

sPlot open - An environmentally-balanced, open-access, global dataset of vegetation plots

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Abstract

Vegetation provides the foundation of life on Earth. Assessing biodiversity status and trends in plant communities is therefore critical to understand and quantify the effects of global change on ecosystems. Here, we present the largest dataset of vegetation plots (i.e. species co-occurrence or community composition data) ever released in open access. It contains information on 91,031 vegetation plots recording the cover or abundance of each plant species that occurs in a plot of a given surface area at the date of the botanical survey. Plots were derived from 103 local to regional datasets. To improve the representation of Earth's environmental conditions, plots were resampled from a larger pool of vegetation plots using an environmentally stratified sampling design. Each vegetation plot comes with information on community-weighted means and variances of key plant functional traits. Our open-access dataset can be used to explore global patterns of diversity at the plant community level, as ground truthing data in remote sensing applications or as a baseline for biodiversity monitoring.

Background & Summary

Biodiversity is facing a global crisis (1). As many as 1 million species are estimated to be already facing extinction, mostly as a consequence of anthropogenic impacts, land-use and climate change (1). The rates of biodiversity redistribution and homogenization are also accelerating (2; 3). Biological assemblages are becoming progressively more similar to each other globally, as local biodiversity and endemic species go extinct and are replaced by introduced exotic species or by more widespread and competitive native species (1; 3). This has profound potential impacts on human and ecosystem health (4; 5). For instance, many terrestrial and marine species are shifting their geographical distribution as a response to climate change (2), including animals hosting pathogens transmissible to humans (6; 7; 8).

Vegetation, i.e., the assemblage of plant species, is no exception to this biodiversity crisis (9; 10; 3). This is worrisome, since terrestrial vegetation accounts for 80% (450 Gt C) of the living biomass on Earth (11). Given the central role of vegetation in ecosystem productivity, stability and functioning (10), assessing biodiversity status and trends in plant communities is paramount, for other life compartments and human societies alike.

Monitoring plant biodiversity trends requires adequate data across a range of scales (12). Large independent collections of plant occurrence data do exist at the global or continental extent via the Botanical Information and Ecology Network (BIEN) (13), the Global Inventory of Floras and Traits (GIFT) (14) or the Global Biodiversity Information Facility (GBIF) (<https://www.gbif.org/>). However, all these occurrence-only databases either neglect how individual plant species co-occur and interact locally to form plant communities, or are collected at spatial resolutions (e.g., one-degree grid cells) which are too coarse to assess biodiversity trends at the most relevant scale of local plant communities (15).

Yet, there is a long-lasting 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. 16). Compared to species-level data, vegetation-plot data present many advantages. First, they contain information on which plant species co-occur together in the same locality at a given moment in time (17). This built-in feature of vegetation plots is a necessary prerequisite for testing hypotheses related to biotic interactions among plant species (i.e. plant-plant interactions). It can also provide crucial information on where and when a species is absent, therefore improving current species distribution models (18). Being spatially explicit, vegetation plots can be resurveyed through time to assess potential changes in plant species composition relative to a baseline (19; 20, 3). As they normally contain also information on the relative cover or abundance of each species, vegetation plots are more adequate to detect subtle biodiversity changes, compared to data based on the occurrence of individual species only (21).

Vegetation-plot data are very fragmented, though, as they typically stem from a myriad of research projects. As such, these data often suffer from the usual trade-off in biodiversity data: Collections have either fine-grain spatial resolutions but small spatial extents, or vice versa (22). Furthermore, with their disparate sampling protocols, standards and taxonomic resolutions, aggregating and harmonizing vegetation plot data proves extremely challenging (23). It is not surprising, therefore, that these data have only been rarely used in global-scale biodiversity research until recently (24; 25).

The sPlot initiative tries to close this data gap. It leverages on several existing local to regional vegetation-plot datasets, to create a harmonized and comprehensive global geo-database of terrestrial plant species assemblages (26). Established in 2013, sPlot currently contains more than 1.9 million vegetation plots, and is fully integrated with the TRY database (27), 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 ([28](#)), the mechanisms underlying the spread and abundance of native vs. invasive tree species ([29](#)), and worldwide trait–environment relationships in plant communities ([23](#)).

Here, we provide an open-access data set composed of 91,031 plots, which is representative of the environmental space covered by the sPlot database. Plots stem from 103 databases, and span across 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 species composition of each vegetation plot, including species cover or abundance information when available; and (3) community-level functional diversity indices derived from the TRY database ([27](#)).

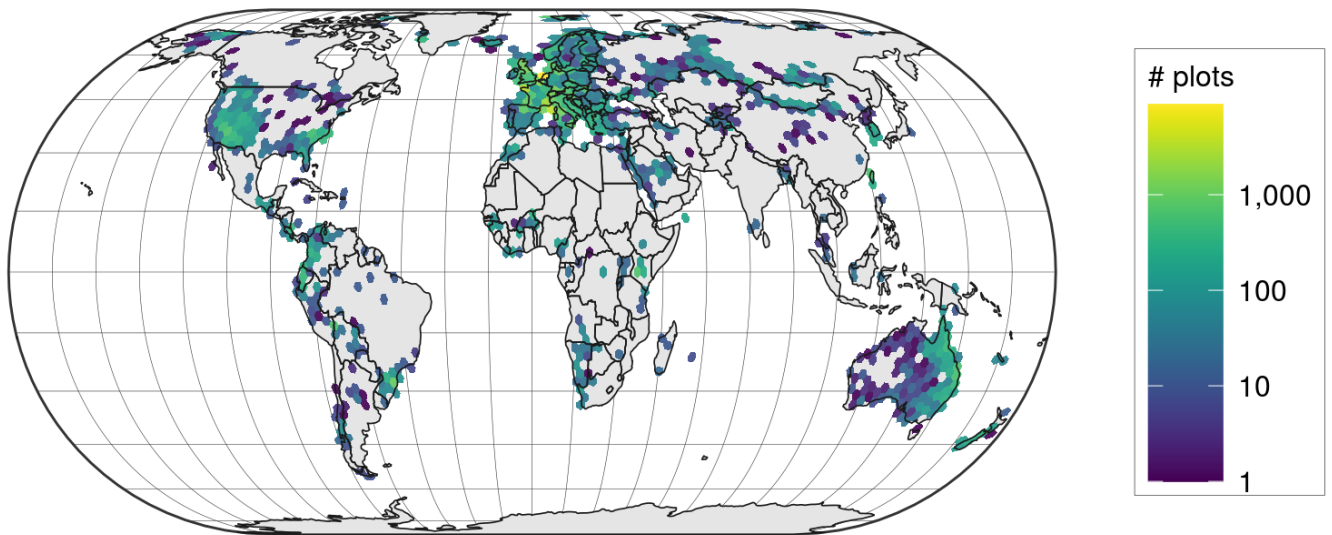


Figure 1: Global map of sPlot Open ($n = 91,031$) and spatial distribution of vegetation plot density per hexagonal cell with a spatial resolution of approximately 70,000 km². Map projection is Eckert IV.

Methods

Vegetation plot data sources

We started from the sPlot database v2.1 (created October 2016), which contains 1,121,244 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 [26](#) for more information). For instance: 48 vegetation-plot datasets derive from the European Vegetation Archive (EVA) ([17](#)), three major African datasets from the Tropical African Vegetation Archive (TAVA), multiple vegetation datasets in the USA from the VegBank archive ([30](#); [31](#)). Data from other continents (South America, Asia) or countries were contributed as separate datasets. The metadata of each of the 110 vegetation-plot datasets stored in sPlot are managed through the Global Index of Vegetation-Plot Databases (GIVD; [32](#)), using the GIVD identifier as the unique dataset identifier.

Resampling method

Data in the sPlot database are unevenly distributed across continents and biomes (see [23](#)). Mid-latitude regions in developing 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., [33](#); [2](#)). To reduce this imbalance to the extent possible, we performed a stratified resampling approach, using several environmental variables available at the global extent as sampling strata. We considered 30 climatic and soil variables. For climate we complemented the 19 bioclimatic variables from CHELSA ([34](#)), 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 - [35](#)). For soil, we extracted seven variables from the SOILGRIDS database ([36](#)), 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) of 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 environmental space represented by the first two principal components (PC1-PC2), accounting for 47% and 23% of the total variation on PC1 and PC2, respectively, into a 100 × 100 grid. This PC1-PC2 bidimensional space was subsequently used to balance our sampling effort across all PC1-PC2 grid cells for which vegetation plots are available. Before projecting vegetation plots from the sPlot database v2.1 onto this PC1-PC2 environmental space, we removed vegetation plots: from wetlands; from anthropogenic vegetation types; without geographical coordinates; and with a location uncertainty higher than 3 km for those having geographical coordinates. This led to 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 randomly selected up to 50 vegetation plots using the heterogeneity-constrained random resampling algorithm from [[37](#)]. 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 ([38](#)) between all possible pairs of vegetation plots for a given random selection of 50 vegetation plots ($n = 1225$). We chose this

dissimilarity index because it is not influenced by differences in species richness among vegetation plots. 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 the 1,000 random selections according to the mean (ascending order) and variance (descending order) value. Ranks from both sortings were summed for each random selection, and the random selection with the lowest summed rank was considered as the most representative of the focal grid cell. In case a grid cell contained fewer than 50 plots, we retained all of them. In this way, we reduced the imbalance towards over-sampled climate types, while ensuring the resampled dataset to be representative of the entire environmental gradient covered by the sPlot database. We repeated the resampling procedure three times to get three different possibilities of a random 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 the 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). Being the sPlot database 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, for instance because 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 vegetation plot in the reserve should belong to the same environmental strata, i.e., the same PC1-PC2 grid cell, of the confidential vegetation plot. Note that a given PC1-PC2 grid cell may have one or more confidential vegetation plots (max = xx) 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 ([27](#)). These traits were selected among those traits that describe the leaf, wood and seed economics spectra ([39](#); [40](#)), and are known to either affect different key ecosystem processes or respond to macroclimatic drivers or both ([26](#)). The eighteen plant functional traits 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', [41](#); [42](#)). Gap-filling was performed at the level of individual observations. We then loge-transformed 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 are available in [\[26\]](#).

Community-weighted means (CWM) and the variances (CWV) were calculated for every plant functional trait j and every vegetation plot k as follows (43):

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

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

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

The final dataset that is provided here as open access contains 91,031 vegetation plots from 115 countries and all continents except Antarctica (Figure 1) and stems from 103 constitutive datasets (Table 1). Information on the size (surface area) of the vegetation survey is available for 61,898 vegetation plots, and ranges between 0.01 m² and 4 ha (mean = 270 m²; median = 78.5 m²). 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). Most plots only include information on vascular plants, while a minority also includes information on lichens (n = 3,045) or mosses (n = 4,963). By reducing the overrepresentation of vegetation plots in specific 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, Africa and South America have only 4507 and 5515 vegetation plots, respectively. The representation of biomes is equally unbalanced. The biomes 'Temperate midlatitudes' and 'Subtropics with winter rain' have 37,507 and 16,510 vegetation plots, respectively, while none of the other biomes have more than 10,000 vegetation plots (Figure 2).



Figure 2: Distribution in environmental space (R package plotbiomes by Valentin Stefan).

Finally, the dataset contains a relatively balanced number of forest (n = 25,832) vs. non forest (n = 38,203) vegetation plots, with a minor proportion of plots remaining unassigned (n = 10,050). The assignment of plots to forests and non-forests is 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 a forest if the cover of the tree layer, or alternatively, the sum of relative cover of all tree taxa, was greater than 0.25. It was instead

considered a non-forest record if the sum of relative cover of low-stature, non-tree and non-shrub taxa was greater than 0.90. For an extensive explanation on this classification scheme, we refer the reader to [\[26\]](#). 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 individual 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 invite potential users to carefully read the description of each individual dataset before using this open-access dataset.

Database Organization

The open-access dataset is organized into three matrices.

The *'header'* matrix contains plot level information for the 91,031 vegetation plots provided in this open access dataset, 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). A brief description of all the xx variables contained 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 1,607,826 records, from 39,922 taxa, mostly resolved at the species level. For each record we report both the taxon name as originally contributed by the data custodian (column *'Matched_concept'*), and the taxon name after taxonomic standardization (column *'Species_name_harmonized'*). For each entry, 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, which often is given on a cover//abundance scale (column *'Cover'*), and a *'Relative_cover'* field, i.e., the cover//abundance of each taxon in each vegetation layer divided by the total cover//abundance of all taxa in that vegetation layer. Finally, for each entry, we provide a *'Taxon_group'* field, reporting whether the corresponding taxa is a vascular plant, moss, lichen or alga.

Finally the *'CWM'* 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 only in one plot we have no functional trait information for any of the occurring species. 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.

Technical Validation

The sPlot database has a nested structure, and is composed of several individual datasets, each validated and maintained by its respective dataset custodian. Each individual dataset also has individual vegetation plots, each provided by its owner (the person who performed the actual vegetation survey) or by someone who digitized the original data from the scientific or grey literature. We obviously have no direct control on the individual vegetation plots that we provide here in an open access dataset. Yet, each of these vegetation plots are stemming 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 having been integrated into the sPlot database, each dataset was further checked for consistency and, if having a different format, was converted to a Turboveg 2 database ([44](#)). During this conversion into a Turboveg format, we checked that all datasets contained the required metadata information and we converted this information to the sPlot database standards, if necessary. Furthermore we cross-checked that each plot is located within the geographic scopes of its respective dataset. Finally, we harmonized all the taxonomic names from a dataset, based on the sPlot's taxonomic backbone (Purschke 2017). This backbone matched all the taxonomic names (without nomenclatural authors) from all datasets in sPlot 2.1 and TRY v3.0 ([27](#)) to their resolved version based on the Taxonomic Name Resolution Service web application (TNRS version 4.0; [45](#); iPlant Collaborative, 2015). This allowed to (1) harmonize all datasets to a common nomenclature, and (2) to link the sPlot database to the TRY database ([27](#)). All taxa originally denoted at taxonomic ranks lower than species, were aggregated at species level. Additional detail on the taxonomic resolution is reported in [[26](#)], while a description of the workflow, including R-code, is available in [[46](#)]

Usage Notes

The sPlot Open database can be downloaded from <https://www.idiv.de> (link to PlantHub). The use of data contained in BioTIME should cite original data citations in addition to the present paper. The data included in the present paper represent the subset of sPlot for which we were able to secure licences for making these data open. The additional studies 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 found here (<https://portal.idiv.de/nextcloud/index.php/s/YjMZtwFDwtoefGi>).

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Author contributions

FMS wrote the first draft of the manuscript, with considerable input by JL and HB. JL and TH wrote the resampling algorithm. FMS set up the GitHub project 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. All authors contributed to revising the manuscript.

Competing interests

[A competing interests statement is required for all papers accepted by and published in Scientific Data. If there is no conflict of interest, a statement declaring this must still be included in the manuscript.]

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

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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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Viktoria Wagner, Toby Spribille, Stefan Abrahamczyk, Erwin Bergmeier

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Supplementary Material

Table 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	DB_name GVID	Nr Plots	Contributed plots	Citation
00-00-004	Vegetation Database of Eurasian Tundra	1132	600	
00-RU-003	Database Meadows and Steppes of Southern Ural	2354	99	
00-TR-001	Forest Vegetation Database of Turkey - FVDT	919	15	
EU-00-002	Nordic-Baltic Grassland Vegetation Database (NBGVD)	7675	931	47
EU-00-011	Vegetation-Plot Database of the University of the Basque Country (BIOVEG)	18441	1694	48
EU-00-013	Balkan Dry Grasslands Database	7683	224	49
EU-00-016	Mediterranean Ammophiletea Database	7359	3713	50
EU-00-017	European Coastal Vegetation Database	4624	1369	
EU-00-018	The Nordic Vegetation Database	5477	1755	51
EU-00-019	Balkan Vegetation Database	9118	211	52
EU-00-020	WetVegEurope	14111	61	53
EU-00-022	European Mire Vegetation Database	10147	1843	54
EU-AL-001	Vegetation Database of Albania	290	99	55
EU-AT-001	Austrian Vegetation Database	34458	950	56
EU-BE-002	INBOVEG	25665	48	
EU-BG-001	Bulgarian Vegetation Database	5254	74	57
EU-CH-005	Swiss Forest Vegetation Database	14193	1409	58
EU-CZ-001	Czech National Phytosociological Database	10469 7	579	59
EU-DE-001	VegMV	53822	5	60
EU-DE-013	VegetWeb Germany	23078	199	61

GIVD ID	DB_name GIVD	Nr Plots	Contributed plots	Citation
EU-DE-014	German Vegetation Reference Database (GVRD)	30840	286	62
EU-DK-002	National Vegetation Database of Denmark	24264	1181	
EU-ES-001	Iberian and Macaronesian Vegetation Information System (SIVIM) - Wetlands	6560	292	
EU-FR-003	SOPHY	20986 4	13322	
EU-GB-001	UK National Vegetation Classification Database	28533	5457	
EU-GR-001	KRITI	292	43	
EU-GR-005	Hellenic Natura 2000 Vegetation Database (HelNatVeg)	5168	777	63
EU-GR-006	Hellenic Woodland Database	3199	4	64
EU-HR-001	Phytosociological Database of Non-Forest Vegetation in Croatia	5057	213	65
EU-HR-002	Croatian Vegetation Database	8734	688	
EU-HU-003	CoenoDat Hungarian Phytosociological Database	8505	17	66
EU-IT-001	VegItaly	15332	2712	67
EU-IT-010	Italian National Vegetation Database (BVN/ISPRA)	3562	155	68
EU-IT-011	Vegetation-Plot Database Sapienza University of Rome (VPD-Sapienza)	12780	1003	69
EU-LT-001	Lithuanian Vegetation Database	7821	119	
EU-LV-001	Semi-natural Grassland Vegetation Database of Latvia	5594	306	70
EU-MK-001	Vegetation Database of the Republic of Macedonia	1417	10	
EU-NL-001	Dutch National Vegetation Database	10232 7	10223	71
EU-PL-001	Polish Vegetation Database	22229	464	72
EU-RO-007	Romanian Forest Database	6017	60	73
EU-RO-008	Romanian Grassland Database	1921	44	74
EU-RS-002	Vegetation Database Grassland Vegetation of Serbia	5587	57	75

GIVD ID	DB_name GIVD	Nr Plots	Contributed plots	Citation
EU-RU-002	Lower Volga Valley Phytosociological Database	14853	149	76
EU-RU-003	Vegetation Database of the Volga and the Ural Rivers Basins	1516	96	77
EU-RU-011	Vegetation Database of Tatarstan	7471	94	78
EU-SI-001	Vegetation Database of Slovenia	10986	435	79
EU-SK-001	Slovak Vegetation Database	36405	893	80
EU-UA-006	Vegetation Database of Ukraine and Adjacent Parts of Russia	3326	479	
AF-00-001	West African Vegetation Database	3129	184	81
AF-00-008	PANAF Vegetation Database	2469	942	
AF-BF-001	Sahel Vegetation Database	1079	279	82
00-00-001	ForestPlots.net	1827	108	83
00-00-003	SALVIAS	4883	2860	
00-00-005	Tundra Vegetation Plots (TundraPlot)	577	227	84
00-RU-002	Database of Masaryk University`s Vegetation Research in Siberia	1547	128	85
AF-00-003	BIOTA Southern Africa Biodiversity Observatories Vegetation Database	1666	562	86
AF-00-006	SWEA-Dataveg	2704	1211	
AF-00-009	Vegetation Database of the Okavango Basin	590	202	87
AF-CD-001	Forest Database of Central Congo Basin	292	97	88
AF-ET-001	Vegetation Database of Ethiopia	74	59	89
AF-MA-001	Vegetation Database of Southern Morocco	1337	266	90
AF-ZW-001	Vegetation Database of Zimbabwe	36	17	91
AS-00-001	Korean Forest Database	4885	766	92
AS-00-003	Vegetation of Middle Asia	1381	128	93

GIVD ID	DB_name GIVD	Nr Plots	Contributed plots	Citation
AS-00-004	Rice Field Vegetation Database	179	31	
AS-BD-001	Tropical Forest Dataset of Bangladesh	211	82	
AS-CN-001	China Forest-Steppe Ecotone Database	148	97	94
AS-CN-002	Tibet-PaDeMoS Grazing Transect	146	27	95
AS-CN-003	Vegetation Database of the BEF China Project	27	18	96
AS-CN-004	Vegetation Database of the Northern Mountains in China	485	70	
AS-EG-001	Vegetation Database of Sinai in Egypt	926	98	97
AS-ID-001	Sulawesi Vegetation Database	24	24	
AS-IR-001	Vegetation Database of Iran	2335	105	
AS-KZ-001	Database of Meadow Vegetation in the NW Tien Shan Mountains	94	3	98
AS-MN-001	Southern Gobi Protected Areas Database	1516	688	99
AS-RU-001	Wetland Vegetation Database of Baikal Siberia (WETBS)	2381	6	100
AS-RU-002	Database of Siberian Vegetation (DSV)	9116	2150	
AS-RU-004	Database of the University of Münster - Biodiversity and Ecosystem Research Group's Vegetation Research in Western Siberia and Kazakhstan	445	85	
AS-SA-001	Vegetation Database of Saudi Arabia	919	607	
AS-TJ-001	Eastern Pamirs	282	174	101
AS-TW-001	National Vegetation Database of Taiwan	930	897	
AS-YE-001	Socotra Vegetation Database	396	190	102
AU-AU-002	AEKOS	21261	7443	103
AU-NC-001	New Caledonian Plant Inventory and Permanent Plot Network (NC-PIPPN)	201	98	104
AU-NZ-001	New Zealand National Vegetation Databank	1895	983	105
AU-PG-001	Forest Plots from Papua New Guinea	63	53	106

GIVD ID	DB_name GIVD	Nr Plots	Contributed plots	Citation
NA-00-002	Tree Biodiversity Network (BIOTREE-NET)	1757	208	107
NA-CA-003	Database of Timberline Vegetation in NW North America	110	38	108
NA-CA-004	Understory of Sugar Maple Dominated Stands in Quebec and Ontario (Canada)	156	9	109
NA-CA-005	Boreal Forest of Canada	89	44	
NA-GL-001	Vegetation Database of Greenland	664	340	110
NA-US-002	VegBank	67352	6456	111
NA-US-006	Carolina Vegetation Survey Database	17221	2317	112
NA-US-014	Alaska-Arctic Vegetation Archive	1363	467	113
SA-00-002	VegPáramo	2643	1591	114
SA-AR-002	Vegetation Database of Central Argentina	218	42	
SA-BO-003	Bolivia Forest Plots	75	18	
SA-BR-002	Forest Inventory, State of Santa Catarina, Brazil (IFFSC Project)	1669	1345	115
SA-BR-003	Grasslands of Rio Grande do Sul, Brazil	320	271	
SA-BR-004	Grassland Database of Campos Sulinos	161	111	
SA-CL-002	SSAForests_Plots_db	261	163	
SA-CL-003	Chilean Park Transects - Fondecyt 1040528	165	33	116
SA-EC-001	Ecuador Forest Plot Database	172	156	

Table Table 2: Description of the variables contained in the 'header' matrix, together with their range (if numeric) or possible levels (if nominal or boolean). Variable type can be c - character (i.e. text), f - factor (i.e. qualitative or ordinal variable), i - integer (e.g. binomial), n - numeric (i.e., double) or l - logical (i.e., boolean).

Variable	Range/Levels	Unit of Measurement	Nr. Records	Type
GIVD_ID			91031	character
Dataset			91031	character
Continent	Africa, Asia, Australia, Europe, North America, Oceania, South America		90729	factor

Variable	Range/Levels	Unit of Measurement	Nr. Records	Type
Country			91031	character
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		91031	factor
Date	-29764 - 16469		75798	numeric
Latitude	-54.73863 - 80.149116	° (WGS84)	91031	numeric
Longitude	-162.741433 - 179.590053	° (WGS84)	91031	numeric
Location_uncertainty	1 - 2500	m	91002	integer
Releve_area	0.01 - 40000	m ²	61898	numeric
Herbs_identified	FALSE = 4876; TRUE = 6323		11199	logical
Plant_recorded	All trees & dominant understory, All vascular plants, All vascular plants and dominant cryptogams, All woody plants, Dominant trees, Only dominant species, Dominant woody plants >= 2.5 cm dbh, Woody plants >= 10 cm dbh, Woody plants >= 1 m height, Woody plants >= 1 cm dbh, Woody plants >= 20 cm dbh, Woody plants >= 2.5 cm dbh, Woody plants >= 5 cm dbh, NA		91015	factor
Elevation	-25 - 4819	m a.s.l.	52121	numeric
Aspect	0 - 360	°	30796	numeric
Slope	0 - 99	°	37784	numeric
is_forest	FALSE = 20396; TRUE = 25832		46228	logical
is_nonforest	FALSE = 50870; TRUE = 38203		89073	logical
ESY			55457	character
Naturalness	1 - 2		68011	integer
Forest	FALSE = 38295; TRUE = 23735		62030	logical
Shrubland	FALSE = 38233; TRUE = 11081		49314	logical
Grassland	FALSE = 10213; TRUE = 46947		57160	logical
Sparse_vegetation	FALSE = 33381; TRUE = 11315		44696	logical
Wetland	FALSE = 29078; TRUE = 18038		47116	logical
Cover_total	1 - 313	%	24712	integer

Variable	Range/Levels	Unit of Measurement	Nr. Records	Type
Cover_tree_layer	0.5 - 150	%	7245	numeric
Cover_shrub_layer	0.5 - 145	%	10197	numeric
Cover_herb_layer	0.2 - 180	%	26679	numeric
Cover_moss_layer	1 - 100	%	9643	integer
Cover_lichen_layer	1 - 95	%	734	integer
Cover_algae_layer	1 - 100	%	221	integer
Cover_litter_layer	1 - 100	%	4500	integer
Cover_bare_rocks	1 - 100	%	1897	integer
Cover_cryptogams	1 - 95	%	593	integer
Cover_bare_soil	0.1 - 99	%	1412	numeric
Height_trees_highest	1 - 99	m	6115	numeric
Height_trees_lowest	1 - 90	m	221	numeric
Height_shrubs_highest	0.1 - 9.9	m	2880	numeric
Height_shrubs_lowest	0.1 - 9	m	328	numeric
Height_herbs_average	0.1 - 440	cm	10125	numeric
Height_herbs_lowest	1 - 250	cm	2785	integer
Height_herbs_highest	1 - 600	cm	1733	integer