Detailed data sources and treatments

1) Species occurrences data

Occurrences data were extracted from the Global Biodiversity Information Facility platform (GBIF: https://www.gbif.org/). To achieve precise species prediction from a geolocation, the geolocations in question must be as precise as possible. However, a high number of occurrences from the GBIF have a spatially degraded geolocation for conservation reasons. Thus, we have chosen source datasets with undegraded geolocations in France, which are:

- Carnet en ligne
 (https://www.gbif.org/dataset/baa86fb2-7346-4507-a34f-44e4c1bd0d57) from Tela

 Botanica.
- Cartographie des Leguminosae (Fabaceae) en France (https://www.gbif.org/dataset/cbd241aa-a115-4856-af66-fac5cb90f2cc) from Tela Botanica
- Naturgucker dataset
 (https://www.qbif.org/dataset/6ac3f774-d9fb-4796-b3e9-92bf6c81c084)
- iNaturalist Research-grade Observations (https://www.gbif.org/dataset/50c9509d-22c7-4a22-a47d-8c48425ef4a7).

In order to maximize the size of data while ensuring homogeneous and rich environmental data (see next part), we restricted the domain of study to the French metropolitan territory, as can be seen of Figure 1. Thus, observations falling outside of it were removed. As some occurrences submitted to the GBIF have been generated from species checklists, several observations of a same species can be at the exact same location. To avoid model bias due to the heterogeneity of the datasets on this matter, we removed the redundancies of a single species at a single location. We chose to work at the species level, while GBIF occurrences can be at any taxonomic level, even inferior to the species (sub-species, varieties), with the accepted name of the bdtfx taxonomic referential for western Europe plant species. We eliminated observations for which the taxonomic matching with the GBIF backbone taxonomy was fuzzy or with higher rank than the species (genus, family etc.). For the remaining observations, we added a field espece_retenue_bdtfx which is the name of the species in the referential bdtfx 4.01(from 15/03/2017). The matching was done from the field scientificname with an internal exact matching algorithm. If the rank was inferior to the species, we took the corresponding species accepted name. If there was no exact match, the observation was excluded, but this case was rare. The field **species_glc_id** attributes an id for each possible value of espece_retenue_bdtfx and will be our class identifier for the task.

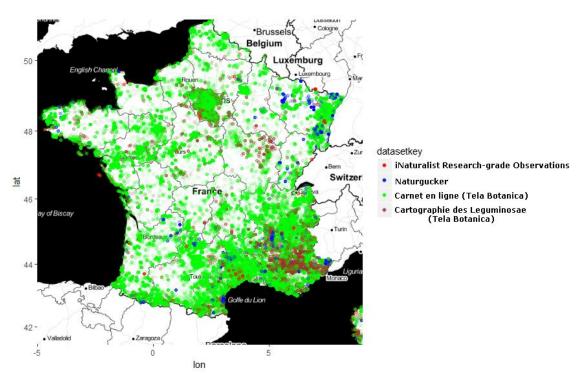


Figure 1: Spatial distribution and source dataset of the occurrences

2) Environmental data

Specimens of a plant species tend to aggregate spatially due to dispersal phenomenons. However, a very important filter for their survival is their adequacy to the abiotic environment. Thus, in addition to know geolocations of species observations, information about the nature of the environment is very useful for modeling species presence or abundance. Those approaches, establishing statistical links between environmental descriptors and species presence, are widely used and usually called Species Distribution Models (SDM). In this task, we supply the participants with a set of Environmental Variables (EV) along with every observation. General informations concerning the nature and range of provided EVs are described in **Table 1**. Each observation **o** is associated a **TIFF** file containing 33 spatial rasters (centered on the location of **o** and of identical dimensions). Each raster is associated with an EV **j**, and must be understood **as a spatial map of the values of j** around the location of **o**. More precisely, it covers the extent:

Where **x_o** is the longitude of **o**, **y_o** is its latitude, **res_x_j** is the initial resolution in longitude of EV **j**, and **res_y_j**, its resolution in latitude. The TIFF file of **o** is located in the directory "**patchTrain**" if **o** is a train observation, or in "**patchTest**" otherwise. Inside each of those directories, there are many subdirectories with names "256", "512" etc. The field **patch_dirname** gives the name of the subdirectory where is the file of **o**. Then, the file is identified with the field **patch_id**. Values stored in each pixel of a TIFF file are between 0 and 255, thus for "awc_top", "bs_top", "cec_top", "crusting", "dgh", "dimp", "erodi", "oc_top", "pd_top", "text", "clc", and "proxi_eau" there is no transformation, so the pixel value corresponds exactly to the original value of the EV. However, for

"chbio_xx","alti" and "etp" one will recover the original value by applying the following operation :

original_value = min + (max-min) [(pixel_value/255) -0.1] /0.8

Here are the max/min values of the EVs and their order in the TIFF:

| variables max min 1 etp 1176.00000 133.000000 2 chbio_1 18.36730 -10.600984 3 chbio_2 20.94560 7.846126 4 chbio_3 59.95573 41.182110 5 chbio_4 777.74048 302.772980 6 chbio_5 36.54550 6.182446 7 chbio_6 5.33183 -28.248663 8 chbio_7 41.94211 16.744829 9 chbio_8 22.96798 -14.122952 10 chbio_9 26.45534 -17.672335 11 chbio_10 26.45534 -2.738379 12 chbio_11 11.73241 -17.672335 13 chbio_12 2543.30225 318.297485 14 chbio_13 285.43790 43.063732 15 chbio_14 135.58406 3.022581 16 chbio_15 57.78888 8.283675 17 chbio_18 851.60620 | | | | |
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Note: Some participants could be only interested in the punctual value of environmental variables at the occurrence geolocation, as it is often done in ecological modeling. To facilitate the use of the data, we stored those values as columns in the occurrences CSV files. All these values are on the variable original scale.

Note: patch_dirname = -1 * euclidian_division_quotient(-1*patch_id / 256) * 256.

| Name | Description | Nature | Values |
|--------------|-------------------------------------|---------|--|
| CHBIO_1 | Annual Mean Temp. | quanti. | [-10.7,18.4] |
| | Mean of monthly | | 10.7014 O 2015 S 10.0012 O 2015 O |
| CHBIO_2 | max(temp)-min(temp) | quanti. | [7.8,21.0] |
| CHBIO_3 | Isothermality (100*2/7) | quanti. | [41.1,60.0] |
| $CHBIO_4$ | Temp. Seasonality (std.dev.*100) | quanti. | [302.7,777.8] |
| CHBIO_5 | Max Temp. of Warmest Month | quanti. | [6.1,36.6] |
| CHBIO_6 | Min Temp. of Coldest Month | quanti. | [-28.3,5.4] |
| CHBIO_7 | Temp. Annual Range (5-6) | quanti. | [16.7,42.0] |
| CHBIO_8 | Mean Temp. of Wettest Quarter | quanti. | [-14.2,23.0] |
| CHBIO_9 | Mean Temp. of Driest Quarter | quanti. | [-17.7,26.5] |
| CHBIO_10 | Mean Temp. of Warmest Quarter | quanti. | [-2.8,26.5] |
| CHBIO_11 | Mean Temp. of Coldest Quarter | quanti. | [-17.7,11.8] |
| $CHBIO_{12}$ | Annual Precip. | quanti. | [318.3,2543.3] |
| CHBIO_13 | Precip. of Wettest Month | quanti. | [43.0,285.5] |
| CHBIO_14 | Precip. of Driest Month | quanti. | [3.0,135.6] |
| CHBIO_15 | Precip. Seasonality (Coef. of Var.) | quanti. | [8.2,26.5] |
| CHBIO_16 | Precip. of Wettest Quarter | quanti. | [121.6,855.6] |
| CHBIO_17 | Precip. of Driest Quarter | quanti. | [19.8,421.3] |
| CHBIO_18 | Precip. of Warmest Quarter | quanti. | [19.8,851.7] |
| CHBIO_19 | Precip. of Coldest Quarter | quanti. | [60.5,520.4] |
| etp | Potential Evapo Transpiration | quanti. | [133,1176] |
| alti | Elevation | quanti. | [-188,4672] |
| awc_top | Topsoil available water capacity | ordinal | {0, 120, 165, 210} |
| bs_top | Base saturation of the topsoil | ordinal | $\{35, 62, 85\}$ |
| cec_top | Topsoil cation exchange capacity | ordinal | $\{7, 22, 50\}$ |
| crusting | Soil crusting class | ordinal | [0,5] |
| dgh | Depth to a gleyed horizon | ordinal | {20, 60, 140} |
| dimp | Depth to an impermeable layer | ordinal | {60, 100} |
| erodi | Soil erodibility class | ordinal | [0, 5] |
| oc_top | Topsoil organic carbon content | ordinal | $\{1, 2, 4, 8\}$ |
| pd_top | Topsoil packing density | ordinal | |
| text | Dominant surface textural class | ordinal | [[0,5]] |
| proxi_eau_fa | ast $<$ 50 meters to fresh water | bool. | $\{0, 1\}$ |
| clc | ground occupation | categ. | [1,48] |

Table 1: The environmental variables supplied for the task.

Note: To open the TIFF files correctly and access to all the 33 channels, we advise participants to use a TIFF dedicated library like **tifffile** for **python** or **TIFF** for **R**.

3) Details on the source and production method of the original data

-- Chelsea Climate data 1.1: those are raster data with worldwide coverage and 1km resolution. A mechanistic climatic model is used to make spatial predictions of monthly

mean-max-min temperatures, mean precipitations and 19 bioclimatic variables, which are downscaled with statistical models

integrating historical measures of meteorologic stations from 1979 to today. The exact method is explained in the reference paper Karger et al. [1]. The data is under Creative Commons Attribution 4.0 International License and downloadable at (http://chelsa-climate.org/downloads/).

-- The ESDB v2 - 1kmx1km Raster Library (Panagos [2],Van Liedekerke et al. [3]): The library contains multiple soil pedology descriptor raster layers covering Eurasia at a resolution of 1km. We selected 11 descriptors from the library. More precisely, those variables have ordinal format, representing physico-chemical properties of the soil, and come from the PTRDB. The PTRDB variables have been directly derived from the initial Soil Geographical Data Base of Europe (SGDBE) using expert rules. SGDBE was a spatial semantic data base relating spatial units to a diverse pedological attributes of categorical nature, which is not useful for our purpose. For more details, see Panagos et al. [4]. The data is maintained and distributed freely for scientific use by the European Soil Data Centre (ESDAC) at http://eusoils.jrc.ec.europa.eu/content/european-soil-database-v2-raster....
-- Corine Land Cover 2012, version 18.5.1, 12/2016: It is a raster layer describing soil occupation with 48 categories across Europe (25 countries) at a resolution of 100 meters. This classification is the result of an automated interpretation process applied to earth

http://land.copernicus.eu/pan-european/corine-land-cover/clc-2012.

accessible online for all use at

-- CGIAR-CSI ETP data: The CGIAR-CSI distributes this worldwide monthly potential-evapotranspiration raster data. It is pulled from a model developed by Antonio Trabucco (see Zomer et al. [5,6]). Those are estimated by the Hargreaves formula, using mean monthly surface temperatures and standard deviation from WorldClim 1:4 (http://www.worldclim.org/version1), and radiation on top of atmosphere. The raster is at a 1km resolution, and is freely downloadable for a nonprofit use at http://www.cgiar-csi.org/data/global-aridity-and-pet-database#description.

surface high resolution satellite images. This data base of the European Union is freely

- -- USGS Digital Elevation data: The Shuttle Radar Topography Mission achieved in 2010 by Endeavour shuttle managed to measure digital elevation at 3 arc second resolution over most of the earth surface. Raw measures have been post-processed by NASA and NGA in order to correct detection anomalies. The data is available from the U.S. Geological Survey, and downloadable on the Earthexplorer (https://earthexplorer.usgs.gov/). See https://lta.cr.usgs.gov/SRTMVF for more informations.
- -- BD Carthage v3: BD Carthage is a spatial semantic database holding many informations on the structure and nature of the french metropolitan hydrological network. For the purpose of plants ecological niche, we focus on the geometric segments representing watercourses, polygons representing hydrographic fresh surfaces and seas. The data has been produced by the Institut National de l'information Géographique et forestière (IGN) from an interpretation of the BD Ortho IGN. It is maintained by the SANDRE under free license for non-profit use and downloadable at

http://services.sandre.eaufrance.fr/telechargement/geo/ETH/BDCarthage/FX....

Note about variables building: For reproducibility, we explain the treatments applied to the original data. As a first treatment, we crop the source layers as short as possible to make calculations faster, making sure the extent contains all the metropolitan French territory. Then, as the original coordinate system of the layer vary among sources, we change it to WGS84, which is the occurrences coordinate system of the GBIF occurrences. Additional processing were necessary to get **proxi_eau**: Its raster have been made from a vector shapefile according to the following procedure. We use qgis to rasterize to a 12.5 meters resolution, with a buffer of 50 meters, the shapefile COURS_D_EAU.shp on one hand, and the polygons of SURFACES_HYDROGRAPHIQUES.shp with attribute NATURE="Eau douce permanente" on the other hand. We then create the maximum raster of the previous ones (So the value of 1 correspond to an approximate distance of less than 50 meters to a watercourse or hydrographic surface of fresh water).

4) Train & test sets

The total of 291,392 occurrences were randomly splitted in a train (218,543) and test (72,849) set with the constraints that :

- For each species in the test set, there is at least one observation of it in the train set.
- An observation of a species in the test set is distant of more than 100 meters from all observations of this species in the train set to avoid major reporting dependances.

Train observations are listed in the file **occurrences_train.csv**, with every field from the GBIF, plus the ones described previously. Test observations are in the file **occurrences_test.csv** whose taxonomic informations were removed.

5) Task description

Data for the task can be downloaded at http://otmedia.lirmm.fr/LifeCLEF/GeoLifeCLEF2018/.

Given the test set of plants observations, the goal of the task is to return for each observation a ranked list of plant species sorted according to the likelihood that they might have been observed at that location. The test set as well as the list of the 3,203 candidate species will be provided within the registration/submission system (see main page of LifeCLEF 2018 for registration).

Each participating group is allowed to submit several run files built from different methods. Each run file has to be named as "teamname_runX.run" where X is the identifier of the run (i.e. 1, 2, 3 or 4). The run file has to contain as much lines as the total number of species recommendations, with at least one recommendation per occurrence of the test set and a maximum of XX recommendations per occurrence (3,203 being the total number of species). Each recommendation (i.e. each line of the run file) has to respect the following format: < glc_id;species_glc_id;probability;rank>

where **glc_id** is the identifier of an occurrence in the test set, **species_glc_id** is the identifier of one of the 3,203 possible species, probability is a real value in [0;1] decreasing with the confidence in that recommendation, rank is the rank of that recommendation among all recommended species for the test occurrence **glc_id**.

Here is a short fake run example respecting this format on only 3 test occurrences:

2777702;2888740;0.955044784350321;1 2777702;2808330;0.0201473159249872;4 2777702;2926110;0.345950950868428;3 2775920;3152583;0.255595623515546;2 2775920;2888740;0.109065826749429;3 2775920;2808330;0.99270333093591;1 2775920;5415039;0.772525980370119;1 2775920;8324121;0.462680039694533;2

For each submitted run, please give within the submission system a short description of the run in the dedicated text area (in addition to the working note to be written later on).

References

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