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Abstract

Assessing biodiversity status and trends in plant communities is critical for understanding, quantifying and predicting effects of global change on ecosystems. Vegetation plots record occurrence or abundance of all plant species present (community composition) in a delimited area of 0.01 to 40,000 m². Absences can be inferred, allowing analyses not possible with presence-only data. Recently the first global vegetation plot database was compiled ('sPlot'). However, this large dataset is environmentally and spatially unbalanced, and not open-access. We address both issues by (a) resampling the vegetation plots using a novel algorithm; (b) securing permission to openly release data from the holders of 104 specific datasets. We present the largest open-access vegetation plot dataset ever released: 91,205 plots globally, recording abundance of each vascular plant species (total 39,997 taxa). Plot-level data include location, date, size, biome, elevation, slope aspect, vegetation type and naturalness. Based on values for 18 traits per species from the 'TRY' database, community-weighted mean and variance of traits per plot are presented. The dataset can be used to explore plant community diversity patterns globally, as ground truth data in remote sensing applications or as baselines for biodiversity monitoring.

Background & Summary

Biodiversity is facing a global crisis. As many as 1 million species are currently threatened with extinction, the vast majority due to anthropogenic impacts such as land-use and climate change ([1](#), [2](#)). In addition, the rates of biodiversity homogenization and redistribution are accelerating ([3](#), [4](#); [5](#)). Biological assemblages are becoming progressively more similar to each other globally, as local and endemic species go extinct and are replaced by more widespread and competitive native or alien species ([1](#); [5](#)). Many terrestrial and marine species are also shifting their geographical distribution as a response to climate change ([4](#)), including animals hosting pathogens transmissible to humans ([6](#); [7](#)). This has profound potential impacts on human and ecosystem health ([8](#); [9](#)).

Plant communities are no exception to this biodiversity crisis ([10](#); [11](#); [5](#)). This is particularly worrying since terrestrial vegetation accounts for 80% (450 Gt C) of the living biomass on Earth ([12](#)). Given the central role of vegetation in ecosystem productivity, stability and functioning ([11](#)), assessing biodiversity status and trends in plant communities is paramount for other kingdoms of life and human societies alike.

Monitoring trends in plant biodiversity requires adequate data across a range of spatiotemporal scales ([13](#), [14](#)). Large independent collections of plant occurrence data do exist at the global or continental extent via the Botanical Information and Ecology Network (BIEN) ([\[???\]](#)), the Global Inventory of Floras and Traits (GIFT) ([15](#)) or the Global Biodiversity Information Facility (GBIF) (<https://www.gbif.org/>). However, these presence-only databases either neglect how individual plant species co-occur and interact locally to form plant communities, or are collected at spatial resolutions which are too coarse to assess biodiversity trends (e.g., one-degree grid cells) at the plant community scale ([16](#)).

Yet, there is a long tradition among botanists to record the cover or abundance of each plant species that occurs in a vegetation plot of a given size (i.e. surface area) at a given time (e.g. [17](#)). Compared to presence-only data, vegetation-plot data (termed 'presence-absence' here) present many advantages. First, they contain information on which plant species do, and do not, co-occur in the same locality at a given moment in time ([18](#)). This is important for testing hypotheses related to biotic interactions among plant species. Vegetation-plot data also provide crucial information on where and when a species was absent, therefore improving predictions from current species distribution models ([19](#)). Being spatially explicit, vegetation plots can be resurveyed through time to assess potential changes in plant species composition relative to a baseline ([20](#); [21](#), [5](#)). As they normally contain information on the relative cover or abundance of each species, vegetation plots are also more appropriate for detecting biodiversity changes than data representing only the occurrence of individual species ([22](#)).

Globally, however, vegetation-plot data are very fragmented, as they typically stem from a myriad of local research and survey projects ([23](#)). Consequently, these data often have either high fine-grain spatial resolutions but small spatial extents, or vice versa ([24](#)). Furthermore, with their disparate sampling protocols, standards and taxonomic resolutions, aggregating and harmonizing vegetation plot data proves extremely challenging ([25](#)). It is not surprising, therefore, that these data are rarely used in global-scale biodiversity research ([26](#); [27](#); [28](#)).

The sPlot initiative tries to close this data gap. It leverages numerous local-to-regional vegetation-plot datasets to create a harmonized and comprehensive global database of georeferenced terrestrial plant species assemblages ([23](#)). Established in 2013, sPlot (version 3) currently contains more than 1.9 million vegetation plots, and is fully integrated with the TRY database ([29](#)), from which it derives information on plant functional traits. The sPlot database is increasingly being used to study continental-to-global scale vegetation patterns, such as the relative contribution of regional vs. local factors on the global patterns of fern richness ([30](#)), the mechanisms underlying the spread and

abundance of native vs. invasive tree species ([31](#)), and worldwide trait–environment relationships in plant communities ([25](#)).

Here, we provide an open-access data set composed of 91,205 vegetation plots, that represent the entire environmental space covered by the sPlot database, to provide a standardized dataset for ecological research. The environmental stratification of the sampling of these plots maximises the benefits of this large dataset for a wide range of potential uses of the data. The selected vegetation plots stem from 104 databases and span 115 countries (Figure [1](#)). This resampled dataset (sPlot Open - hereafter) is composed of: (1) plot-level information, including metadata and basic vegetation structure descriptors; (2) the vascular plant species composition of each vegetation plot, including species cover or abundance information when available; and (3) community-level functional information derived from the TRY database ([29](#)).



Figure 1: Global map of sPlot Open ($n = 91,205$) and spatial distribution of vegetation plot density per hexagonal cell with a spatial resolution of approximately 70,000 km². Map projection is Eckert IV.

Methods

Vegetation plot data sources

We started from the sPlot database v2.1 (created October 2016), which contains 1,121,244 unique vegetation plots and 23,586,216 species records stemming from 110 different vegetation-plot datasets of regional, national or continental extent. Some of the 110 datasets stem from regional or continental initiatives (see [23](#) for more information). For instance: 48 vegetation-plot datasets derive from the European Vegetation Archive (EVA) ([18](#)); three major African datasets derive from the Tropical African Vegetation Archive (TAVA); and multiple vegetation datasets in the USA derive from the VegBank archive ([32](#); [33](#)). Data from other continents (South America, Asia) or countries were contributed as separate standalone datasets. The metadata of each individual vegetation-plot dataset stored in sPlot are managed through the Global Index of Vegetation-Plot Databases [GIVD](#) ([34](#)), using the GIVD code as the unique dataset identifier.

Resampling method

Data in the sPlot database are unevenly distributed across continents and biomes (see [25](#)). Mid-latitude regions in developed countries (mostly Europe, the USA and Australia) are overrepresented, while regions in the tropics and subtropics are underrepresented, which is a typical geographical bias in biodiversity data (e.g., [35](#); [4](#)). To reduce this imbalance as much as possible, we performed a stratified resampling approach, using several environmental variables available at global extent as sampling strata. We considered 30 climatic and soil variables. For climate, we complemented the 19 bioclimatic variables from CHELSA ([36](#)), as well as two variables reflecting growing-season warmth (growing degree days above 1 °C - GDD1 - and 5 °C - GDD5), which we calculated based on CHELSA bioclimatic variables. In addition, we considered an index of aridity (AR) and a model for Potential Evapotranspiration (PET - [37](#)). For soil, we extracted seven variables from the SOILGRIDS database ([38](#)), namely: soil organic carbon content in the fine earth fraction, cation exchange capacity, pH, as well as the fractions of coarse fragments, sand, silt and clay.

We stratified our sampling effort based on the following procedure. First, we ran a global principal component analysis (PCA) on a matrix of terrestrial grid cells by the 30 above-mentioned environmental variables. We considered the full environmental space of all terrestrial habitats on Earth at a spatial resolution of 2.5 arcmin, totaling 8,384,404 terrestrial grid cells, irrespective of whether a grid cell hosted vegetation plots from the sPlot database v2.1 or not. We then subdivided the PCA ordination space, represented by the first two principal components (PC1–PC2), which accounted for 47% and 23% of the total environmental variation in terrestrial grid cells, into a regular 100 × 100 grid. This PC1-PC2 two-dimensional space was subsequently used to balance our sampling effort across all PC1-PC2 grid cells for which vegetation plots were available. Before projecting vegetation plots from the sPlot database v2.1 onto this PC1-PC2 environmental space, we removed vegetation plots: a) from wetlands; b) from anthropogenic vegetation types; c) without geographical coordinates; and d) with a location uncertainty higher than 3 km for those having geographical coordinates. This resulted in a total of 799,400 out of the initial set of 1,121,244 vegetation plots. When projecting the 799,400 vegetation plots in the PC1-PC2 grid, we calculated how many vegetation plots occurred in each PC1-PC2 grid cell. For those grid cells with more than 50 vegetation plots ($n = 858$), we selected up to 50 vegetation plots using the heterogeneity-constrained random resampling algorithm from Lengyel et al. (2011) [[39](#)]. This approach optimizes the selection of a random subset of vegetation plots that encompasses the highest variability in species composition while avoiding peculiar and rare communities, which may represent outliers. We based the quantification of variability in plant species composition among the 50 randomly selected vegetation plots by computing the mean and the variance of the Jaccard's dissimilarity index ([40](#)) between all possible

pairs of these 50 vegetation plots ($n = 1225$). More precisely, for a given PC1-PC2 grid cell containing more than 50 vegetation plots, we generated 1,000 random selections of 50 vegetation plots and ranked them according to the mean (ascending order) and variance (descending order) value. Ranks from both sortings were summed for each random selection, and the selection with the lowest summed rank was considered to provide the most balanced/even representation of vegetation types within the focal grid cell. Where a grid cell contained less than 50 plots, we retained all of them. In this way, we reduced the imbalance towards over-sampled climate types while ensuring that the resampled dataset represents the entire environmental gradient covered by the sPlot database. We repeated the resampling procedure three times to get three different possibilities of a heterogeneity-constrained selection of 50 vegetation plots per PC1-PC2 grid cell with, initially, more than 50 vegetation plots. Vegetation plots selected during the first iteration were our first choice, while we considered the vegetation plots additionally selected in the second and third iteration as reserves when asking for permission to release the data as open access to each dataset's contributor(s).

Permission to release the data as open access

The resampling procedure resulted in a preliminary potential selection of 98,383 vegetation plots (first choice) and 51,634 vegetation plots flagged as reserves (second or third choice for the subset of PC1-PC2 grid cells with more than 50 vegetation plots available). Since the sPlot database is a consortium of independent datasets whose copyright belongs to the data contributor, we used this preliminary potential selection to ask each dataset's custodian (i.e., either the owner of a dataset or its authorized representative in case of a collective dataset) for permission to release the data of each selected vegetation plot as open access. For 8,070 vegetation plots, permission could not be granted because, for instance, the data are unpublished, confidential or sensitive. For these vegetation plots, we used the reserve pool to randomly select replacements, for which such permission could be granted. We imposed the constraint that each candidate vegetation plot in the reserve pool should belong to the same environmental stratum, i.e., the same PC1-PC2 grid cell, of the confidential vegetation plot. Note that 2,380 PC1-PC2 grid cells (11.7% of total) had one or more confidential vegetation plots (median = 1, mean = 3.4, max = 171) that could not be replaced from the reserve pool.

Trait information

For each vegetation plot for which open access has been granted, we computed the community weighted means for eighteen plant functional traits derived from the TRY database v3.0 ([29](#)). These traits were selected among those that describe the leaf, wood and seed economics spectra ([41](#); [42](#)), and are known to either affect different key ecosystem processes or respond to macroclimatic drivers, or both ([23](#)). The eighteen plant functional traits (all concentrations based on dry weight) were: (1) leaf area [mm^2]; (2) stem specific density [g cm^{-3}]; (3) specific leaf area [m^2kg^{-1}]; (4) leaf carbon concentration [mg g^{-1}]; (5) leaf nitrogen concentration [mg g^{-1}]; (6) leaf phosphorus concentration [mg g^{-1}]; (7) plant height [m]; (8) seed mass [mg]; (9) seed length [mm]; (10) leaf dry matter content [g g^{-1}]; (11) leaf nitrogen per area [g m^{-2}]; (12) leaf N:P ratio [g g^{-1}]; (13) leaf $\delta^{15}\text{N}$ [per million]; (14) seed number per reproductive unit; (15) leaf fresh mass [g]; (16) stem conduit density [mm^{-2}]; (17) dispersal unit length [mm]; and (18) conduit element length [μm].

Because missing values were particularly widespread in the species-trait matrix, we employed a gap-filling procedure based on hierarchical Bayesian modeling (R package 'BHPMF', [43](#); [44](#)). Gap-filling was performed at the level of individual observations. We then transformed to the natural logarithm all gap-filled trait values and averaged each trait by taxon (i.e., at species, or genus level). Additional information on the gap-filling procedure is available in [[23](#)].

Community-weighted means (CWM) and variances (CWV) were calculated for every plant functional trait j and every vegetation plot k as follows ([45](#)):

$$CWM_{j,k} = \sum_i^{n_k} p_{i,k} t_{i,j} \quad (1)$$

$$CWW_{j,k} = \sum_i^{n_k} p_{i,k} (t_{i,j} - CWM_{j,k})^2 \quad (2)$$

where n_k is the number of species with trait information in vegetation plot k , $p_{i,k}$ is the relative abundance of species i in vegetation plot k calculated as the species' fraction in cover or abundance of total cover or abundance, and $t_{i,j}$ is the mean value of species i for trait j .

Data Records

sPlot Open contains 91,205 vegetation plots (also called 'relevés') from 115 countries and all continents except Antarctica (Figure 1). This randomized selection comes from 105 constitutive datasets (Table 1). It only contains the species composition of vascular plants; information on the composition of bryophytes and lichens was discarded since it was only available for a minority of plots ($n = 4,963$ and $n = 3,045$, respectively). Information on the size (surface area) of the vegetation survey is available for 61,898 vegetation plots, and ranges between 0.01 and 40,000 m² (mean = 270 m²; median = 78.5 m²). Similarly, only for a minority of plots ($n = 17,757$) information on the exact group of plants sampled in the field is available (e.g., complete vegetation, only trees, only trees > 1 m height, and so on). However, as most data were collected using the phytosociological method, we deem safe to assume that, unless otherwise specified, plots contain information on all vascular plants. The average number of vascular plant species per vegetation plot ranges between 1 (i.e. monospecific stands) and 270 species (mean = 17.6; median = 13).

By capping the number of vegetation plots in overrepresented environmental conditions, the resampling procedure described above strongly reduced the bias in the distribution of vegetation plots within the environmental niche space. Yet, due to the lack or scarcity of data from some geographical regions, like the tropics, the spatial distribution of vegetation plots remains unbalanced across geographical regions (Figure 1). This is evident when comparing the number of plots across continents or biomes. Europe is by far the best represented continent, with 53,884 vegetation plots. In contrast, in Africa and South America the remaining plots after data edition and selection were 4,507 and 5,533 vegetation plots, respectively. The representation of biomes is also unbalanced (Figure 2). Despite these imbalances, all the Whittaker biomes are covered by sPlot Open, and our resampling algorithm has resulted in a much more balanced dataset than many other large global datasets that are available, such as GBIF.



Figure 2: Distribution of all the vegetation plots provided by sPlot Open ($n = 91,205$) in the two-dimensional climatic space represented by mean annual temperature and mean annual precipitation superimposed onto Whittaker biomes

Almost one third of vegetation plots in sPlot Open belong to forest ($n = 25,740$), two thirds to non-forest vegetation ($n = 58,145$) vegetation, with 8 % of plots remaining unassigned ($n = 7,320$). When not directly done by data providers, the assignment of plots to forests and non-forests was based on multiple lines of evidence, including the plot-level information on the cover of the tree layer, as well as traits of species composing a plot, such as growth form and height. In short, a plot record was considered as forest if the cover of the tree layer, or alternatively, the sum of the (relative) cover of all tree taxa (normalized to 100%), was greater than 25%. It was considered a non-forest record if the sum of relative cover of low-stature, non-tree and non-shrub taxa was greater than 90%. For an extensive explanation of this classification scheme, we refer the reader to Brulheide et al. (2019) [23]. Even if the proportion of forest vs. non-forest vegetation plots is relatively well-balanced, the geographical distribution of vegetation plots belonging to different vegetation types is likely not balanced in the geographical space, as it depends on the idiosyncrasies of the constitutive datasets composing the sPlot database. For instance, the data from New Zealand only include plots collected in non-forest ecosystems, while data from Chile only refer to forests. We urge potential users to carefully read the description of each individual dataset in [GIVD](#), or to contact the custodians of each dataset before using sPlot Open.

Database Organization

sPlot Open is organized into three main matrices, relationally linked through the key column *'PlotObservationID'*.

The **'header'** matrix contains plot-level information for the 91,205 vegetation plots provided in sPlot Open, including: metadata (e.g., plot ID, ownership, sampling date, geographical location, positional accuracy); sampling design information (e.g., the total surface area used during the vegetation survey); and a plot-level description of vegetation structure (e.g., vegetation type, percentage cover of each vegetation layer) and vegetation type. Plots in Europe are also classified according to the EUNIS habitat classification (column *'ESY'*), based on the habitat classification expert system described in Chytrý et al. (2020) ([47](#)). For each vegetation plot, we further provide information on the dataset it originates from, based on the IDs used in [GIVD](#). A brief description of all the 43 variables in the header matrix is provided in Table [2](#).

The **'DT'** matrix contains data on the species composition of each plot. It is structured in a long format and contains 1,608,610 records from 39,997 vascular plant taxa, mostly resolved at the species level. For each record, we report both the taxon name as originally contributed by the data custodian (column *'Original_species'*), and the taxon name after taxonomic standardization (column *'Species'*). For each record, we report the species cover/abundance values. These follow different standards across the datasets constituting the sPlot database. We, therefore, provide both the cover/abundance value as reported in the original data (column *'Original_abundance'*), together with the abundance scale that was originally used (column *'Abundance_scale'*). This can take seven values: *'CoverPerc'* = percentage cover, *'pa'* = presence-absence, *'x_BA'* = basal area (m²/ha, only for woody species), *'x_IC'* = individual count, i.e., number of individuals in plot, *'x_SC'* = stem count, i.e., number of stems in plot, *'x_IV'* = importance value index, *'x_PF'* = presence frequency. The great majority of entries, however, use the percentage cover scale (n= 1,397,109). Finally, for each entry, we calculated a *'Relative_cover'*, i.e., the cover/abundance of a given taxon divided by the total cover/abundance of all taxa in that vegetation plot.

The **'CWM_CWV'** matrix contains the community-weighted means and variances calculated for each of the 18 functional traits mentioned above. It also contains three additional columns. The column *'Species_richness'* returns the number of species recorded in each plot. The columns *'Trait_coverage_cover'* and *'Trait_coverage_pa'* return, respectively, the proportion of total cover and species in a plot for which functional trait information was available. Functional trait information was available for 20,932 species. The average proportion of species in each plot for which we have functional trait information is 0.88 (median = 1). For 47,177 plots, the coverage is complete, while for only one plot do we have no functional trait information for any of the species occurring in it. When considering relative cover, the average trait coverage is 0.89. As many as 68,234 and 74,388 plots have functional trait information for more than 80% of the species or 80% of relative cover, respectively.

sPlot Open contains two additional objects. The **'metadata'** matrix contains plot-level metadata, which provide information on the origin of each individual vegetation plot. This object contains 15 columns, with information on the dataset of origin (column *'GIVD_ID'* - [34](#)), author or surveyor names (columns *'Releve_author'* and *'Releve_coauthor'*), bibliographic references both at the dataset (column *'DB_BIBTEXKEY'*) and plot level (*'Plot_Biblioreference'* and *'BIBTEXKEY'*), when available. Similarly, the column *'Project_name'* provides information on the project in which a vegetation plot was collected. When available, we also provide information on the numbering of the plots in the publication where they originally appeared (columns *'Nr_table_in_publ'*, *'Nr_releve_in_table'*), or in the dataset where they were initially stored (*'Original_nr_in_database'*). In the case of nested plots (n = 1,786), we also provide the original plot and subplot IDs (columns: *'Original_plotID'*, *'Original_subplotID'*). The last two columns

report plot-level *'Remarks'*, and the unique identifier produced by Turboveg when the vegetation plot was first stored (*'GUID'*).

Finally, the object **'references'**, contains all the bibliographic references formatted according to a BibTex standard. Each reference is tagged with a key corresponding to the fields *'DB_BIBTEXKEY'* and *'BIBTEXKEY'* in the metadata. We further provide an R function (*'sPlotOpen_citation'*) to create reference lists, based on a selection of plots and/or datasets.

Except for the 'reference' file (format .bib), all objects/matrices are provided in tab-delimited .txt files. All objects, including the *'sPlotOpen_citation'* function, are also compiled inside an .RData object.

Technical Validation

The original sPlot database has a nested structure and consists of several individual datasets, each validated and maintained by its respective dataset custodian. In many cases, individual datasets are also collections whose vegetation plots were provided by their respective owners (the person who performed the actual vegetation survey) or by someone who digitized the original data from the scientific published or grey literature. We obviously have no direct control over the individual vegetation plots that we provide here in sPlot Open. Yet, all these vegetation plots stem from trained professional botanists, or published scientific work, and are accompanied by detailed information on the sampling protocols used, thus ensuring data quality and reliability.

Before integration into the sPlot database, each dataset was further checked for consistency and, if it was in a different format, we converted it to a Turboveg 2 database (48). During this conversion, we checked that all datasets contained the required metadata information, and cross-checked that each plot was located within the geographic scopes of its respective dataset. Finally, we harmonized all the taxonomic names from all datasets, based on the sPlot's taxonomic backbone (49). This backbone matched all the taxonomic names (without nomenclatural authors) from all datasets in sPlot 2.1 and TRY v3.0 (29) to their resolved version based on the Taxonomic Name Resolution Service web application (TNRS version 4.0; 50; iPlant Collaborative, 2015). This allowed us to (1) harmonize all datasets to a common nomenclature, and (2) link the sPlot database to the TRY database (29). All taxa originally denoted at taxonomic ranks lower than species were aggregated at species level. Additional detail on the taxonomic resolution is reported in Bruelheide et al. (2019) [23], while a description of the workflow, including R-code, is available in Purschke (2017) [49].

Usage Notes

The sPlot Open database can be downloaded from <https://www.idiv.de> (link to PlantHub). Users are urged to cite the original sources when using sPlot Open in addition to the present paper, particularly when using data contained in BioTIME (51). For two datasets (AF-00-009, AF-CD-001), the identification of taxa at species level is still in progress. As most of the constitutive datasets remain under continuous development, sPlot Open users are encouraged to get in touch with the custodian(s) of the data they are planning to use (custodian names are reported in <https://www.idiv.de/sPlot>).

The data included in the present paper represent the subset of sPlot for which we were able to secure permission for making these data open. The additional data in sPlot are available under sPlot's Governance and Data Property Rules (www.idiv.de/sPlot).

Code Availability

The R code used to produce sPlot Open from the sPlot 2.1 database is contained in the *sPlotOpen_code* GitHub repository: (https://github.com/fmsabatini/sPlotOpen_Code/). This manuscript was produced using the Manubot workflow (52). The code for reproducing this manuscript is stored in the *sPlotOpen_manuscript* GitHub repository: (https://github.com/fmsabatini/sPlotOpen_Manuscript).

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Author contributions

FMS wrote the first draft of the manuscript, with considerable input from JL and HB. JL and TH wrote the resampling algorithm. FMS set up the GitHub projects, curated the database, and produced the graphs. He also coordinated the sPlot consortium. SMH wrote the Turboveg v3 software, which holds the sPlot database. JK provided the trait data from TRY and FS performed the trait data gap filling. HB secured the funding for sPlot as a strategic project of iDiv. All other authors contributed data and/or helped set up the database and/or helped develop the resampling algorithm. All authors contributed to revising the manuscript.

Competing interests

The authors declare no competing interests.

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DOI: [10.1201/9781315368252](https://doi.org/10.1201/9781315368252)

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Thomas Ibanez, Jérôme Munzinger, Gilles Dagostini, Vanessa Hequet, Frédéric Rigault, Tanguy Jaffré, Philippe Birnbaum

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Jonathan Lenoir, Bente Jessen Graae, Per Arild Aarrestad, Inger Greve Alsos, W. Scott Armbruster, Gunnar Austrheim, Claes Bergendorff, H. John B. Birks, Kari Anne Bråthen, Jörg Brunet, ... Jens-Christian Svenning
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DOI: [10.1111/gcb.12129](https://doi.org/10.1111/gcb.12129) · PMID: [23504984](https://pubmed.ncbi.nlm.nih.gov/23504984/)

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Kiril Vassilev, Hristo Pedashenko, Alexandra Alexandrova, Alexandar Tashev, Anna Ganeva, Anna Gavrilova, Asya Gradevska, Assen Assenov, Antonina Vitkova, Borislav Grigorov, ... Vladimir Vulchev
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Flavia Landucci, Marcela Řezníčková, Kateřina Šumberová, Milan Chytrý, Liene Aunina, Claudia Biță-Nicolae, Alexander Bobrov, Lyubov Borsukevych, Henry Brisse, Andraž Čarni, ... Wolfgang Willner
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DOI: [10.7809/b-e.00074](https://doi.org/10.7809/b-e.00074)
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DOI: [10.7809/b-e.00177](https://doi.org/10.7809/b-e.00177)
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DOI: [10.7809/b-e.00178](https://doi.org/10.7809/b-e.00178)
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Biodiversity & Ecology (2012-09-10) <https://doi.org/ghgt9f>
DOI: [10.7809/b-e.00180](https://doi.org/10.7809/b-e.00180)
101. **Hungarian Phytosociological database (COENODATREF): sampling methodology, nomenclature and its actual stage**
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DOI: [10.7809/b-e.00192](https://doi.org/10.7809/b-e.00192)

104. Nationwide Vegetation Plot Database – Sapienza University of Rome: state of the art, basic figures and future perspectives

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Solvita Rūsiņa

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Adrian Indreica, Pavel Dan Turtureanu, Anna Szabó, Irina Irimia

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DOI: [10.1127/phyto/2017/0201](https://doi.org/10.1127/phyto/2017/0201)

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Kiril Vassilev, Eszter Ruprecht, Valeriu Alexiu, Thomas Becker, Monica Beldean, Claudia Biță-

Nicolae, Anna Mária Csörgő, Iliana Dzhovanova, Eva Filipova, József Pál Frink, ... Jürgen Dengler

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DOI: [10.1127/phyto/2017/0229](https://doi.org/10.1127/phyto/2017/0229)

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Svetlana Ačić, Milica Petrović, Urban Šilc, Zora Dajić Stevanović

Biodiversity & Ecology (2012-09-10) <https://doi.org/ghgt9h>

DOI: [10.7809/b-e.00206](https://doi.org/10.7809/b-e.00206)

111. Lower Volga Valley Phytosociological Database

Alexey Sorokin, Valentin Golub, Kseniya Starichkova, Lyudmila Nikolaychuk, Viktoria Bondareva,

Tatyana Ivakhnova

Biodiversity & Ecology (2012-09-10) <https://doi.org/ghgt9j>
DOI: [10.7809/b-e.00207](https://doi.org/10.7809/b-e.00207)

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Tatiana Lysenko, Olga Kalmykova, Anna Mitroshenkova
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DOI: [10.7809/b-e.00208](https://doi.org/10.7809/b-e.00208)

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Vadim Prokhorov, Tatiana Rogova, Maria Kozhevnikova
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DOI: [10.7809/b-e.00216](https://doi.org/10.7809/b-e.00216)

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DOI: [10.7809/b-e.00217](https://doi.org/10.7809/b-e.00217)

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Luis Cayuela, Lucía Gálvez-Bravo, Ramón Pérez Pérez, Fábio de Albuquerque, Duncan Golicher, Rakan Zahawi, Neptalí Ramírez-Marcial, Cristina Garibaldi, Richard Field, José Rey Benayas, ... Regino Zamora
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DOI: [10.7809/b-e.00078](https://doi.org/10.7809/b-e.00078)

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Viktoria Wagner, Toby Spribille, Stefan Abrahamczyk, Erwin Bergmeier
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Robert Peet, Michael Lee, Michael Jennings, Don Faber-Langendoen

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DOI: [10.7809/b-e.00080](https://doi.org/10.7809/b-e.00080)

122. Vegetation-plot database of the Carolina Vegetation Survey

Robert Peet, Michael Lee, Forbes Boyle, Thomas Wentworth, Michael Schafale, Alan Weakley
Biodiversity & Ecology (2012-09-10) <https://doi.org/ghgvcm>
DOI: [10.7809/b-e.00081](https://doi.org/10.7809/b-e.00081)

123. The Alaska Arctic Vegetation Archive (AVA-AK)

Donald A. Walker, Amy L. Breen, Lisa A. Druckenmiller, Lisa W. Wirth, Will Fisher, Martha K. Reynolds, Jozef Šibík, Marilyn D. Walker, Stephan Hennekens, Keith Boggs, ... Donatella Zona
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DOI: [10.1127/phyto/2016/0128](https://doi.org/10.1127/phyto/2016/0128)

124. VegPáramo, a flora and vegetation database for the Andean páramo

Gwendolyn Peyre, Henrik Balslev, David Martí, Petr Sklenář, Paul Ramsay, Pablo Lozano, Nidia Cuello, Rainer Bussmann, Omar Cabrera, Xavier Font
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DOI: [10.1007/s00035-016-0182-6](https://doi.org/10.1007/s00035-016-0182-6)

Supplementary Material

Table 1: List of databases contributing to the open access dataset extracted from the sPlot database. Databases are ordered based on their ID in the Global Index of Vegetation Databases (GVID ID).

| GVID ID | Dataset name | Custodian | Deputy custodian | Nr. OA plots | Ref |
|-----------|--|--------------------------|------------------|--------------|--------------------|
| 00-00-001 | ForestPlots.net | Oliver L. Phillips | Aurora Levesley | 108 | 53 |
| 00-00-003 | SALVIAS | Brian Enquist | Brad Boyle | 2860 | |
| 00-00-004 | Vegetation Database of Eurasian Tundra | Risto Virtanen | | 600 | |
| 00-00-005 | Tundra Vegetation Plots (TundraPlot) | Anne D. Bjorkman | Sarah Elmendorf | 227 | 54 |
| 00-RU-001 | Vegetation Database Forest of Southern Ural | Vasiliy Martynenko | Pavel Shirokikh | 25 | |
| 00-RU-002 | Database of Masaryk University`s Vegetation Research in Siberia | Milan Chytrý | | 128 | 55 |
| 00-RU-003 | Database Meadows and Steppes of Southern Ural | Sergey Yamalov | Mariya Lebedeva | 99 | |
| 00-TR-001 | Forest Vegetation Database of Turkey - FVDT | Ali Kavgacı | | 15 | |
| AF-00-001 | West African Vegetation Database | Marco Schmidt | Georg Zizka | 184 | 56 |
| AF-00-003 | BIOTA Southern Africa Biodiversity Observatories Vegetation Database | Norbert Jürgens | Ute Schmiedel | 562 | 57 |
| AF-00-006 | SWEA-Dataveg | Miguel Alvarez | Michael Curran | 1211 | |
| AF-00-008 | PANAF Vegetation Database | Hjalmar Kühl | TeneKwetché Sop | 942 | |
| AF-00-009 | Vegetation Database of the Okavango Basin | Rasmus Revermann | Manfred Finckh | 202 | 58 |
| AF-BF-001 | Sahel Vegetation Database | Jonas V. Müller | Marco Schmidt | 279 | 59 |
| AF-CD-001 | Forest Database of Central Congo Basin | Kim Sarah Jacobsen | Hans Verbeeck | 97 | 60 |
| AF-ET-001 | Vegetation Database of Ethiopia | Desalegn Wana | Anke Jentsch | 59 | 61 |
| AF-MA-001 | Vegetation Database of Southern Morocco | Manfred Finckh | | 266 | 62 |
| AF-ZW-001 | Vegetation Database of Zimbabwe | Cyrus Samimi | | 17 | 63 |
| AS-00-001 | Korean Forest Database | Tomáš Černý | Jiri Dolezal | 766 | 64 |
| AS-00-003 | Vegetation of Middle Asia | Arkadiusz Nowak | Marcin Nobis | 128 | 65 |
| AS-00-004 | Rice Field Vegetation Database | Arkadiusz Nowak | | 31 | |
| AS-BD-001 | Tropical Forest Dataset of Bangladesh | Mohammed A.S. Arfin Khan | Fahmida Sultana | 82 | |
| AS-CN-001 | China Forest-Steppe Ecotone Database | Hongyan Liu | Fengjun Zhao | 97 | 66 |
| AS-CN-002 | Tibet-PaDeMoS Grazing Transect | Karsten Wesche | | 27 | 67 |
| AS-CN-003 | Vegetation Database of the BEF China Project | Helge Bruehlheide | | 18 | 68 |
| AS-CN-004 | Vegetation Database of the Northern Mountains in China | Zhiyao Tang | | 70 | |

| GIVD ID | Dataset name | Custodian | Deputy custodian | Nr. OA plots | Ref |
|----------------|---|-------------------------------------|-------------------------|---------------------|--------------------|
| AS-EG-001 | Vegetation Database of Sinai in Egypt | Mohamed Z. Hatim | | 98 | 69 |
| AS-ID-001 | Sulawesi Vegetation Database | Michael Kessler | | 24 | |
| AS-IR-001 | Vegetation Database of Iran | Jalil Noroozi | Parastoo Mahdavi | 105 | |
| AS-KZ-001 | Database of Meadow Vegetation in the NW Tien Shan Mountains | Viktoria Wagner | | 3 | 70 |
| AS-MN-001 | Southern Gobi Protected Areas Database | Henrik von Wehrden | Karsten Wesche | 688 | 71 |
| AS-RU-001 | Wetland Vegetation Database of Baikal Siberia (WETBS) | Victor Chepinoga | | 6 | 72 |
| AS-RU-002 | Database of Siberian Vegetation (DSV) | Andrey Korolyuk | Andrei Zverev | 2150 | 73 |
| AS-RU-004 | Database of the University of Münster - Biodiversity and Ecosystem Research Group's Vegetation Research in Western Siberia and Kazakhstan | Norbert Hölzel | Wanja Mathar | 85 | |
| AS-SA-001 | Vegetation Database of Saudi Arabia | Mohamed Abd El-Rouf Mousa El-Sheikh | | 607 | 74 |
| AS-TJ-001 | Eastern Pamirs | Kim André Vanselow | | 174 | 75 |
| AS-TW-001 | National Vegetation Database of Taiwan | Ching-Feng Li | Chang-Fu Hsieh | 897 | |
| AS-YE-001 | Socotra Vegetation Database | Michele De Sanctis | Fabio Attorre | 190 | 76 |
| AU-AU-002 | AEKOS | Ben Sparrow | | 7443 | 77 |
| AU-NC-001 | New Caledonian Plant Inventory and Permanent Plot Network (NC-PIPPN) | Jérôme Munzinger | Philippe Birnbaum | 98 | 78 |
| AU-NZ-001 | New Zealand National Vegetation Databank | Susan K. Wiser | | 983 | 79 |
| AU-PG-001 | Forest Plots from Papua New Guinea | Timothy Whitfeld | George D. Weiblen | 53 | 80 |
| EU-00-002 | Nordic-Baltic Grassland Vegetation Database (NBGVD) | Jürgen Dengler | Łukasz Kozub | 931 | 81 |
| EU-00-011 | Vegetation-Plot Database of the University of the Basque Country (BIOVEG) | Idoia Biurrun | Itziar García-Mijangos | 1694 | 82 |
| EU-00-013 | Balkan Dry Grasslands Database | Kiril Vassilev | Armin Macanović | 224 | 83 |
| EU-00-016 | Mediterranean Ammophiletea Database | Corrado Marcenò | Borja Jiménez-Alfaro | 3713 | 84 |
| EU-00-017 | European Coastal Vegetation Database | John Janssen | | 1369 | |
| EU-00-018 | The Nordic Vegetation Database | Jonathan Lenoir | Jens-Christian Svenning | 1755 | 85 |
| EU-00-019 | Balkan Vegetation Database | Kiril Vassilev | Hristo Pedashenko | 211 | 86 |
| EU-00-020 | WetVegEurope | Flavia Landucci | | 61 | 87 |
| EU-00-022 | European Mire Vegetation Database | Tomáš Peterka | Martin Jiroušek | 1843 | 88 |
| EU-AL-001 | Vegetation Database of Albania | Michele De Sanctis | Giuliano Fanelli | 99 | 89 |

| GIVD ID | Dataset name | Custodian | Deputy custodian | Nr. OA plots | Ref |
|----------------|---|----------------------------|-------------------------|---------------------|---------------------|
| EU-AT-001 | Austrian Vegetation Database | Wolfgang Willner | Christian Berg | 950 | 90 |
| EU-BE-002 | INBOVEG | Els De Bie | | 48 | |
| EU-BG-001 | Bulgarian Vegetation Database | Iva Apostolova | Desislava Sopotlieva | 74 | 91 |
| EU-CH-005 | Swiss Forest Vegetation Database | Thomas Wohlgemuth | | 1409 | 92 |
| EU-CZ-001 | Czech National Phytosociological Database | Milan Chytrý | Ilona Knollová | 579 | 93 |
| EU-DE-001 | VegMV | Florian Jansen | Christian Berg | 5 | 94 |
| EU-DE-013 | VegetWeb Germany | Florian Jansen | Jörg Ewald | 199 | 95 |
| EU-DE-014 | German Vegetation Reference Database (GVRD) | Ute Jandt | Helge Bruelheide | 286 | 96 |
| EU-DK-002 | National Vegetation Database of Denmark | Jesper Erenskjold Moeslund | Rasmus Ejrnæs | 1181 | |
| EU-ES-001 | Iberian and Macaronesian Vegetation Information System (SIVIM) - Wetlands | Aaron Pérez-Haase | Xavier Font | 292 | |
| EU-FR-003 | SOPHY | Emmanuel Garbolino | Patrice De Ruffray | 13322 | 97 |
| EU-GB-001 | UK National Vegetation Classification Database | John S. Rodwell | | 5457 | |
| EU-GR-001 | KRITI | Erwin Bergmeier | | 43 | |
| EU-GR-005 | Hellenic Natura 2000 Vegetation Database (HelNatVeg) | Panayotis Dimopoulos | Ioannis Tsiripidis | 777 | 98 |
| EU-GR-006 | Hellenic Woodland Database | Ioannis Tsiripidis | Georgios Fotiadis | 4 | 99 |
| EU-HR-001 | Phytosociological Database of Non-Forest Vegetation in Croatia | Zvezdana Stančić | | 213 | 100 |
| EU-HR-002 | Croatian Vegetation Database | Željko Škvorc | Daniel Krstonošić | 688 | |
| EU-HU-003 | CoenoDat Hungarian Phytosociological Database | János Csiky | Zoltán Botta-Dukát | 17 | 101 |
| EU-IT-001 | VegItaly | Roberto Venanzoni | Flavia Landucci | 2712 | 102 |
| EU-IT-010 | Vegetation database of Habitats in the Italian Alps – HabItAlp | Laura Casella | Pierangela Angelini | 155 | 103 |
| EU-IT-011 | Vegetation-Plot Database Sapienza University of Rome (VPD-Sapienza) | Emiliano Agrillo | Fabio Attorre | 1003 | 104 |
| EU-LT-001 | Lithuanian Vegetation Database | Valerijus Rašomavičius | Domas Uogintas | 119 | |
| EU-LV-001 | Semi-natural Grassland Vegetation Database of Latvia | Solvita Rūsiņa | | 306 | 105 |
| EU-MK-001 | Vegetation Database of the Republic of Macedonia | Renata Ćušterevska | | 10 | |
| EU-NL-001 | Dutch National Vegetation Database | Stephan M. Hennekens | Joop H.J. Schaminée | 10223 | 106 |

| GIVD ID | Dataset name | Custodian | Deputy custodian | Nr. OA plots | Ref |
|----------------|---|-----------------------------|-------------------------|-----------------------------|---------------------|
| EU-PL-001 | Polish Vegetation Database | Zygmunt Kącki | Grzegorz Swacha | 464 | 107 |
| EU-RO-007 | Romanian Forest Database | Adrian Indreica | Pavel Dan Turtureanu | 60 | 108 |
| EU-RO-008 | Romanian Grassland Database | Eszter Ruprecht | Kiril Vassilev | 44 | 109 |
| EU-RS-002 | Vegetation Database Grassland Vegetation of Serbia | Svetlana Ačić | Zora Dajić Stevanović | 57 | 110 |
| EU-RU-002 | Lower Volga Valley Phytosociological Database | Valentin Golub | Andrey Chuvashov | 149 | 111 |
| EU-RU-003 | Vegetation Database of the Volga and the Ural Rivers Basins | Tatiana Lysenko | | 96 | 112 |
| EU-RU-011 | Vegetation Database of Tatarstan | Vadim Prokhorov | Maria Kozhevnikova | 94 | 113 |
| EU-SI-001 | Vegetation Database of Slovenia | Urban Šilc | Filip Kuzmič | 435 | 114 |
| EU-SK-001 | Slovak Vegetation Database | Milan Valachovič | Jozef Šibík | 893 | 115 |
| EU-UA-001 | Ukrainian Grasslands Database | Anna Kuzemko | Yulia Vashenyak | 149 | 116 |
| EU-UA-006 | Vegetation Database of Ukraine and Adjacent Parts of Russia | Viktor Onyshchenko | Vitaliy Kolomiychuk | 479 | |
| NA-00-002 | Tree Biodiversity Network (BIOTREE-NET) | Luis Cayuela | | 208 | 117 |
| NA-CA-003 | Database of Timberline Vegetation in NW North America | Viktoria Wagner | Toby Spribille | 38 | 118 |
| NA-CA-004 | Understory of Sugar Maple Dominated Stands in Quebec and Ontario (Canada) | Isabelle Aubin | | 9 | 119 |
| NA-CA-005 | Boreal Forest of Canada | Yves Bergeron | Louis De Grandpré | 44 | |
| NA-GL-001 | Vegetation Database of Greenland | Birgit Jedrzejek | Fred J.A. Daniëls | 340 | 120 |
| NA-US-002 | VegBank | Robert K. Peet | Michael T. Lee | 6456 | 121 |
| NA-US-006 | Carolina Vegetation Survey Database | Robert K. Peet | Michael T. Lee | 2317 | 122 |
| NA-US-014 | Alaska-Arctic Vegetation Archive | Donald A. Walker | Amy Breen | 467 | 123 |
| SA-00-002 | VegPáramo | Gwendolyn Peyre | Xavier Font | 1591 | 124 |
| SA-AR-002 | Vegetation Database of Central Argentina | Melisa Giorgis | Alicia T.R. Acosta | 42 | |
| SA-BO-003 | Bolivia Forest Plots | Michael Kessler | Sebastian Herzog | 18 | |
| SA-BR-002 | Forest Inventory, State of Santa Catarina, Brazil (IFFSC Project) | Alexander Christian Vibrans | André Luís de Gasper | 1345 | 125 |
| SA-BR-003 | Grasslands of Rio Grande do Sul, Brazil | Eduardo Vélez-Martin | Valério De Patta Pillar | 271 | |
| SA-BR-004 | Grassland Database of Campos Sulinos | Gerhard E. Overbeck | Valério De Patta Pillar | 111 | |
| SA-CL-002 | SSAForests_Plots_db | Alvaro G. Gutierrez | | 163 | |
| SA-CL-003 | Chilean Park Transects - Fondecyt 1040528 | Aníbal Pauchard | Alicia Marticorena | 33 | 126 |
| SA-EC-001 | Ecuador Forest Plot Database | Jürgen Homeier | | 156 | |

Table 2: Description of the variables contained in the ‘header’ matrix, together with their range (if numeric) or possible levels (if nominal or binary), and the number of non-empty (i.e., non NA) records. Variable types can be n - nominal (i.e., qualitative variable), o - ordinal, q - quantitative, or b - binary (i.e., boolean), or d - date. . Additional details on the variables is in Bruelheide et al. (2019) [23]. GIVD codes derive from Dengler et al. (2011) [34]. Biomes refer to Schultz 2005 [127], modified to include also the world mountain regions by Körner et al. (2017)[128]. The column ESY refers to the EUNIS Habitat Classification Expert system described in Chytrý et al. (2020) [47].

| Variable | Range/Levels | Unit of Measurement | Nr. non-NA Records | Type |
|----------------------|---|---------------------|--------------------|------|
| GIVD_ID | | | 91205 | n |
| Dataset | | | 91205 | n |
| Continent | Africa, Asia, Europe, North America, Oceania, South America | | 91205 | n |
| Country | | | 91205 | n |
| Biome | Alpine, Boreal zone, Dry midlatitudes, Dry tropics and subtropics, Polar and subpolar zone, Subtrop. with year-round rain, Subtropics with winter rain, Temperate midlatitudes, Tropics with summer rain, Tropics with year-round rain | | 91205 | n |
| Date_of_recording | 1888-07-05 - 2015-02-03 | dd-mm-yyyy | 75971 | d |
| Latitude | -54.73863 - 80.149116 | ° (WGS84) | 91205 | q |
| Longitude | -162.741433 - 179.590053 | ° (WGS84) | 91205 | q |
| Location_uncertainty | 1 - 2500 | m | 91176 | q |
| Releve_area | 0.01 - 40000 | m ² | 62063 | q |
| Plant_recorded | All vascular plants, All trees & dominant understory, Dominant trees, Only dominant species, Dominant woody plants >= 2.5 cm dbh, All woody plants, Woody plants >= 1 cm dbh, Woody plants >= 2.5 cm dbh, Woody plants >= 5 cm dbh, Woody plants >= 10 cm dbh, Woody plants >= 20 cm dbh, Woody plants >= 1 m height, Not specified | | 91205 | n |
| Elevation | -25 - 4819 | m a.s.l. | 52277 | q |
| Aspect | 0 - 360 | ° | 30842 | q |
| Slope | 0 - 90 | ° | 37817 | q |
| is_forest | FALSE = 58145; TRUE = 25740 | | 83885 | b |
| ESY | | | 55631 | n |
| Forest | FALSE = 50071; TRUE = 23979 | | 74050 | b |
| Shrubland | FALSE = 62967; TRUE = 11083 | | 74050 | b |
| Grassland | FALSE = 26974; TRUE = 47076 | | 74050 | b |
| Wetland | FALSE = 55970; TRUE = 18080 | | 74050 | b |
| Sparse_vegetation | FALSE = 62728; TRUE = 11322 | | 74050 | b |
| Cover_total | 1 - 313 | % | 24850 | q |
| Cover_tree_layer | 0.5 - 150 | % | 7270 | q |
| Cover_shrub_layer | 0.5 - 145 | % | 10209 | q |
| Cover_herb_layer | 0.2 - 180 | % | 26846 | q |

| Variable | Range/Levels | Unit of Measurement | Nr. non-NA Records | Type |
|-----------------------|-------------------------------|---------------------|--------------------|------|
| Cover_moss_layer | 1 - 100 | % | 9685 | q |
| Cover_lichen_layer | 1 - 95 | % | 739 | q |
| Cover_algae_layer | 1 - 100 | % | 221 | q |
| Cover_litter_layer | 1 - 100 | % | 4510 | q |
| Cover_bare_rocks | 1 - 100 | % | 1904 | q |
| Cover_cryptogams | 1 - 95 | % | 593 | q |
| Cover_bare_soil | 0.1 - 99 | % | 1414 | q |
| Height_trees_highest | 1 - 99 | m | 6140 | q |
| Height_trees_lowest | 1 - 90 | m | 246 | q |
| Height_shrubs_highest | 0.1 - 9.9 | m | 2902 | q |
| Height_shrubs_lowest | 0.1 - 9 | m | 350 | q |
| Height_herbs_average | 0.1 - 440 | cm | 10161 | q |
| Height_herbs_lowest | 1 - 250 | cm | 2809 | q |
| Height_herbs_highest | 1 - 600 | cm | 1744 | q |
| Naturalness | 1 = Natural, 2 = Semi-natural | | 68179 | o |