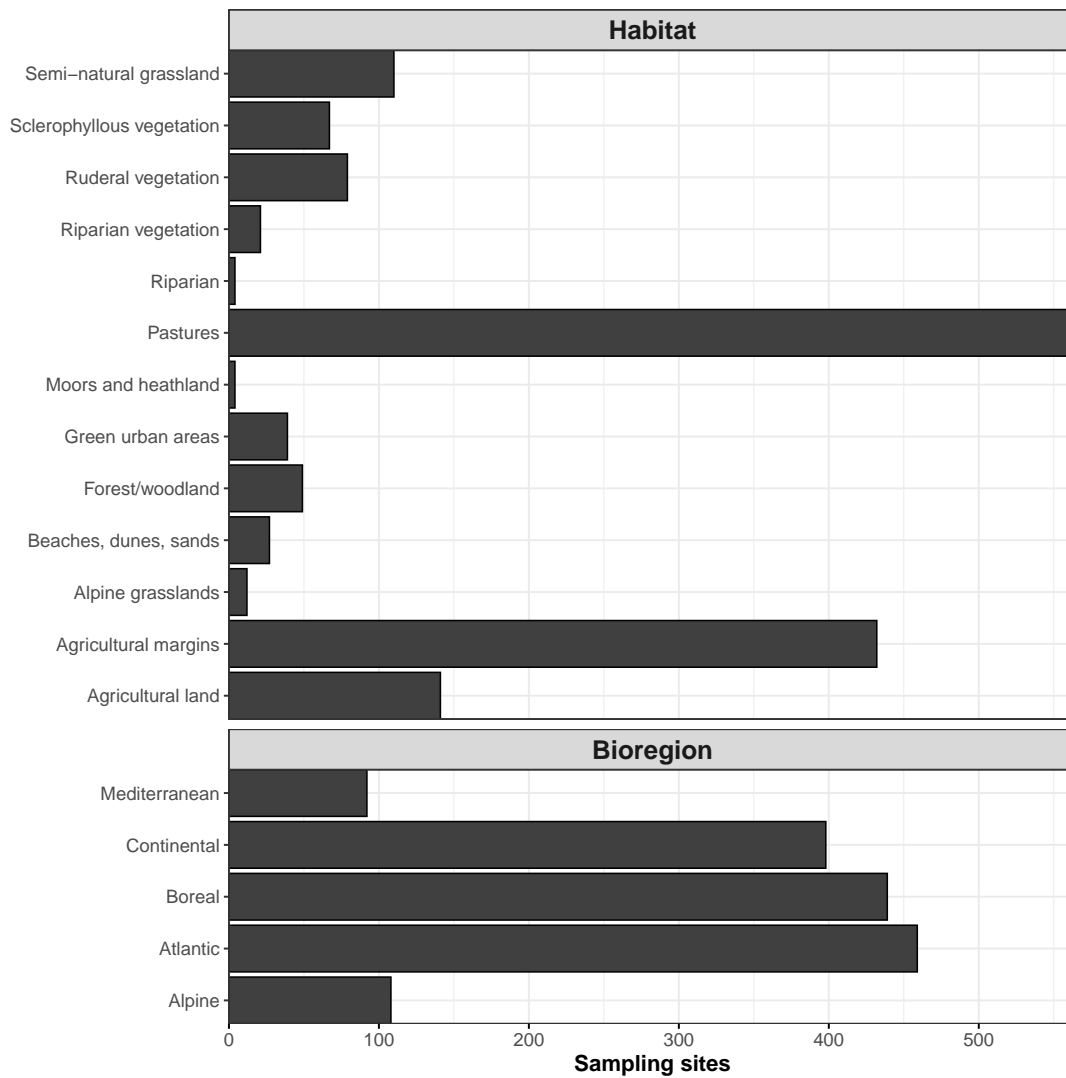


Figure S2. Barplot indicating the number of sampling sites by habitat and bioclimatic region within the database.

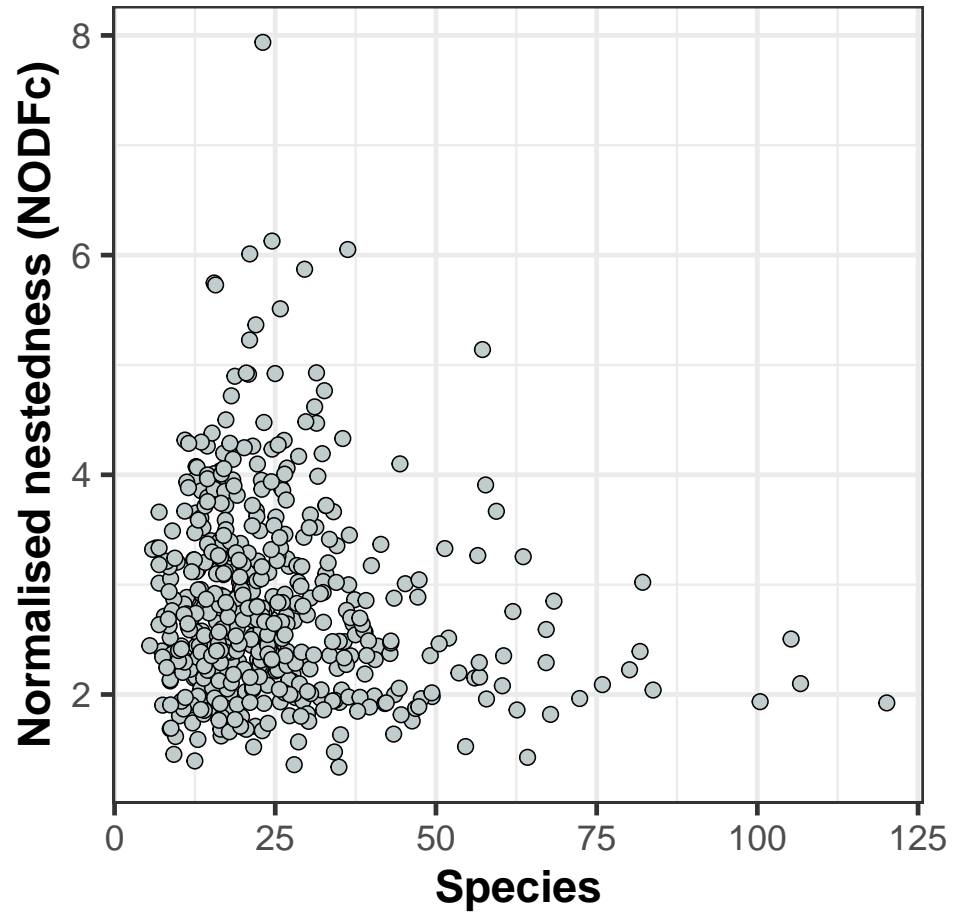


Figure S3. Association between nestedness (NODFc) and the geometric mean of plan and pollinator species per network