

Appendix 1 - sPlotOpen - Demo

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Appendix to the paper: Sabatini, Lenoir et al., sPlotOpen – An environmentally-balanced, open-access, global dataset of vegetation plots. *Global Ecology and Biogeography*.

This demo illustrates how to import and manipulate sPlotOpen data to create some basic graphics or tables together with a reference list. As a worked example, the code below will:

1. select all plots containing at least a species of *Quercus* from sPlotOpen's resampled iteration #1
2. show some summary at biome level
3. graph the distribution of the community weighted mean of a selected functional trait
4. show the geographical location of all selected plots
5. create a reference list based on the plots effectively selected.

```
#load libraries
library(tidyverse)
library(sf)
library(raster)
library(rnaturalearth)
library(RefManager)
```

Import data

```
load("_sPlotOpenDB/sPlotOpen.RData")
ls()

## [1] "CWM_CWV.oa"          "DT2.oa"              "header.oa"
## [4] "metadata.oa"         "reference.oa"         "sPlotOpen_citation"
```

Extract all plots containing at least a *Quercus* species

Use only the first resampled iteration of sPlotOpen

```
#select only the first resample
header.oa1 <- header.oa %>%
  filter(Resample_1 == T)
DT2.oa1 <- DT2.oa %>%
  filter(PlotObservationID %in% header.oa1$PlotObservationID)
CWM_CWV.oa1 <- CWM_CWV.oa %>%
  filter(PlotObservationID %in% header.oa1$PlotObservationID)

#get all plots containing at least one Quercus species
plotlist.quercus <- DT2.oa1 %>%
```

```

filter(str_detect(Species, "^Quercus")) %>%
distinct(PlotObservationID) %>%
pull(PlotObservationID)

header.quercus <- header.oa1 %>%
  filter(PlotObservationID %in% plotlist.quercus &
    Resample_1 == T)

DT2.quercus <- DT2.oa1 %>%
  filter(PlotObservationID %in% plotlist.quercus)

CWM_CWV.quercus <- CWM_CWV.oa1 %>%
  mutate(Quercus=ifelse(PlotObservationID %in% plotlist.quercus, T, F))

```

There are 5143 plots containing at least a *Quercus* species in sPlotOpen's resampled iteration 1.

Number of plots with *Quercus* across biomes

Summarize the number of plots containing at least one *Quercus* species across biomes

```

header.quercus %>%
  group_by(Biome) %>%
  summarize(n = n())

```

```

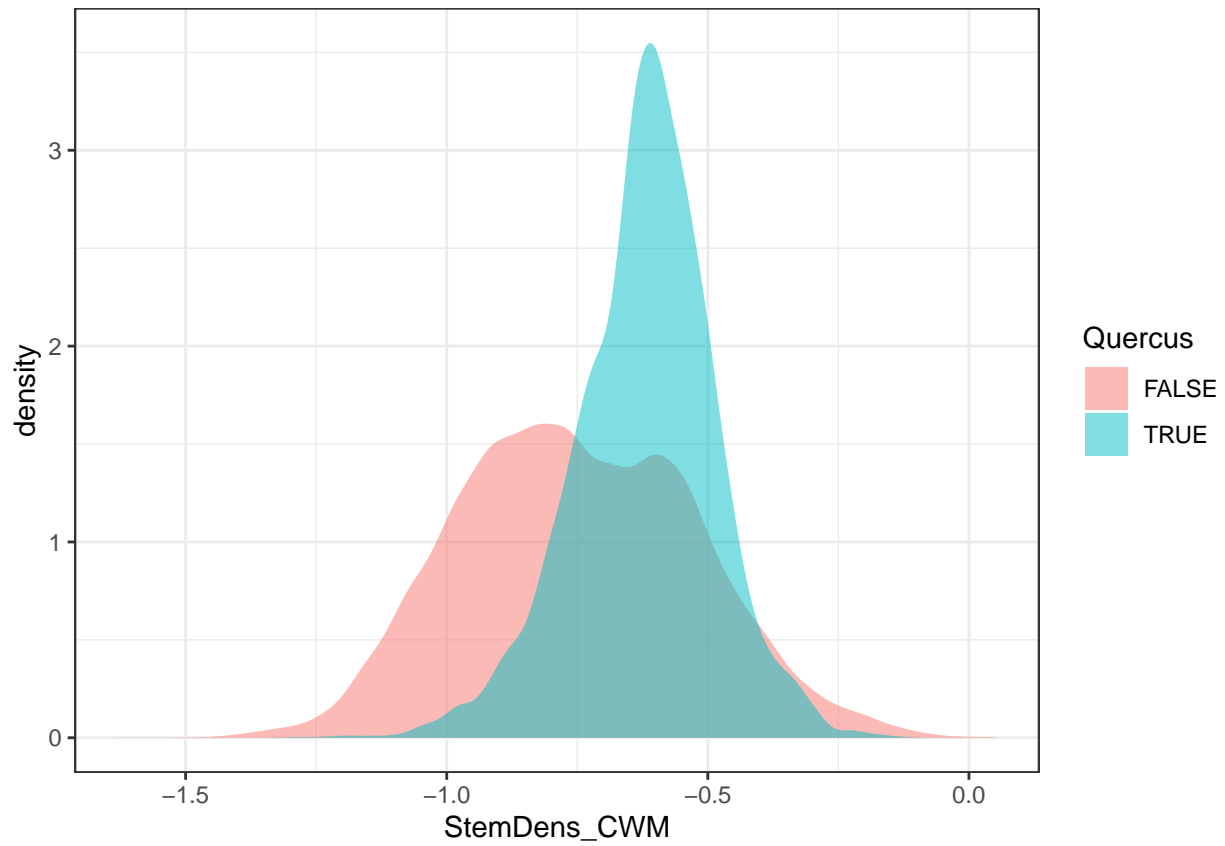
## # A tibble: 9 x 2
##   Biome                                n
## * <fct>                                <int>
## 1 Alpine                                6
## 2 Boreal zone                            7
## 3 Dry midlatitudes                       76
## 4 Dry tropics and subtropics             183
## 5 Subtropics with year-round rain       1157
## 6 Subtropics with winter rain           748
## 7 Temperate midlatitudes                1952
## 8 Tropics with summer rain               580
## 9 Tropics with year-round rain          434

```

Compare Community Weighted Means

Compare the distribution of the community weighted means of Stem density, between plots containing and not containing a *Quercus* species.

```
ggplot(data = CWM_CWV.quercus) +  
  geom_density(aes(x = StemDens_CWM, fill = Quercus), col = NA, alpha = 0.5) +  
  theme_bw()
```



Geographical distribution of plots containing a *Quercus* species

Download some spatial data of the world and create a template map using the r package `rnaturalearth`, first. Transform all geographical data to Eckert IV projection.

```
countries <- ne_countries(returnclass = "sf") %>%
  st_transform(crs = "+proj=eck4") %>%
  st_geometry()
graticules <- ne_download(type = "graticules_15", category = "physical",
  returnclass = "sf") %>%
  st_transform(crs = "+proj=eck4") %>%
  st_geometry()
bb <- ne_download(type = "wgs84_bounding_box", category = "physical",
  returnclass = "sf") %>%
  st_transform(crs = "+proj=eck4") %>%
  st_geometry()
```

Template of Global map - with country borders

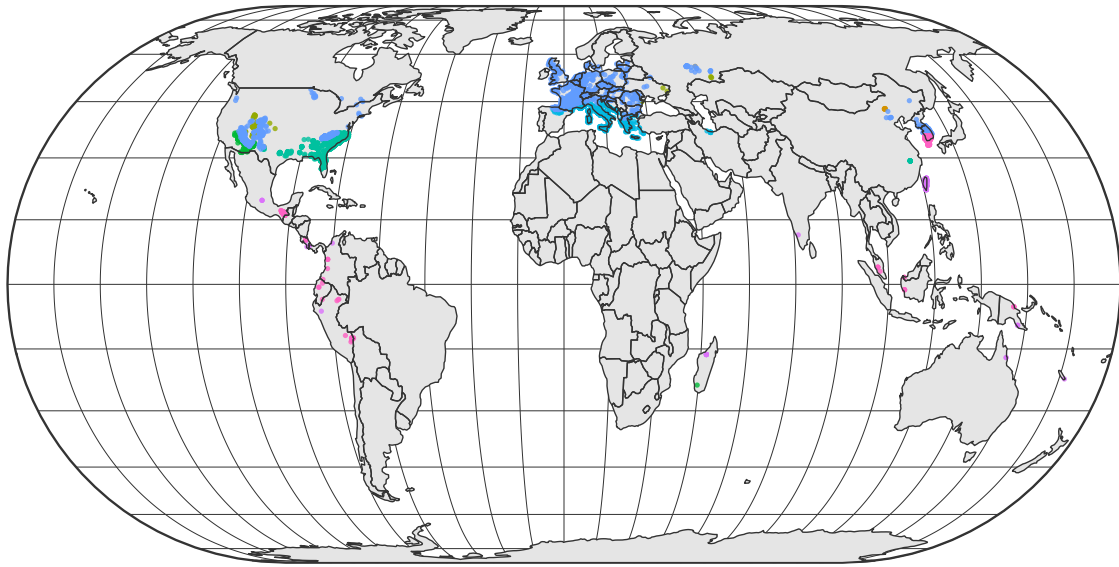
```
w3a <- ggplot() +
  geom_sf(data = bb, col = "grey20", fill = "white") +
  geom_sf(data = graticules, col = "grey20", lwd = 0.1) +
  geom_sf(data = countries, fill = "grey90", col = NA, lwd = 0.3) +
  coord_sf(crs = "+proj=eck4") +
  theme_minimal() +
  theme(axis.text = element_blank(),
    legend.title = element_text(size=12),
    legend.text = element_text(size=12),
    legend.background = element_rect(size = 0.1, linetype = "solid", colour = 1),
    legend.key.height = unit(1.1, "cm"),
    legend.key.width = unit(1.1, "cm"))
```

Project selected plots to Eckert IV and transform them to sf, before plotting.

```
header.quercus.sf <- SpatialPointsDataFrame(coords = header.quercus %>%
  dplyr::select(Longitude, Latitude),
  proj4string = CRS("+init=epsg:4326"),
  data=header.quercus %>%
  dplyr::select(-Longitude, -Latitude)) %>%
  st_as_sf() %>%
  st_transform(crs = "+proj=eck4")
```

Show all plots containing at least one *Quercus* species. Color code based on biomes.

```
(Figure1a <- w3a +
  geom_sf(data = header.quercus.sf, aes(color = Biome),
    pch = 16, size = 0.8, alpha = 0.8) +
  geom_sf(data = countries, col = "grey20", fill=NA, lwd = 0.3) +
  theme(legend.position = "bottom",
    legend.title = element_blank()) +
  guides(color = guide_legend(ncol = 2,
    override.aes = list(size = 2))))
```



● Alpine	● Subtropics with winter rain
● Boreal zone	● Temperate midlatitudes
● Dry midlatitudes	● Tropics with summer rain
● Dry tropics and subtropics	● Tropics with year-round rain
● Subtropics with year-round rain	

Create reference list as BibText

```
## Loading required package: bib2df
```

```
## Warning: package 'bib2df' was built under R version 4.0.3
```

```
## WARNING: This is a beta-version. References were parsed and converted automatically. They might need
```

```
# show first few lines of output file
```

```
read_lines("output/demo.bib", n_max = 25)
```

```
## [1] "@Article{dengler2012a,"
## [2] "  Author = {Dengler, J{\\"u\"rger and R{\\"u\"si{\\"c{n}}a, Solvita},"
## [3] "  Editor = {Dengler, J. and Oldeland, J. and Jansen, F. and Chytr<fd>, M. and Ewald, J. and Fi
## [4] "  Journal = {Biodiversity & Ecology},"
## [5] "  Pages = {319<U+0096>320},"
## [6] "  Title = {Database Dry Grasslands in the Nordic and Baltic region},"
## [7] "  Volume = {4},"
## [8] "  Year = {2012},"
## [9] "  Doi = {10.7809/b-e.00114},"
## [10] "  Url = {https://doi.org/10.7809/b-e.00114},"
## [11] "  Language = {en}"
## [12] "}"
## [13] ""
## [14] ""
## [15] "@Article{biurrun2012a,"
## [16] "  Author = {Biurrun, Idoia and Garc{\\"{\\i}}a-Mijangos, Itziar and Campos, Juan A and Herrera, M
## [17] "  Editor = {Dengler, J. and Oldeland, J. and Jansen, F. and Chytr<fd>, M. and Ewald, J. and Fi
## [18] "  Journal = {Biodiversity & Ecology},"
## [19] "  Pages = {328},"
## [20] "  Title = {Vegetation-Plot Database of the University of the Basque Country (BIOVEG)},"
## [21] "  Volume = {4},"
## [22] "  Year = {2012},"
## [23] "  Doi = {10.7809/b-e.00121},"
## [24] "  Language = {en}"
## [25] "}"
```

Convert to reference list

```
mybib <- RefManagerR::ReadBib("_output/demo.bib", check = FALSE)
```

```
## Loading required namespace: bibtex
```

mybib

6

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<https://doi.org/10.7809/b-e.00069>>.

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En. In: *_Ecological Monographs_* 81 (2011), p. 25-41. DOI:
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##

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[11] J. Dengler and S. R\=usi\cna. "Database Dry Grasslands in the
Nordic and Baltic region". En. In: *_Biodiversity & Ecology_* 4 (2012).
Ed. by J. Dengler, J. Oldeland, F. Jansen, M. Chytrý, J. Ewald, M.
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Database (HelNatVeg)". En. In: *_Biodiversity & Ecology_* 4 (2012). Ed.
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and J. Schaminée, p. 388. DOI: 10.7809/b-e.00177. <URL:
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```
## 10.1127/phyto/2016/0109.
##
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## _Biodiversity & Ecology_ 4 (2012). Ed. by J. Dengler, J. Oldeland, F.
## Jansen, M. Chytrý, J. Ewald, M. Finckh and J. Schaminée, p. 340. DOI:
## 10.7809/b-e.00131.
```

sessionInfo()

```
sessionInfo()
```

```
## R version 4.0.1 (2020-06-06)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19042)
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=English_United Kingdom.1252
## [2] LC_CTYPE=English_United Kingdom.1252
## [3] LC_MONETARY=English_United Kingdom.1252
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United Kingdom.1252
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] bib2df_1.1.1      RefManager_1.3.0  rnatualearth_0.1.0
## [4] raster_3.4-5      sp_1.4-5          sf_0.9-7
## [7] forcats_0.5.1     stringr_1.4.0     dplyr_1.0.4
## [10] purrr_0.3.4       readr_1.4.0       tidyr_1.1.2
## [13] tibble_3.0.6      ggplot2_3.3.3     tidyverse_1.3.0
##
## loaded via a namespace (and not attached):
## [1] Rcpp_1.0.6        lubridate_1.7.9.2 lattice_0.20-41    class_7.3-18
## [5] assertthat_0.2.1  digest_0.6.27     utf8_1.1.4        plyr_1.8.6
## [9] R6_2.5.0          cellranger_1.1.0  backports_1.2.1    reprex_1.0.0
## [13] evaluate_0.14     e1071_1.7-4       highr_0.8          httr_1.4.2
## [17] pillar_1.5.0      rlang_0.4.10      readxl_1.3.1       rstudioapi_0.13
## [21] rmarkdown_2.7     labeling_0.4.2    rgdal_1.5-23       munsell_0.5.0
## [25] broom_0.7.5       compiler_4.0.1    modelr_0.1.8       xfun_0.21
```

## [29] pkgconfig_2.0.3	rgeos_0.5-5	htmltools_0.5.1.1	tidyselect_1.1.0
## [33] codetools_0.2-18	fansi_0.4.2	crayon_1.4.1	dbplyr_2.1.0
## [37] withr_2.4.1	humaniformat_0.6.0	grid_4.0.1	jsonlite_1.7.2
## [41] gtable_0.3.0	lifecycle_1.0.0	DBI_1.1.1	magrittr_2.0.1
## [45] units_0.7-0	scales_1.1.1	bibtex_0.4.2.3	KernSmooth_2.23-18
## [49] cli_2.3.0	stringi_1.5.3	farver_2.0.3	fs_1.5.0
## [53] xml2_1.3.2	ellipsis_0.3.1	generics_0.1.0	vctrs_0.3.6
## [57] tools_4.0.1	glue_1.4.2	hms_1.0.0	yaml_2.2.1
## [61] colorspace_2.0-0	classInt_0.4-3	rvest_0.3.6	knitr_1.31
## [65] haven_2.3.1			