

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

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## Authors

✉ Francesco Maria Sabatini<sup>1,2</sup>, Jonathan Lenoir<sup>3</sup>, Tarek Hattab<sup>4</sup>, Elise Arnst<sup>5</sup>, Milan Chytrý<sup>6</sup>, Jürgen Dengler<sup>7,8,9</sup>, Valério De Patta Pillar<sup>10</sup>, Patrice De Ruffray<sup>11</sup>, Stephan M. Hennekens<sup>12</sup>, Ute Jandt<sup>2</sup>, Florian Jansen<sup>13</sup>, Borja Jiménez-Alfaro<sup>14</sup>, Jens Kattge<sup>15</sup>, Aurora Levesley<sup>16</sup>, Oliver Purschke<sup>17</sup>, Brody Sandel<sup>18</sup>, Fahmida Sultana<sup>19</sup>, Marten Winter<sup>9</sup>, Svetlana Aćić<sup>20</sup>, Emiliano Agrillo<sup>21</sup>, Miguel Alvarez<sup>22</sup>, Iva Apostolova<sup>23</sup>, Mohammed A.S. Arfin Khan<sup>24</sup>, Isabelle Aubin<sup>25</sup>, Yves Bergeron<sup>26</sup>, Erwin Bergmeier<sup>27</sup>, Idoia Biurrun<sup>28</sup>, Anne D. Bjorkman<sup>29</sup>, Laura Casella<sup>30</sup>, Luis Cayuela<sup>31</sup>, Tomáš Černý<sup>32</sup>, Victor Chepinoga<sup>33</sup>, János Csiky<sup>34</sup>, Renata Čušterevska<sup>35</sup>, Els De Bie<sup>36</sup>, Michele De Sanctis<sup>21</sup>, Panayotis Dimopoulos<sup>37</sup>, Mohamed Abd El-Rouf Mousa El-Sheikh<sup>38,39</sup>, Brian Enquist<sup>40</sup>, Manfred Finckh<sup>41</sup>, Emmanuel Garbolino<sup>42</sup>, Melisa Giorgis<sup>43</sup>, Valentin Golub<sup>44</sup>, Alvaro G. Gutierrez<sup>45</sup>, Mohamed Z. Hatim<sup>46</sup>, Norbert Hölzel<sup>47</sup>, Jürgen Homeier<sup>48</sup>, Adrian Indreica<sup>49</sup>, Kim Sarah Jacobsen<sup>50</sup>, John Janssen<sup>12</sup>, Birgit Jedrzejek<sup>51</sup>, Norbert Jürgens<sup>41</sup>, Zygmunt Kaćki<sup>52</sup>, Ali Kavgaci<sup>53</sup>, Michael Kessler<sup>54</sup>, Andrey Korolyuk<sup>55</sup>, Hjalmar Kühl<sup>9,56</sup>, Flavia Landucci<sup>6</sup>, Ching-Feng Li<sup>57</sup>, Hongyan Liu<sup>58</sup>, Tatiana Lysenko<sup>59</sup>, Corrado Marcenò<sup>28</sup>, Jesper Erenskjold Moeslund<sup>60</sup>, Jonas V. Müller<sup>61</sup>, Jérôme Munzinger<sup>62</sup>, Jalil Noroozi<sup>63</sup>, Arkadiusz Nowak<sup>64</sup>, Viktor Onyshchenko<sup>65</sup>, Gerhard E. Overbeck<sup>66</sup>, Aníbal Pauchard<sup>67</sup>, Robert K. Peet<sup>68</sup>, Aaron Pérez-Haase<sup>69,70</sup>, Tomáš Peterka<sup>6</sup>, Gwendolyn Peyre<sup>71</sup>, Oliver L. Phillips<sup>16</sup>, Vadim Prokhorov<sup>72</sup>, Valerijus Rašomavičius<sup>73</sup>, Rasmus Revermann<sup>41</sup>, John S. Rodwell<sup>74</sup>, Eszter Ruprecht<sup>75</sup>, Solvita Rūsiņa<sup>76</sup>, Cyrus Samimi<sup>77</sup>, Joop H.J. Schaminée<sup>12</sup>, Marco Schmidt<sup>78</sup>, Urban Šilc<sup>79</sup>, Željko Škvorc<sup>80</sup>, Anita Smyth<sup>81</sup>, Zvezdana Stančić<sup>82</sup>, Zhiyao Tang<sup>58</sup>, Ioannis Tsiripidis<sup>83</sup>, Milan Valachovič<sup>84</sup>, Kim André Vanselow<sup>85</sup>, Kiril Vassilev<sup>23</sup>, Eduardo Vélez-Martin<sup>86</sup>, Roberto Venanzoni<sup>87</sup>, Alexander Christian Vibrans<sup>88</sup>, Risto Virtanen<sup>9,89,90</sup>, Henrik von Wehrden<sup>91</sup>, Viktoria Wagner<sup>92</sup>, Donald A. Walker<sup>93</sup>, Desalegn Wana<sup>94</sup>, Karsten Wesche<sup>9,95,96</sup>, Timothy Whitfeld<sup>97</sup>, Wolfgang Willner<sup>98</sup>, Susan Wiser<sup>5</sup>, Thomas Wohlgemuth<sup>99</sup>, Sergey Yamalov<sup>100</sup>, Helge Bruelheide<sup>1,2</sup>,

✉ — To whom correspondence should be addressed: francesco.sabatini@botanik.uni-halle.de

1. German Centre for Integrative Biodiversity Research (iDiv) - Halle-Jena-Leipzig. Germany
2. Martin-Luther University Halle-Wittenberg, Institute of Biology, Am Kirchtor 1, 06108, Halle, Germany
3. Unité de Recherche "Ecologie et Dynamique des Systèmes Anthropisés" (EDYSAN), UMR 7058 CNRS, Université de Picardie Jules Verne, 80037 Amiens Cedex 1, France
4. MARBEC, University of Montpellier, CNRS, IFREMER and IRD, Sète, France
5. Manaaki Whenua – Landcare Research, PO Box 69040, 7640, Lincoln, New Zealand
6. Masaryk University, Department of Botany and Zoology, Kotlářská 2, 611 37, Brno, Czech Republic
7. Zurich University of Applied Sciences (ZHAW), Vegetation Ecology Group, Institute of Natural Resource Sciences (IUNR), Grüentalstr. 14, 8820, Wädenswil, Switzerland
8. University of Bayreuth, Plant Ecology, Bayreuth Center of Ecology and Environmental Research (BayCEER), Universitätsstr. 30, 95447, Bayreuth, Germany
9. German Centre for Integrative Biodiversity Research (iDiv) Halle-Jena-Leipzig, Deutscher Platz 5e, 04103, Leipzig, Germany
10. Federal University of Rio Grande do Sul, Ecology, Av. Bento Gonçalves 9500, 91501-970, Porto Alegre, Brazil
11. IBMP, 12, rue du Général-Zimmer, 67084, Strasburg, France
12. Wageningen University and Research, Wageningen Environmental Research (Alterra), P.O.Box 47, 6700 AA, Wageningen, Netherlands

13. University of Rostock, Faculty of Agricultural and Environmental Sciences, Justus-von-Liebig-Weg 6, 18059, Rostock, Germany
14. University of Oviedo, Research Unit of Biodiversity (CSIC/UO/PA), C. Gonzalo Gutiérrez Quirós s/n, 33600, Mieres, Spain
15. Max Planck Institute for Biogeochemistry, Hans Knöll Str. 10, 07745, Jena, Germany
16. University of Leeds, School of Geography, Woodhouse Lane, LS2 9JT, Leeds, United Kingdom
17. NA,
18. Aarhus University, Aarhus, Denmark
19. Shahjalal University of Science & Technology, Forestry & Environmental Science, 3114, Sylhet, Bangladesh
20. Faculty of Agriculture, Department of Agrobotany, Nemanjina 6, 11080, Belgrade-Zemun, Serbia
21. Sapienza University of Rome, Department of Environmental Biology, P.le Aldo Moro 5, 00185, Rome, Italy
22. University of Bonn, Plant Nutrition, INRES, Karlrobert-Kreiten-Str., 53115, Bonn, Germany
23. Institute of Biodiversity and Ecosystem Research, Bulgarian Academy of Sciences, Department of Plant and Fungal Diversity and Resources, Acad. Georgi Bonchev 23, 1113, Sofia, Bulgaria
24. Shahjalal University of Science & Technology, Forestry & Environmental Science, Akhalia, 3114, Sylhet, Bangladesh
25. Canadian Forest Service, Natural Resources Canada, Great Lakes Forestry Centre, 1219 Queen St. East, P6A 2E5, Sault Ste Marie (Ontario), Canada
26. Université du Québec en Abitibi-Témiscamingue, Forest Research Institute, 445 boul. de l'Université, J9X5E4, Rouyn-Noranda, Canada
27. University of Göttingen, Vegetation Ecology and Phytodiversity, Untere Karspüle 2, 37073, Göttingen, Germany
28. University of the Basque Country UPV/EHU, Plant Biology and Ecology, P.O. Box 644, 48080, Bilbao, Spain
29. Aarhus University, Section for Ecoinformatics & Biodiversity, Department of Bioscience, Ny Munkegade 114, 8000, Aarhus C, Denmark
30. ISPRA - Italian National Institute for Environmental Protection and Research, Biodiversity Conservation Department, Via Vitaliano Brancati, 60, 00144, Roma, Italy
31. Universidad Rey Juan Carlos, Department of Biology, Geology, Physics and Inorganic Chemistry, c/ Tulipán s/n, 28933, Móstoles, Spain
32. Czech University of Life Sciences Prague, Department of Forest Ecology, Faculty of Forestry and Wood Sciences, Kamýcká 1176, 165 21, Praha 6 - Suchbát, Czech Republic
33. V.B. Sochava Institute of Geography SB RAS, Laboratory of Physical Geography and Biogeography, Ulan-Batorskaya, 1, 664033, Irkutsk, Russian Federation
34. University of Pécs, Department of Ecology, Ifjúság u. 6., 7624, Pécs, Hungary
35. Faculty of Natural Sciences and Mathematics, Institute of Biology, Arhimedova 3, 1000, Skopje, Republic of Macedonia
36. Research Institute for Nature and Forest (INBO), Department of Biodiversity and Natural Environment, Havenlaan 88, bus 73, 1000, Brussels, Belgium
37. University of Patras, Institute of Botany, Division of Plant Biology, Department of Biology, University Campus, 26504, Patras, Greece
38. College of Science, King Saud University, Botany and Microbiology Department, P.O. Box 2455, 11451, Riyadh, Saudi Arabia
39. Damanhour University, Botany Department, Faculty of Science, Damanhour, Egypt
40. University of Arizona, Ecology and Evolutionary Biology, 1041 E. Lowell St., AZ 85721, Tucson, United States
41. University of Hamburg, Biodiversity, Ecology and Evolution of Plants, Institute for Plant Science & Microbiology, Ohnhorststr. 18, 22609, Hamburg, Germany
42. MINES ParisTech, Crisis and Risk research Centre (CRC), 1 rue Claude Daunesse, BP 207, 06904, Sophia Antipolis, France
43. Instituto Multidisciplinario de Biología Vegetal (IMBIV-CONICET), ECOLOGÍA VEGETAL Y FITOGEOGRAFÍA, Av. Vélez Sársfield 1611, 5000, Córdoba, Argentina
44. Institute of Ecology of the Volga River Basin, Laboratory of Phytocoenology, Komzina, 10, 445003, Toljatty, Russia
45. Universidad de Chile, Departamento de Ciencias Ambientales y Recursos Naturales Renovables, Facultad de Ciencias Agronomicas, Santa Rosa 11315, La Pintana, 8820808, Santiago, Chile
46. Tanta University, Botany, Faculty of Science, El Geish St., 31527, Tanta, Egypt
47. University of Muenster, Institute of Landscape Ecology, Heisenbergstr. 2, 48149, Münster, Germany
48. University of Göttingen, Plant Ecology and Ecosystems Research, Untere Karspüle 2, 37073, Göttingen, Germany
49. Transilvania University of Brasov, Department of Silviculture, Sirul Beethoven 1, 500123, Brasov, Romania
50. Ghent University, Dept Environment, Coupure Links 653, 9000, Ghent, Belgium
51. University of Münster, Institute of Landscape Ecology, Heisenbergstr. 2, 48149, Münster, Germany
52. University of Wrocław, Botanical Garden, Sienkiewicza 23, 50-335, Wrocław, Poland
53. Southwest Anatolia Forest Research Institute, Silviculture and Forest Botany, POB 264, 07002, Antalya, Turkey
54. University of Zurich, Department of Systematic and Evolutionary Botany, Zollikerstrasse 107, 8008, Zurich, Switzerland
55. Central Siberian Botanical Garden, Siberian Branch, Russian Academy of Sciences, Geosystem Laboratory, Zolotodolinskaya str. 101, 630090, Novosibirsk, Russian Federation
56. Max Planck Institute for Evolutionary Anthropology (MPI-EVA), Primatology, Deutscher Platz 6, 04103, Leipzig, Germany
57. National Taiwan University, School of Forestry and Resource Conservation, No. 101, Section 2, Kuang-Fu Road, 30013, Hsinchu, Taiwan
58. Peking University, College of Urban and Environmental Sciences, Yiheyuan Rd. 5, 100871, Beijing, China

59. Institute of Ecology of the Volga River Basin RAS, Dept. of the Phytodiversity Problems, Komzin str. 10, 445003, Togliatti, Russia
60. Aarhus University, Department of Bioscience, Grenaaavej 14, 8410, Roende, Denmark
61. Royal Botanic Gardens, Kew, Conservation Science, Wakehurst Place, RH17 6TN, Ardingly, West Sussex, United Kingdom
62. IRD, CIRAD, CNRS, INRA, Université Montpellier, AMAP - Botany and Modelling of Plant Architecture and Vegetation, Boulevard de la Lironde, 34398, Montpellier, France
63. University of Vienna, Department of Botany and Biodiversity Research, Rennweg 14, 1030, Vienna, Austria
64. Polish Academy of Sciences, Botanical Garden - Center for Biological Diversity Conservation, Prawdziwka 2, 02-976, Warszawa, Poland
65. National Academy of Sciences of Ukraine, M.G. Kholodny Institute of Botany, Tereshchenkivska 2, 01601, Kyiv, Ukraine
66. Universidade Federal do Rio Grande do Sul, Department of Botany, Av. Bento Gonçalves 9500, 91501-970, Porto Alegre, Brazil
67. University of Concepción, Laboratorio de Invasiones Biológicas (LIB), Victoria 631, 4030000, Concepción, Chile
68. University of North Carolina, Department of Biology, CB3280, South Road, 27599-3280, Chapel Hill, NC, United States
69. University of Barcelona, Department of Evolutionary Biology, Ecology and Environmental Sciences, Diagonal 643, 08028, Barcelona, Spain
70. Center for Advanced Studies of Blanes, Spanish Research Council (CEAB-CSIC), Continental Ecology, Carrer d'accés a la Cala St. Francesc, 14, 17300, Blanes, Girona, Spain
71. University of the Andes, Department of Civil and Environmental Engineering, Carrera 1 Este No. 19A-40, Edificio Mario Laserna, Piso 6, 111711, Bogota, Colombia
72. Kazan Federal University, Institute of Environmental Sciences, Kremlevskaya 18, 420008, Kazan, Russia
73. Nature Research Centre, Institute of Botany, Zaliuju Ezeru 49, 08406, Vilnius, Lithuania
74. NA, 7 Derwent Road, LA1 3ES, Lancaster, United Kingdom
75. Babeş-Bolyai University, Hungarian Department of Biology and Ecology, Faculty of Biology and Geology, Republicii street 42., 400015, Cluj-Napoca, Romania
76. University of Latvia, Department of Geography, 1 Jelgavas Street, 1004, Riga, Latvia
77. University of Bayreuth, Climatology, Bayreuth Center of Ecology and Environmental Research (BayCEER), Universitätsstr. 30, 95447, Bayreuth, Germany
78. Stadt Frankfurt am Main - Der Magistrat, Palmengarten, Siesmayerstraße 61, 60323, Frankfurt am Main, Germany
79. Research Centre of Slovenian Academy of Sciences and Arts (ZRC SAZU), Institute of Biology, Novi trg 2, 1000, Ljubljana, Slovenia
80. University of Zagreb, Faculty of Forestry, Svetošimunska 25, 10000, Zagreb, Croatia
81. University of Adelaide, TERN, North Terrace, 5005, Adelaide, Australia
82. University of Zagreb, Faculty of Geotechnical Engineering, Hallerova aleja 7, 42000, Varaždin, Croatia
83. Aristotle University of Thessaloniki, School of Biology, 54124, Thessaloniki, Greece
84. Plant Science and Biodiversity Centre Slovak Academy of Sciences, Institute of Botany, Dubravská cesta 9, 84523, Bratislava, Slovakia
85. University of Erlangen-Nuremberg, Department of Geography, Wetterkreuz 15, 91058, Erlangen, Germany
86. Universidade Federal do Rio Grande do Sul, Department of Ecology, Av Bento Gonçalves 9500, 91501-970, Porto Alegre, Brazil
87. University of Perugia, Department of Chemistry, Biology and Biotechnology, Borgo XX giugno 74, 06124, Perugia, Italy
88. Universidade Regional de Blumenau, Departamento de Engenharia Florestal, Rua São Paulo, 3250, 89030-000, Blumenau, Brazil
89. University of Oulu, Ecology and Genetics Research Unit, Biodiversity Unit, Kaitoväylä 5, 90014, Oulu, Finland
90. Helmholtz Center for Environmental Research - UFZ, Department of Physiological Diversity, Permoserstr. 15, 04318, Leipzig, Germany
91. Leuphana University of Lüneburg, Institute of Ecology, Universitätsallee 1, 21335, Lüneburg, Germany
92. University of Alberta, Department of Biological Sciences, Biological Sciences Building, T6G2E9, Edmonton, Canada
93. University of Alaska, Institute of Arctic Biology, P. O. Box 7570000, 99775, Fairbanks, United States
94. Addis Ababa University, Department of Geography & Environmental Studies, Sidist Kilo SQ, 150178, Addis Ababa, Ethiopia
95. Senckenberg Museum of Natural History Görlitz, Botany Department, PO Box 300 154, 02806, Görlitz, Germany
96. Technische Universität Dresden, International Institute Zittau, Markt 23, 02763, Zittau, Germany
97. Brown University, Department of Ecology and Evolutionary Biology/Brown University Herbarium, 34 Olive Street, 02912, Providence, United States
98. Vienna Institute for Nature Conservation & Analyses, Giessergasse 6/7, 1090, Vienna, Austria
99. Swiss Federal Institute for Forest, Snow and Landscape Research WSL, Research Unit Forest Dynamics, Zürcherstrasse 111, 8903, Birmensdorf, Switzerland
00. Ufa Scientific Centre, Russian Academy of Sciences, Laboratory of Wild-Growing Flora, Botanical Garden-Institute, Mendeleev str., 195/3, 450080, Ufa, Russia

# Abstract

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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

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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 ([???]). Large independent collections of plant occurrence data do exist at the global or continental extent via the Botanical Information and Ecology Network (BIEN) (12), the Global Inventory of Floras and Traits (GIFT) (13) 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 (14).

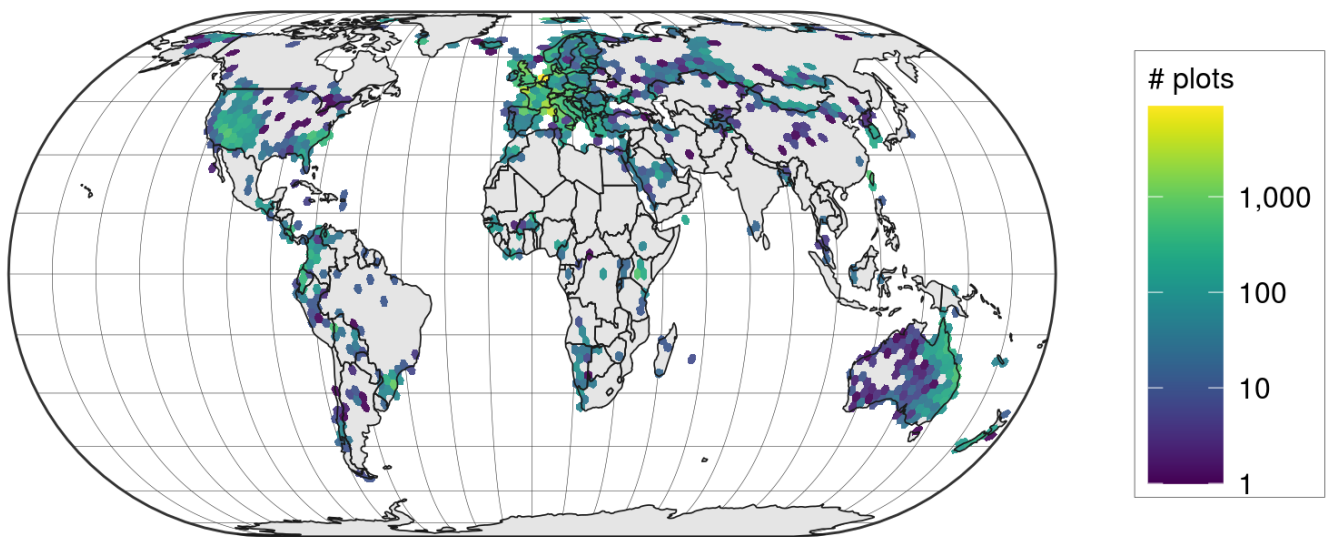
Yet, there is a long-lasting tradition among botanists to record the cover or abundance of each plant species that occurs in a vegetation plot of a given size (i.e. surface area) at a given time (e.g. Stebler & Schröter 1893). Compared to species-level data, vegetation-plot data present many advantages. First, they contain information on which plant species co-occur together in the same locality at a given moment in time (15). 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 (16). Being spatially explicit, vegetation plots can be resurveyed through time to assess potential changes in plant species composition relative to a baseline (17; 18, 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 (19).

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 (20). Furthermore, with their disparate sampling protocols, standards and taxonomic resolutions, aggregating and harmonizing vegetation plot data proves extremely challenging (21). It is not surprising, therefore, that these data have only been rarely used in global-scale biodiversity research until recently (22; 23).

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 (24). Established in 2013, sPlot currently contains more than 1.9 million vegetation plots, and is fully integrated with the TRY database (25), 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 ([26](#)), the mechanisms underlying the spread and abundance of native vs. invasive tree species ([27](#)), and worldwide trait–environment relationships in plant communities ([21](#)).

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 ([25](#)).



**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  $70,000 \text{ km}^2$ . Map projection is Eckert IV.

# Methods

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## 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 [24](#) for more information). For instance: 48 vegetation-plot datasets derive from the European Vegetation Archive (EVA) ([15](#)), three major African datasets from the Tropical African Vegetation Archive (TAVA), multiple vegetation datasets in the USA from the VegBank archive ([28](#); [29](#)). 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; [30](#)), using the GIVD identifier as the unique dataset identifier.

## Resampling method

Data in the sPlot database are unevenly distributed across continents and biomes (see [21](#)). 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., [31](#); [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 ([32](#)), 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 - [33](#)). For soil, we extracted seven variables from the SOILGRIDS database ([34](#)), 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 [[35](#)]. 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 ([36](#)) 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 ([25](#)). These traits were selected among those traits that describe the leaf, wood and seed economics spectra ([37](#); [38](#)), and are known to either affect different key ecosystem processes or respond to macroclimatic drivers or both ([24](#)). The eighteen plant functional traits were: (1) leaf area [ $\text{mm}^2$ ]; (2) stem specific density [ $\text{g cm}^{-3}$ ]; (3) specific leaf area [ $\text{m}^2\text{kg}^{-1}$ ]; (4) leaf carbon concentration [ $\text{mg g}^{-1}$ ]; (5) leaf nitrogen concentration [ $\text{mg g}^{-1}$ ]; (6) leaf phosphorus concentration [ $\text{mg g}^{-1}$ ]; (7) plant height [m]; (8) seed mass [mg]; (9) seed length [mm]; (10) leaf dry matter content [ $\text{g g}^{-1}$ ]; (11) leaf nitrogen per area [ $\text{g m}^{-2}$ ]; (12) leaf N:P ratio [ $\text{g g}^{-1}$ ]; (13) leaf  $\delta^{15}\text{N}$  [per million]; (14) seed number per reproductive unit; (15) leaf fresh mass [g]; (16) stem conduit density [ $\text{mm}^{-2}$ ]; (17) dispersal unit length [mm]; and (18) conduit element length [ $\mu\text{m}$ ].

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', [39](#); [40](#)). 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 [\[24\]](#).



Community-weighted means (CWM) and the variances (CWV) were calculated for every plant functional trait  $j$  and every vegetation plot  $k$  as follows (41):

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

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

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 (11). Information on the size (surface area) of the vegetation survey is available for 61,898 vegetation plots, and ranges between 0.01 m<sup>2</sup> and 4 ha (mean = 270 m<sup>2</sup>; median = 78.5 m<sup>2</sup>). 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 [24]. 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. We recommend for applications that focus on particular vegetation types to retrieve them from the current sPlot version, following sPlot's Governance and Data Property Rules ([www.idiv.de/sPlot](http://www.idiv.de/sPlot)).

# Database Organization

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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 [12](#).

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

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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 (Hennekens & Schaminée 2001). 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 (25) to their resolved version based on the Taxonomic Name Resolution Service web application (TNRS version 4.0; 42; 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 (25). All taxa originally denoted at taxonomic ranks lower than species, were aggregated at species level. Additional detail on the taxonomic resolution is reported in [24], while a description of the workflow, including R-code, is available in Purschke (2017).

## Usage Notes

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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](http://www.idiv.de/sPlot)).

## Code Availability

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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>).

## Acknowledgements

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We are grateful to thousands of vegetation scientists who sampled vegetation plots in the field or digitized them into regional, national or international databases. We also appreciate the support of the German Research Foundation for funding sPlot as one of the iDiv (DFG FZT 118, 202548816) research platforms, and the organization of three workshops through the sDiv calls. We acknowledge this support with naming the database "sPlot", where the "s" refers to the sDiv synthesis workshops. The study was supported by the TRY initiative on plant traits (<http://www.try-db.org>). For all further acknowledgements see Appendix S10 in [24].

## Author contributions

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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

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*[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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**34. SoilGrids250m: Global gridded soil information based on machine learning**

Tomislav Hengl, Jorge Mendes de Jesus, Gerard B. M. Heuvelink, Maria Ruiperez Gonzalez, Milan Kilibarda, Aleksandar Blagotić, Wei Shangguan, Marvin N. Wright, Xiaoyuan Geng, Bernhard Bauer-Marschallinger, ... Bas Kempen

*PLOS ONE* (2017-02-16) <https://doi.org/f9qc5p>

DOI: [10.1371/journal.pone.0169748](https://doi.org/10.1371/journal.pone.0169748) · PMID: [28207752](https://pubmed.ncbi.nlm.nih.gov/28207752/) · PMCID: [PMC5313206](https://pubmed.ncbi.nlm.nih.gov/PMC5313206/)

**35. Heterogeneity-constrained random resampling of phytosociological databases**

Attila Lengyel, Milan Chytrý, Lubomír Tichý

*Journal of Vegetation Science* (2011-02) <https://doi.org/dvjzbz>  
DOI: [10.1111/j.1654-1103.2010.01225.x](https://doi.org/10.1111/j.1654-1103.2010.01225.x)

**36. The relationship between species replacement, dissimilarity derived from nestedness, and nestedness**

Andrés Baselga

*Global Ecology and Biogeography* (2012-12) <https://doi.org/gddc72>

DOI: [10.1111/j.1466-8238.2011.00756.x](https://doi.org/10.1111/j.1466-8238.2011.00756.x)

**37. *unav***

Mark Westoby

*Plant and Soil* (1998) <https://doi.org/bsvqvz>

DOI: [10.1023/a:1004327224729](https://doi.org/10.1023/a:1004327224729)

**38. The world-wide “fast-slow” plant economics spectrum: a traits manifesto**

Peter B. Reich

*Journal of Ecology* (2014-03) <https://doi.org/gfc4z9>

DOI: [10.1111/1365-2745.12211](https://doi.org/10.1111/1365-2745.12211)

**39. Uncertainty Quantified Matrix Completion Using Bayesian Hierarchical Matrix Factorization**

Farideh Fazayeli, Arindam Banerjee, Jens Kattge, Franziska Schrod, Peter B. Reich

*Institute of Electrical and Electronics Engineers (IEEE)* (2014-12) <https://doi.org/ghfnw3>

DOI: [10.1109/icmla.2014.56](https://doi.org/10.1109/icmla.2014.56)

**40. BHPMF - a hierarchical Bayesian approach to gap-filling and trait prediction for macroecology and functional biogeography**

Franziska Schrod, Jens Kattge, Hanhuai Shan, Farideh Fazayeli, Julia Joswig, Arindam Banerjee, Markus Reichstein, Gerhard Bönisch, Sandra Díaz, John Dickie, ... Peter B. Reich

*Global Ecology and Biogeography* (2015-12) <https://doi.org/f76qw8>

DOI: [10.1111/geb.12335](https://doi.org/10.1111/geb.12335)

**41. Scaling from Traits to Ecosystems**

Brian J. Enquist, Jon Norberg, Stephen P. Bonser, Cyrille Violle, Colleen T. Webb, Amanda Henderson, Lindsey L. Sloat, Van M. Savage

*Advances in Ecological Research* (2015) <https://doi.org/ghfnsw>

DOI: [10.1016/bs.aecr.2015.02.001](https://doi.org/10.1016/bs.aecr.2015.02.001)

**42. The taxonomic name resolution service: an online tool for automated standardization of plant names**

Brad Boyle, Nicole Hopkins, Zhenyuan Lu, Juan Antonio Raygoza Garay, Dmitry Mozzherin, Tony Rees, Naim Matasci, Martha L Narro, William H Piel, Sheldon J Mckay, ... Brian J Enquist

*BMC Bioinformatics* (2013-01-16) <https://doi.org/gb8vxz>

DOI: [10.1186/1471-2105-14-16](https://doi.org/10.1186/1471-2105-14-16) · PMID: [23324024](https://pubmed.ncbi.nlm.nih.gov/23324024/) · PMCID: [PMC3554605](https://pubmed.ncbi.nlm.nih.gov/PMC3554605/)

# Supplementary Material

**Table 11:** 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 GIVD	Custodian	Deputy custodian	n - Plots	contributed_plots	Citation
00000004	Vegetation Database of Eurasian Tundra	Risto Virtanen	NA	1132	600	NA
00000003	Database Meadows and Steppes of Southern Ural	Sergey Yamalov	Mariya Lebedeva	2354	99	NA
00000001	Forest Vegetation Database of Turkey - FVDT	Ali Kavgacı	NA	919	15	NA
EU00000002	Nordic-Baltic Grassland Vegetation Database (NBGVD)	Jürgen Dengler	Łukasz Kozub	7675	931	Dengler & Rūsiņa (2012)
EU00000011	Vegetation-Plot Database of the University of the Basque Country (BIOVEG)	Idoia Biurrun	Itziar García-Mijangos	18441	1694	Biurrun et al. (2012)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - 0 0 - 0 1 3	Balkan Dry Grasslands Database	Kiril Vassilev	Armin Macano vić	7 6 8 3	224	Vassilev et al. (2012)
E U - 0 0 - 0 1 6	Mediterranean Ammophiletea Database	Corrado Marcenò	Borja Jiménez -Alfaro	7 3 5 9	3713	Marcenò & Jiménez- Alfaro (2017)
E U - 0 0 - 0 1 7	European Coastal Vegetation Database	John Janssen	NA	4 6 2 4	1369	NA
E U - 0 0 - 0 1 8	The Nordic Vegetation Database	Jonathan Lenoir	Jens- Christia n Svenning	5 4 7 7	1755	Lenoir et al. (2013)
E U - 0 0 - 0 1 9	Balkan Vegetation Database	Kiril Vassilev	Hristo Pedash enko	9 1 1 8	211	Vassilev et al. (2016)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - 0 0 - 0 2 0	WetVegEurope	Flavia Landucci	NA	1 4 1 1 1	61	Landucci et al. (2015)
E U - 0 0 - 0 2 2	European Mire Vegetation Database	Tomáš Peterka	Martin Jiroušek	1 0 1 4 7	1843	Peterka et al. (2015)
E U - A L - 0 0 1	Vegetation Database of Albania	Michele De Sanctis	Giulian o Fanelli	2 9 0	99	De Sanctis et al. (2017)
E U - A T - 0 0 1	Austrian Vegetation Database	Wolfgang Willner	Christia n Berg	3 4 4 5 8	950	Willner et al. (2012)
E U - B E - 0 0 2	INBOVEG	Els De Bie	NA	2 5 6 6 5	48	NA



<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - B G - 0 0 1	Bulgarian Vegetation Database	Iva Apostolova	Desisla va Sopotli eva	5 2 5 4	74	Apostolova et al. (2012)
E U - C H - 0 0 5	Swiss Forest Vegetation Database	Thomas Wohlgemuth	NA	1 4 1 9 3	1409	Wohlgemut h (2012)
E U - C Z - 0 0 1	Czech National Phytosociological Database	Milan Chytrý	Ilona Knollov á	1 0 4 6 9 7	579	Chytrý & Rafajová (2003)
E U - D E - 0 0 1	VegMV	Florian Jansen	Christia n Berg	5 3 8 2 2	5	Jansen et al. (2012)
E U - D E - 0 1 3	VegetWeb Germany	Florian Jansen	Jörg Ewald	2 3 0 7 8	199	Ewald et al. (2012)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - D E - 0 1 4	German Vegetation Reference Database (GVRD)	Ute Jandt	Helge Bruelhe ide	3 0 8 4 0	286	NA
E U - D K - 0 0 2	National Vegetation Database of Denmark	Jesper Erenskjold Moeslund	Rasmus Ejrnæs	2 4 2 6 4	1181	NA
E U - E S - 0 0 1	Iberian and Macaronesian Vegetation Information System (SIVIM) - Wetlands	Aaron Pérez- Haase	Xavier Font	6 5 6 0	292	NA
E U - F R - 0 0 3	SOPHY	Emmanuel Garbolino	Patrice De Ruffray	2 0 9 8 6 4	1332 2	NA
E U - G B - 0 0 1	UK National Vegetation Classification Database	John S. Rodwell	NA	2 8 5 3 3	5457	NA

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - G R - 0 0 1	KRITI	Erwin Bergmeier	NA	2 9 2	43	NA
E U - G R - 0 0 5	Hellenic Natura 2000 Vegetation Database (HelNatVeg)	Panayotis Dimopoulos	Ioannis Tsiripidi s	5 1 6 8	777	Dimopoulos & Tsiripidis (2012)
E U - G R - 0 0 6	Hellenic Woodland Database	Ioannis Tsiripidis	Georgio s Fotiadis	3 1 9 9	4	Fotiadis et al. (2012)
E U - H R - 0 0 1	Phytosociological Database of Non-Forest Vegetation in Croatia	Zvezdana Stančić	NA	5 0 5 7	213	Stančić (2012)
E U - H R - 0 0 2	Croatian Vegetation Database	Željko Škvorc	Daniel Krstono šić	8 7 3 4	688	NA

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - H U - 0 0 3	CoenoDat Hungarian Phytosociological Database	János Csiky	Zoltán Botta- Dukát	8 5 0 5	17	Lájer et al. (2008)
E U - I T - 0 0 1	VegItaly	Roberto Venanzoni	Flavia Landuc ci	1 5 3 3 2	2712	Landucci et al. (2012)
E U - I T - 0 1 0	Italian National Vegetation Database (BVN/ISPRA)	Laura Casella	Pierang ela Angelini	3 5 6 2	155	Casella et al. (2012)
E U - I T - 0 1 1	Vegetation-Plot Database Sapienza University of Rome (VPD-Sapienza)	Emiliano Agrillo	Fabio Attorre	1 2 7 8 0	1003	Agrillo et al. (2017)
E U - L T - 0 0 1	Lithuanian Vegetation Database	Valerijus Rašomavičius	Domas Uoginta s	7 8 2 1	119	NA

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - L V - 0 0 1	Semi-natural Grassland Vegetation Database of Latvia	Solvita Rūsiņa	NA	5 5 9 4	306	Rūsiņa (2012)
E U - M K - 0 0 1	Vegetation Database of the Republic of Macedonia	Renata Ćušterevska	NA	1 4 1 7	10	NA
E U - N L - 0 0 1	Dutch National Vegetation Database	Joop H.J. Schaminée	Stepha n M. Hennek ens	1 0 2 3 2 7	1022 3	Schaminée et al. (2006)
E U - P L - 0 0 1	Polish Vegetation Database	Zygmunt Kącki	Grzegor z Swacha	2 2 2 2 9	464	Kącki & Śliwiński (2012)
E U - R O - 0 0 7	Romanian Forest Database	Adrian Indreica	Pavel Dan Turture anu	6 0 1 7	60	Indreica et al. (2017)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - R O - 0 0 8	Romanian Grassland Database	Eszter Ruprecht	Kiril Vassilev	1 9 2 1	44	Vassilev et al. (2018)
E U - R S - 0 0 2	Vegetation Database Grassland Vegetation of Serbia	Svetlana Aćić	Zora Dajić Stevano vić	5 5 8 7	57	Aćić et al. (2012)
E U - R U - 0 0 2	Lower Volga Valley Phytosociological Database	Valentin Golub	Viktoria Bondar eva	1 4 8 5 3	149	Golub et al. (2012)
E U - R U - 0 0 3	Vegetation Database of the Volga and the Ural Rivers Basins	Tatiana Lysenko	NA	1 5 1 6	96	Lysenko et al. (2012)
E U - R U - 0 1 1	Vegetation Database of Tatarstan	Vadim Prokhorov	Maria Kozhev nikova	7 4 7 1	94	Prokhorov et al. (2017)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
E U - S I - 0 0 1	Vegetation Database of Slovenia	Urban Šilc	Filip Küzmič	1 0 9 8 6	435	Šilc (2012)
E U - S K - 0 0 1	Slovak Vegetation Database	Milan Valachovič	Jozef Šibík	3 6 4 0 5	893	Šibík (2012)
E U - U A - 0 0 6	Vegetation Database of Ukraine and Adjacent Parts of Russia	Viktor Onyshchenko	Vitaliy Kolomiy chuk	3 3 2 6	479	NA
A F - 0 0 - 0 0 1	West African Vegetation Database	Marco Schmidt	Georg Zizka	3 1 2 9	184	Schmidt et al. (2012)
A F - 0 0 - 0 0 8	PANAF Vegetation Database	Hjalmar Kühl	TeneKw etche Sop	2 4 6 9	942	NA



<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A F - B F - 0 0 1	Sahel Vegetation Database	Jonas V. Müller	Marco Schmid t	1 0 7 9	279	Müller (2003)
0 0 - 0 0 - 0 0 1	ForestPlots.net	Oliver L. Phillips	Aurora Levesle y	1 8 2 7	108	Lopez- Gonzalez et al. (2011)
0 0 - 0 0 - 0 0 3	SALVIAS	Brian Enquist	Brad Boyle	4 8 8 3	2860	NA
0 0 - 0 0 - 0 0 5	Tundra Vegetation Plots (TundraPlot)	Anne D. Bjorkman	Sarah Elmend orf	5 7 7	227	Elmendorf et al. (2012)
0 0 - R U - 0 0 2	Database of Masaryk University`s Vegetation Research in Siberia	Milan Chytrý	NA	1 5 4 7	128	Chytrý (2012)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A F - 0 0 - 0 0 3	BIOTA Southern Africa Biodiversity Observatories Vegetation Database	Norbert Jürgens	Ute Schmiedel	1 6 6 6 6	562	Muche et al. (2012)
A F - 0 0 - 0 0 6	SWEA-Dataveg	Miguel Alvarez	Michael Curran	2 7 0 4	1211	NA
A F - 0 0 - 0 0 9	Vegetation Database of the Okavango Basin	Rasmus Revermann	Manfred Finckh	5 9 0	202	Revermann et al. (2016)
A F - C D - 0 0 1	Forest Database of Central Congo Basin	Kim Sarah Jacobsen	Hans Verbeek	2 9 2	97	Kearsley et al. (2013)
A F - E T - 0 0 1	Vegetation Database of Ethiopia	Desalegn Wana	Anke Jentsch	7 4	59	Wana & Beierkuhnlein (2011)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A F - M A - 0 0 1	Vegetation Database of Southern Morocco	Manfred Finckh	NA	1 3 3 7	266	Finckh (2012)
A F - Z W - 0 0 1	Vegetation Database of Zimbabwe	Cyrus Samimi	NA	3 6	17	Samimi (2003)
A S - 0 0 - 0 0 1	Korean Forest Database	Tomáš Černý	Jiri Dolezal	4 8 8 5	766	Černý et al. (2015)
A S - 0 0 - 0 0 3	Vegetation of Middle Asia	Arkadiusz Nowak	Marcin Nobis	1 3 8 1	128	Nowak et al. (2017)
A S - 0 0 - 0 0 4	Rice Field Vegetation Database	Arkadiusz Nowak	NA	1 7 9	31	NA

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A S - B D - 0 0 1	Tropical Forest Dataset of Bangladesh	Mohammed A.S. Arfin Khan	Fahmid a Sultana	2 1 1	82	NA
A S - C N - 0 0 1	China Forest-Steppe Ecotone Database	Hongyan Liu	Fengjun Zhao	1 4 8	97	Liu et al. (2000)
A S - C N - 0 0 2	Tibet-PaDeMoS Grazing Transect	Karsten Wesche	NA	1 4 6	27	Wang et al. (2017)
A S - C N - 0 0 3	Vegetation Database of the BEF China Project	Helge Bruelheide	NA	2 7	18	Bruelheide et al. (2011)
A S - C N - 0 0 4	Vegetation Database of the Northern Mountains in China	Zhiyao Tang	NA	4 8 5	70	NA

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A S - E G - 0 0 1	Vegetation Database of Sinai in Egypt	Mohamed Z. Hatim	NA	9 2 6	98	Hatim (2012)
A S - I D - 0 0 1	Sulawesi Vegetation Database	Michael Kessler	NA	2 4	24	NA
A S - I R - 0 0 1	Vegetation Database of Iran	Jalil Noroozi	Parasto o Mahdav i	2 3 3 5	105	NA
A S - K Z - 0 0 1	Database of Meadow Vegetation in the NW Tien Shan Mountains	Viktoria Wagner	NA	9 4	3	NA
A S - M N - 0 0 1	Southern Gobi Protected Areas Database	Henrik von Wehrden	Karsten Wesche	1 5 1 6	688	von Wehrden et al. (2009)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A S - R U - 0 0 1	Wetland Vegetation Database of Baikal Siberia (WETBS)	Victor Chepinoga	NA	2 3 8 1	6	Chepinoga (2012)
A S - R U - 0 0 2	Database of Siberian Vegetation (DSV)	Andrey Korolyuk	Andrei Zverev	9 1 1 6	2150	NA
A S - R U - 0 0 4	Database of the University of Münster - Biodiversity and Ecosystem Research Group's Vegetation Research in Western Siberia and Kazakhstan	Norbert Hölzel	Wanja Mathar	4 4 5	85	NA
A S - S A - 0 0 1	Vegetation Database of Saudi Arabia	Mohamed Abd El-Rouf Mousa El- Sheikh	NA	9 1 9	607	NA
A S - T J- 0 0 1	Eastern Pamirs	Kim André Vanselow	NA	2 8 2	174	Vanselow (2016)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A S - T W - 0 0 1	National Vegetation Database of Taiwan	Ching-Feng Li	Chang- Fu Hsieh	9 3 0	897	NA
A S - Y E - 0 0 1	Socotra Vegetation Database	Michele De Sanctis	Fabio Attorre	3 9 6	190	De Sanctis & Attorre (2012)
A U - A U - 0 0 2	AEKOS	Anita Smyth	Ben Sparro w	2 1 2 6 1	7443	NA
A U - N C - 0 0 1	New Caledonian Plant Inventory and Permanent Plot Network (NC-PIPPN)	Jérôme Munzinger	Philippe Birnbau m	2 0 1	98	Ibanez et al. (2014)
A U - N Z - 0 0 1	New Zealand National Vegetation Databank	Susan Wiser	NA	1 8 9 5	983	Wiser et al. (2001)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
A U - P G - 0 0 1	Forest Plots from Papua New Guinea	Timothy Whitfeld	George D. Weible n	6 3	53	Whitfeld et al. (2014)
N A - 0 0 - 0 0 2	Tree Biodiversity Network (BIOTREE-NET)	Luis Cayuela	NA	1 7 5 7	208	Cayuela et al. (2012)
N A - C A - 0 0 3	Database of Timberline Vegetation in NW North America	Viktoria Wagner	Toby Spribille	1 1 0	38	NA
N A - C A - 0 0 4	Understory of Sugar Maple Dominated Stands in Quebec and Ontario (Canada)	Isabelle Aubin	NA	1 5 6	9	Aubin et al. (2007)
N A - C A - 0 0 5	Boreal Forest of Canada	Yves Bergeron	Louis De Grandp ré	8 9	44	NA



<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - p l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
N A - G L - 0 0 1	Vegetation Database of Greenland	Birgit Jedrzejek	Fred J.A. Daniëls	6 6 4	340	Sieg et al. (2006)
N A - U S - 0 0 2	VegBank	Robert K. Peet	Michael T. Lee	6 7 3 5 2	6456	Peet et al. (2012a)
N A - U S - 0 0 6	Carolina Vegetation Survey Database	Robert K. Peet	Michael T. Lee	1 7 2 2 1	2317	Peet et al. (2012b)
N A - U S - 0 1 4	Alaska-Arctic Vegetation Archive	Donald A. Walker	Amy Breen	1 3 6 3	467	Walker et al. (2016)
S A - 0 0 - 0 0 2	VegPáramo	Gwendolyn Peyre	Xavier Font	2 6 4 3	1591	Peyre et al. (2015)

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
S A - A R - 0 0 2	Vegetation Database of Central Argentina	Melisa Giorgis	Alicia Acosta	2 1 8	42	NA
S A - B O - 0 0 3	Bolivia Forest Plots	Michael Kessler	Sebasti an Herzog	7 5	18	NA
S A - B R - 0 0 2	Forest Inventory, State of Santa Catarina, Brazil (IFFSC Project)	Alexander Christian Vibrans	André Luis de Gaspar	1 6 6 9	1345	Vibrans et al. (2010)
S A - B R - 0 0 3	Grasslands of Rio Grande do Sul, Brazil	Eduardo Vélez-Martin	Valério De Patta Pillar	3 2 0	271	NA
S A - B R - 0 0 4	Grassland Database of Campos Sulinos	Gerhard E. Overbeck	Valério De Patta Pillar	1 6 1	111	NA

<b>G I V D I D</b>	<b>DB_name GIVD</b>	<b>Custodian</b>	<b>Deputy custodi an</b>	<b>n - P l o t s</b>	<b>cont ribut ed_p lots</b>	<b>Citation</b>
S A - C L - 0 0 2	SSAForests_Plots_db	Alvaro G. Gutierrez	NA	2 6 1	163	NA
S A - C L - 0 0 3	Chilean Park Transects - Fondecyt 1040528	Aníbal Pauchard	Alicia Martico rena	1 6 5	33	NA
S A - E C - 0 0 1	Ecuador Forest Plot Database	Jürgen Homeier	NA	1 7 2	156	NA

**Table 12:** 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).

<b>Var iabl e</b>	<b>Range/Levels</b>	<b>N o · r e c o r d s</b>	<b>Ty pe · of · v ari abl e</b>
GIV D_I D	NA	9 1 0 3 1	ch ar ac ter
Dat ase t	NA	9 1 0 3 1	ch ar ac ter

Variable	Range/Levels	No. records	Type of variable
Continent	Africa, Asia, Australia, Europe, North America, Oceania, South America	90729	factor
Country	NA	91031	character
Bio me	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	91031	numeric
Longitude	-162.741433 - 179.590053	91031	numeric
Location_uncertainty	1 - 2500	91002	integer
Relevance	0.01 - 40000	61898	numeric
Herbs_identified	FALSE = 4876; TRUE = 6323	11199	logical

Variable	Range/Levels	No. records	Type of variable
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
Altitude	-25 - 4819	52121	numeric
Aspect	0 - 360	30796	numeric
Slope	-5 - 99	37878	numeric
is_forest	FALSE = 20396; TRUE = 25832	46228	logical
is_nonforest	FALSE = 50870; TRUE = 38203	89073	logical
ESY	NA	55457	character
Naturalness	1 - 2	68011	integer
Forest	FALSE = 38295; TRUE = 23735	62030	logical

Variable	Range/Levels	No. records	Type of variable
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
Cover_tree_layer	0.5 - 150	7245	numeric
Cover_shrub_layer	0.5 - 145	10197	numeric
Cover_herb_layer	0.2 - 180	26679	numeric

Var iabl e	Range/Levels	N o . r e c o r d s	Ty pe of .v ari abl e
Cov er_ mo ss_l aye r	1 - 100	9 6 4 3	int eg er
Cov er_l ich en_ lay er	1 - 95	7 3 4	int eg er
Cov er_ alg ae_l aye r	1 - 100	2 2 1	int eg er
Cov er_l itte r_la yer	1 - 100	4 5 0 0	int eg er
Cov er_ bar e_r ock s	1 - 100	1 8 9 7	int eg er
Cov er_ cry pto ga ms	1 - 95	5 9 3	int eg er
Cov er_ bar e_s oil	0.1 - 99	1 4 1 2	nu m eri c
Hei ght_ tre es_ hig hes t	1 - 99	6 1 1 5	nu m eri c

Variable	Range/Levels	No. records	Type of variable
Height_trees_lowest	1 - 90	221	numeric
Height_shrubshighest	0.1 - 9.9	2880	numeric
Height_shrubslowest	0.1 - 9	328	numeric
Height_herbs_average	0.1 - 440	10125	numeric
Height_herbs_lowest	1 - 250	2785	integer
Height_herbs_highest	1 - 600	1733	integer