



Towards improving the explanatory power in landscape genomics

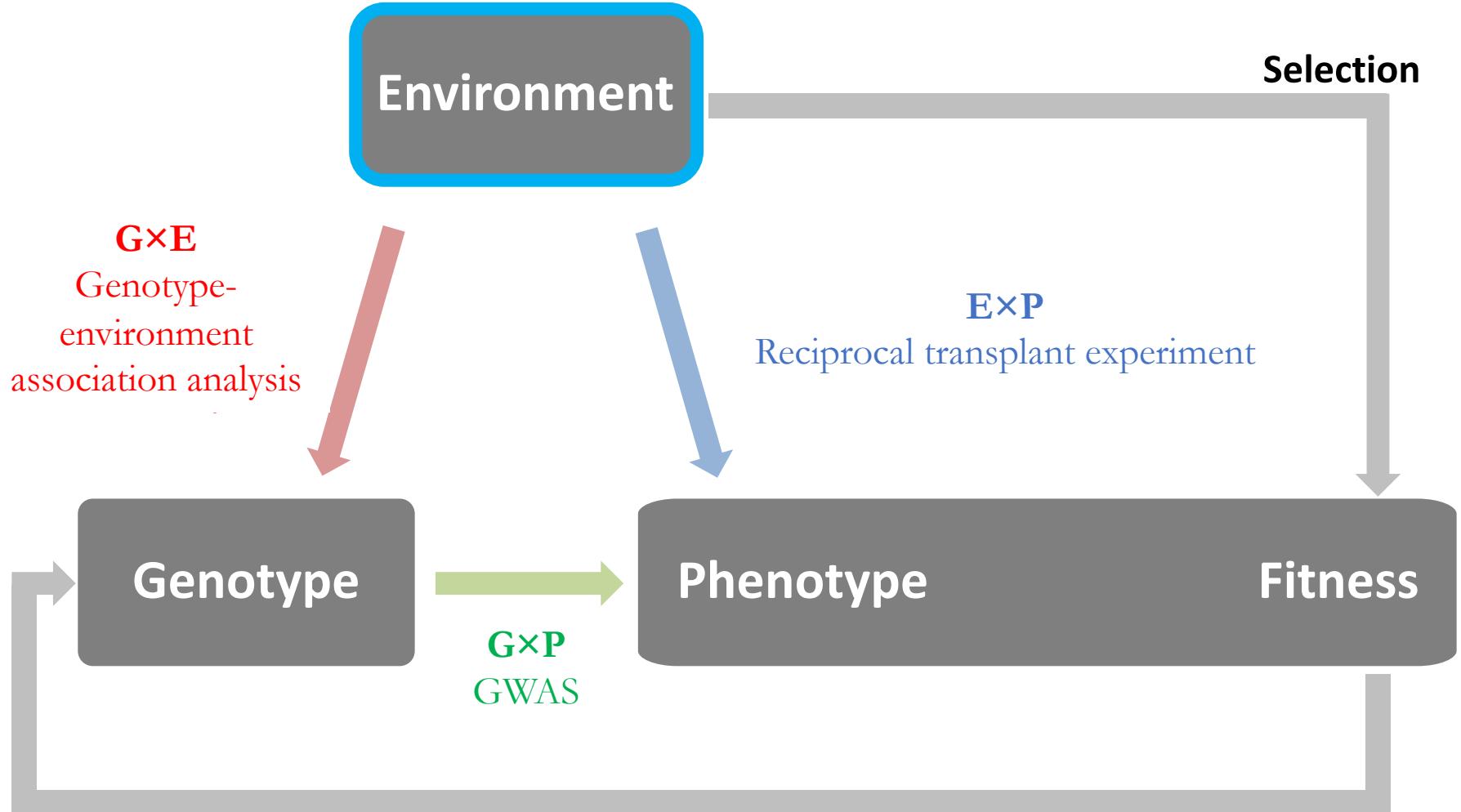
Benjamin Dauphin

Swiss Federal Research Institute WSL

Summer School SSMPG - 21.09.2022



Focus



Outline

I. General introduction

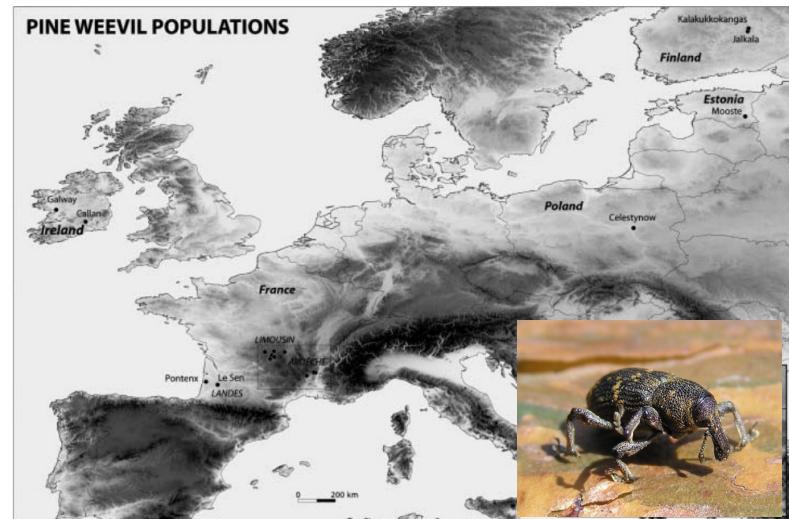
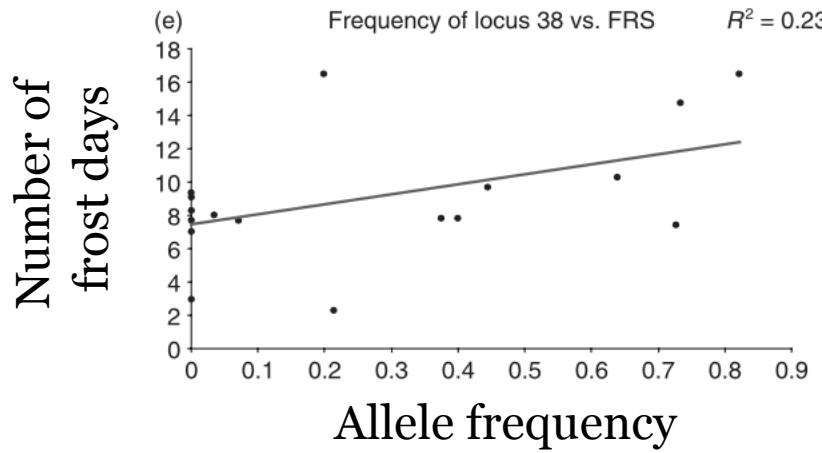
II. A brief portray of environmental data

III. A short tutorial on how to handle environmental data in R

IV. Reduction of environmental variables

Landscape genomics

a field that searches for associations between environmental conditions and adaptive genetic variation in natural populations ([Joost et al. 2007](#)).

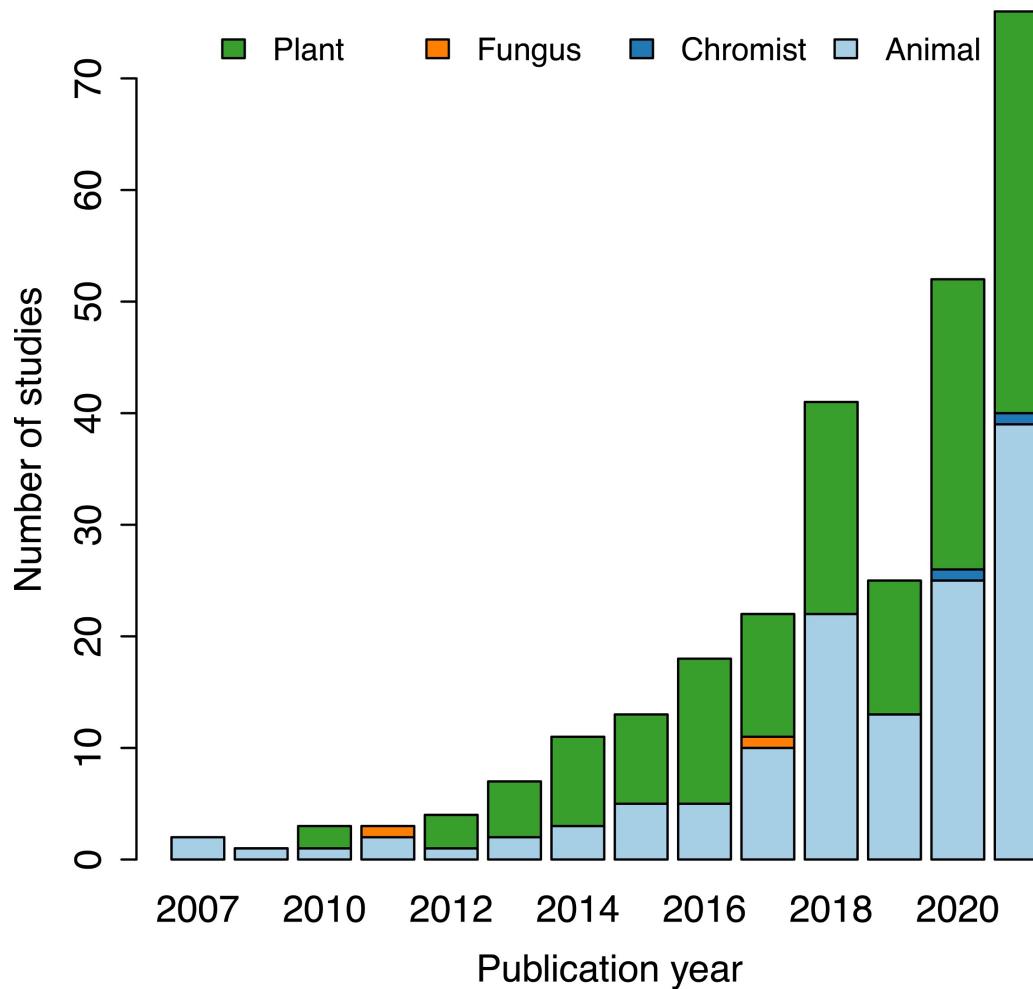


Ecological genomics

a field that aims to understand the genetic mechanisms underlying responses of organisms to their natural environment, **including experimental approaches** ([Ungerer et al. 2008](#))

Where are we with landscape genomics?

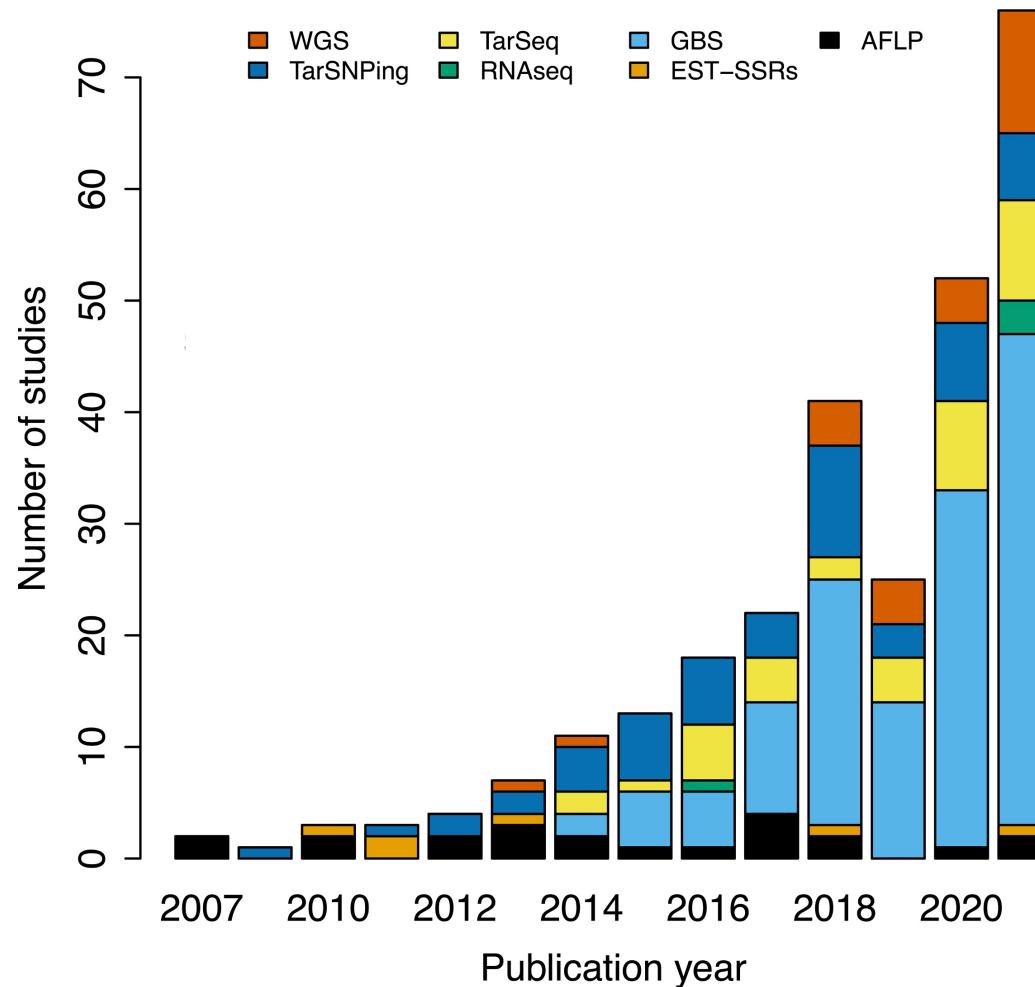
Landscape genomics is still a growing field...



Survey of 278 studies

Dauphin et al. in review

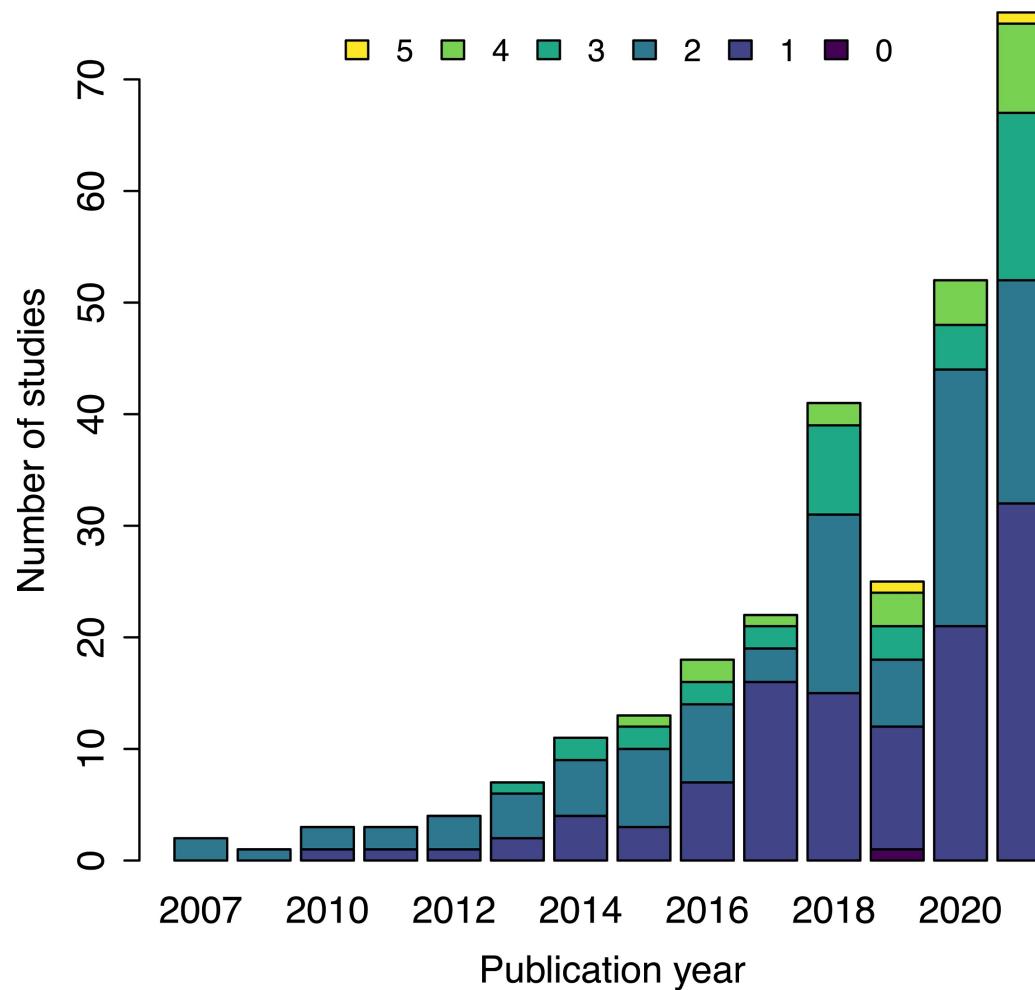
taking benefit of (new) sequencing techniques...



Survey of 278 studies

Dauphin et al. in review

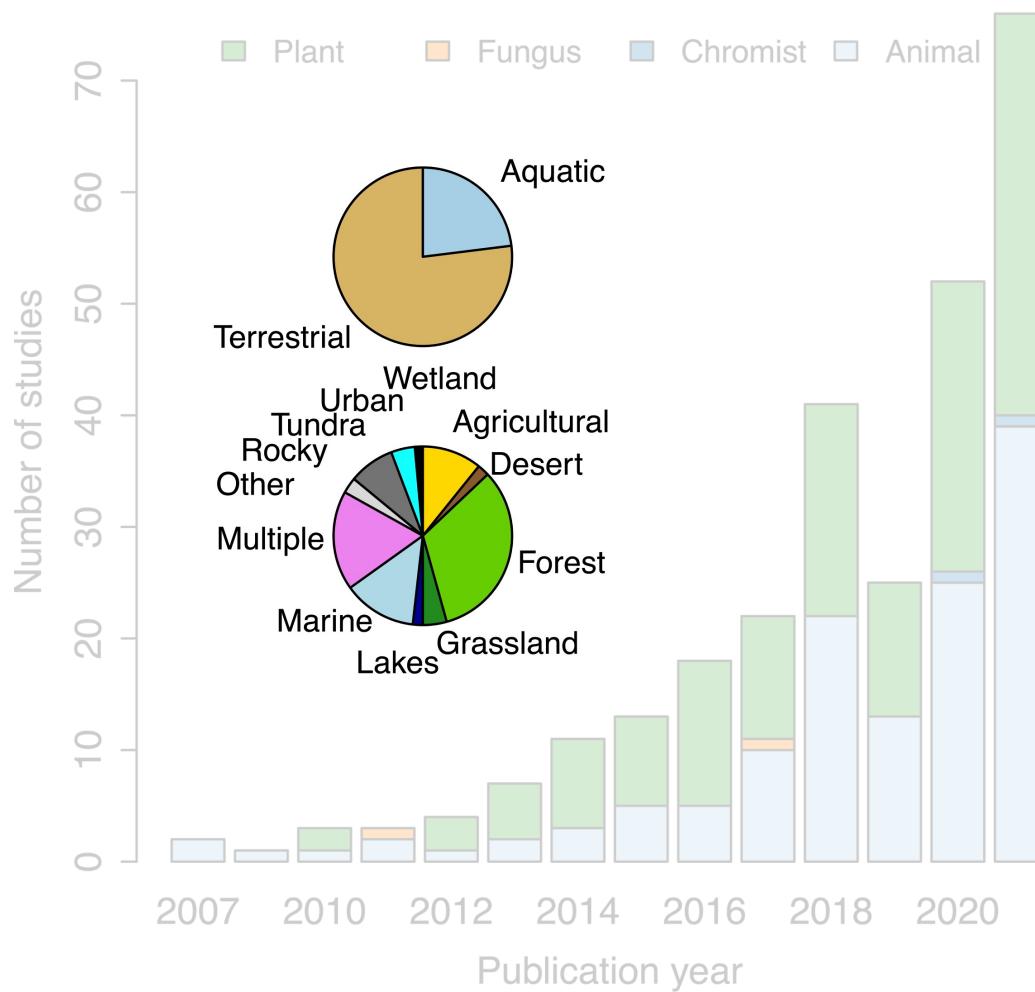
... and increasingly integrative environmental datasets



Survey of 278 studies

Dauphin et al. in review

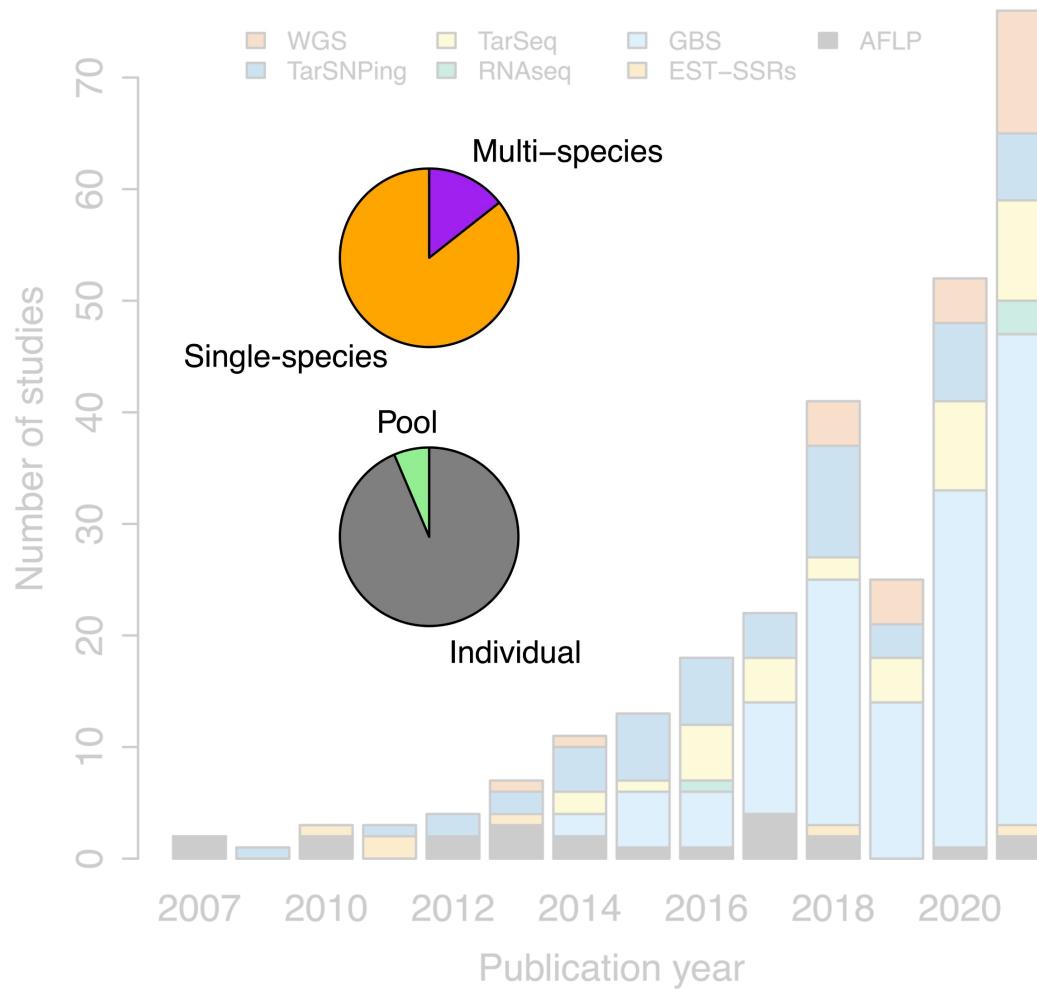
However, some ecosystems are under-represented...



Survey of 278 studies

Dauphin et al. in review

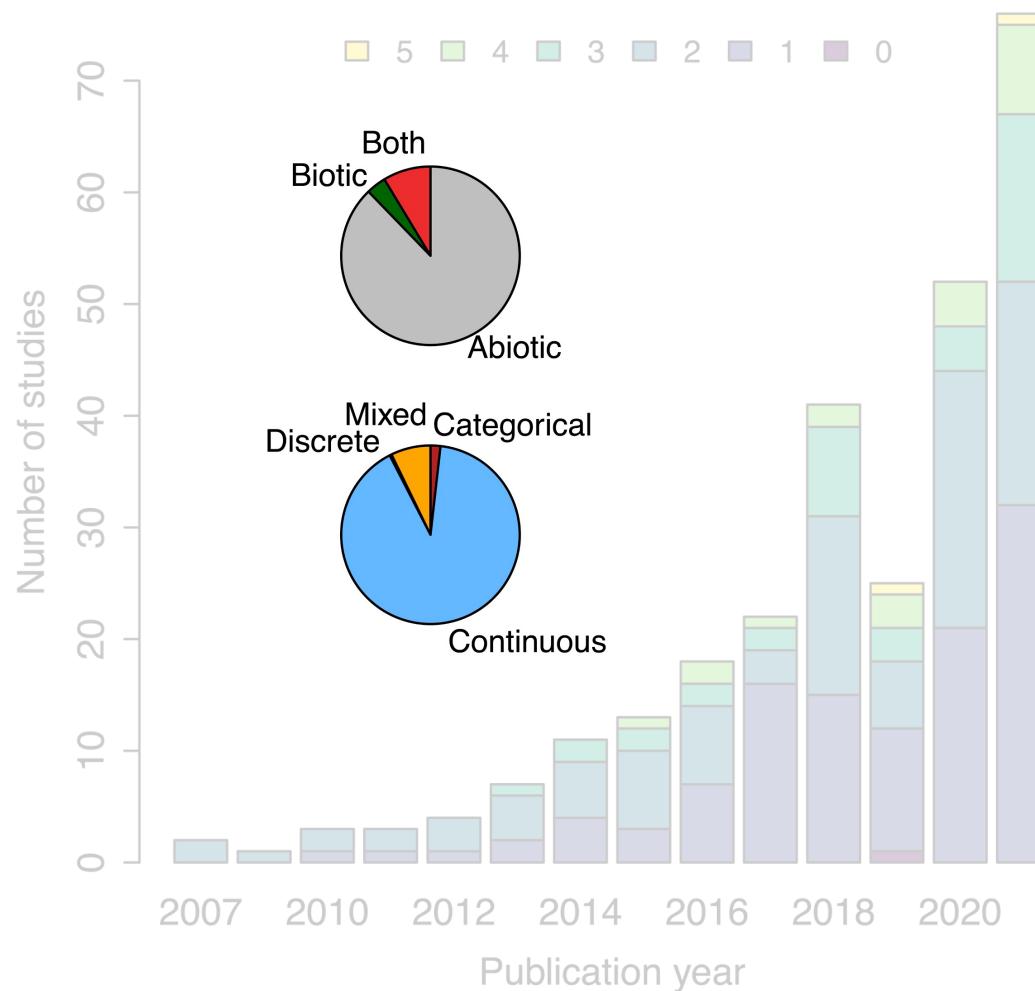
most studies focused on a single species...



Survey of 278 studies

Dauphin et al. in review

... and considered abiotic environmental conditions

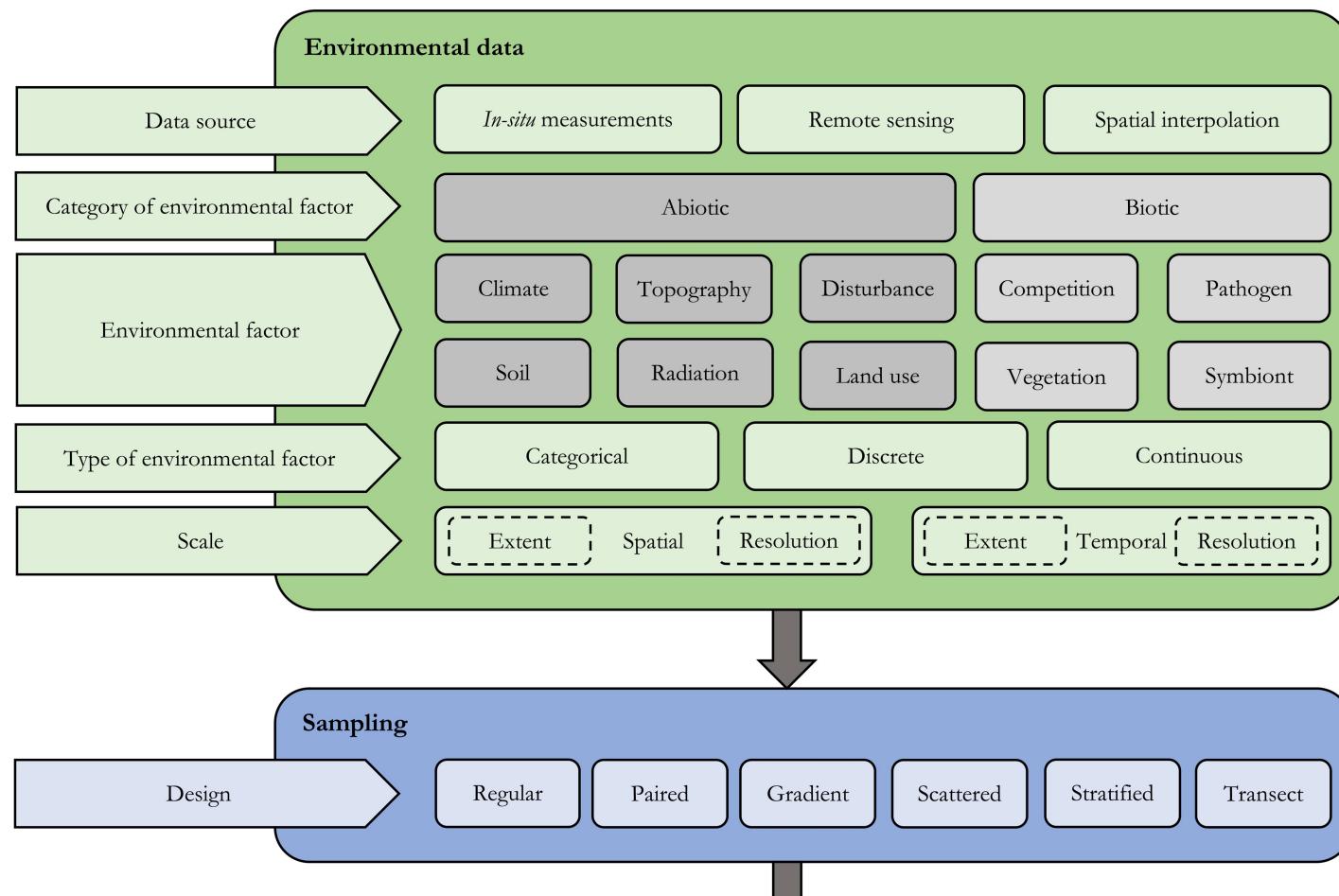


Survey of 278 studies

Dauphin et al. in review

Where shall we start when investigating local adaptation?

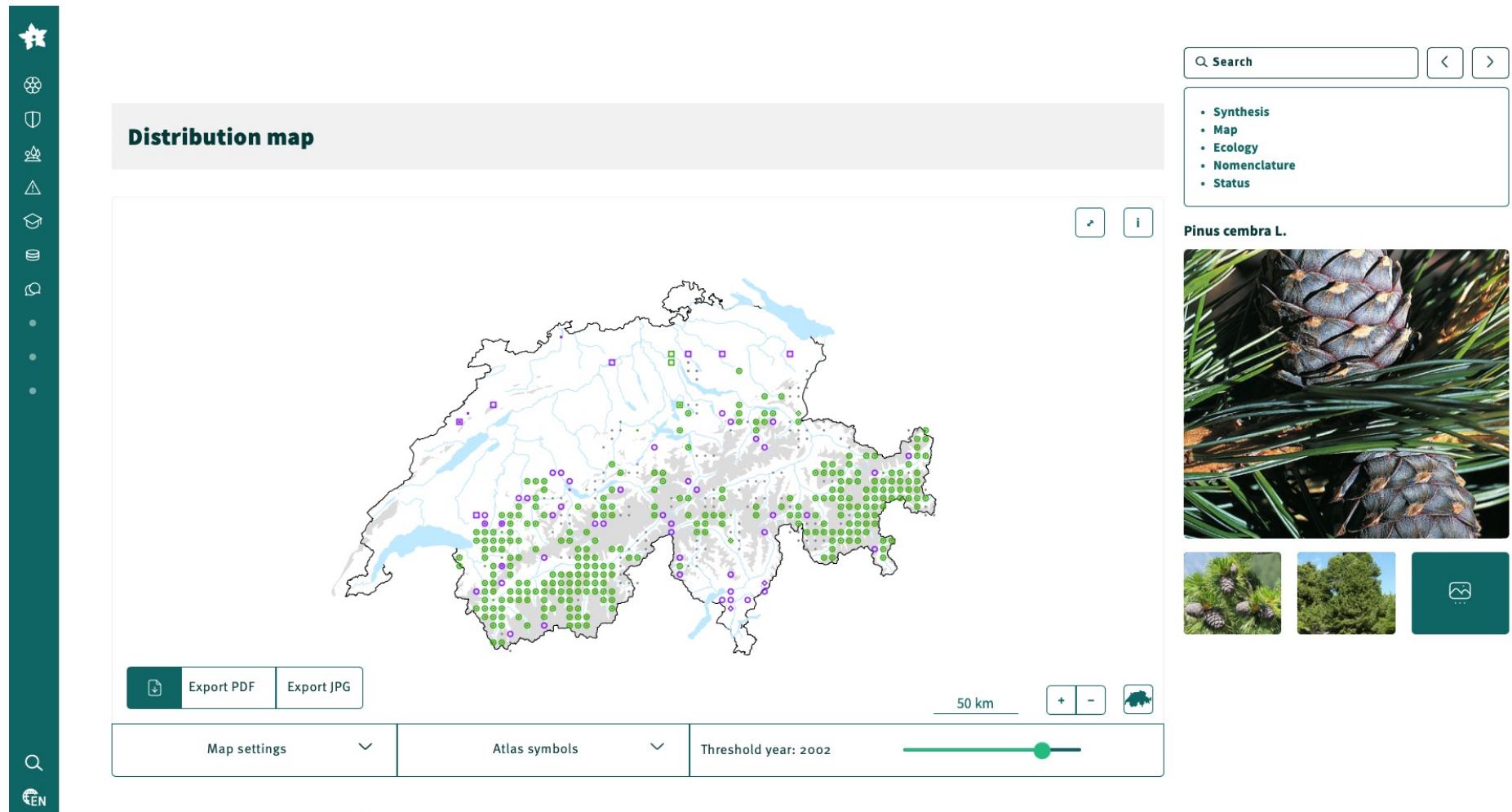
Explore environmental data before sampling



... with some precautions

Dauphin et al. in review

Use the known species range to identify gradient(s) / contrast(s)



<https://www.infoflora.ch/en/flora/pinus-cembra.html>

Although curation of occurrence data may be required

Get data How-to Tools Community About

15,253 OCCURRENCES 2 INFRASPECIES

Classification

Select a species

Kingdom Plantae

Phylum Tracheophyta

Class Pinopsida

Order Pinales

Family Pinaceae

Genus *Pinus* L.

Species *Pinus cembra* L.

- = *Apinus cembra* (L.) Neck.
- = *Apinus cembra* (L.) Neck. ex Rydb.
- = *Cembra montana* Opiz
- = *Pinea cembra* (L.) Opiz
- = *Pinus cembra* f. *columnaris* (Hellm. ex Beissn.) Rehder
- = *Pinus cembra* subsp. *communis* Endl.

OVERVIEW 1 TREATMENT METRICS REFERENCE TAXON

1,317 OCCURRENCES WITH IMAGES

SEE GALLERY

14,114 GEOFERENCED RECORDS

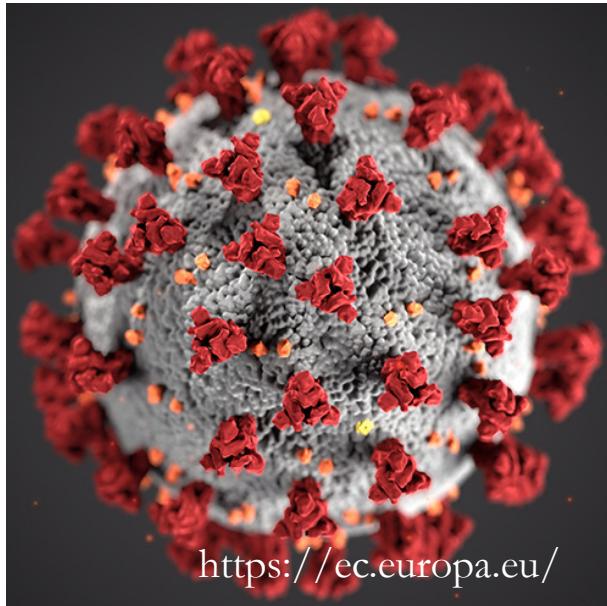
Generated 38 minutes ago © OpenStreetMap contributors, © OpenMapTiles, GBIF.

Any year 1684 - 2022 EXPLORE AREA

<https://www.gbif.org/species/5285134>

Environment, from the infinitely small to the infinitely large

~ 9–12 nanometres



<https://ec.europa.eu/>

~ 115 metres



www.livescience.com/

But the largest organism is not the one we think...

earth

Home Attenborough's Story of Life Big Questions Discoveries Video Earth by email About us

Is This Fungus the World's Biggest Organism?

The 'Humungous Fungus' is older and bigger than previously thought. This enormous honey fungus has been revisited and reanalysed using scientific techniques that had yet to be invented when it was first discovered in the 1980s.

Genomic analysis and GPS show how far the fungus has spread, and surprisingly how little genetic variance it has developed in its long lifespan. The fungus is now thought to be at least 2,500 years old. Researchers say understanding why it lacks genetic mutation might be useful in understanding cancer.

Record Breakers | Fungus

The largest living thing on Earth is a humongous fungus

Forget blue whales and giant redwood trees. The biggest living organism is over 2 miles across, and you'll hardly ever see it

<http://www.bbc.com/>

And the oldest organism is ...



GETTY IMAGES

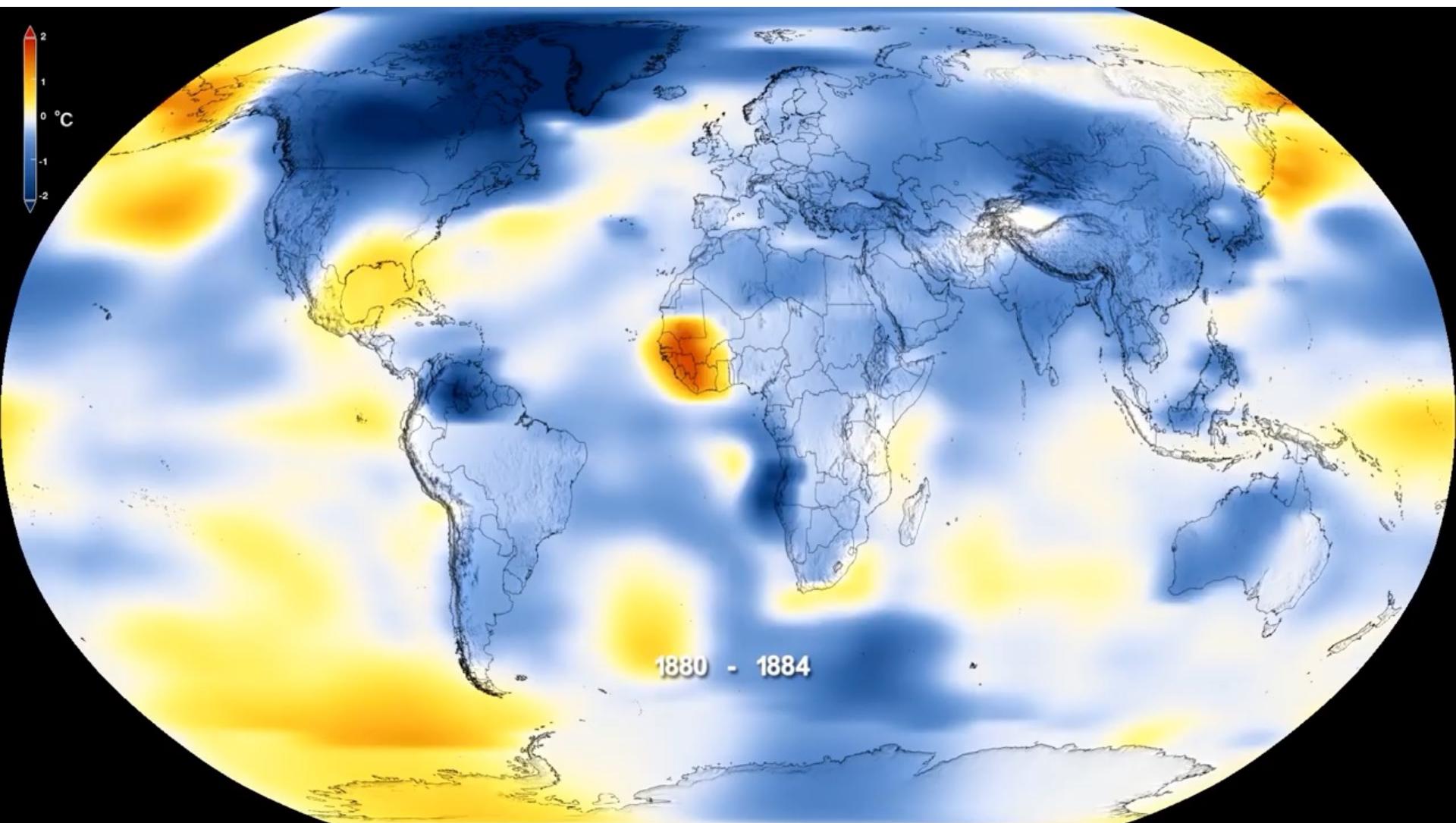
| The Great Basin bristlecone pine is found in the western United States

<http://www.bbc.com/>

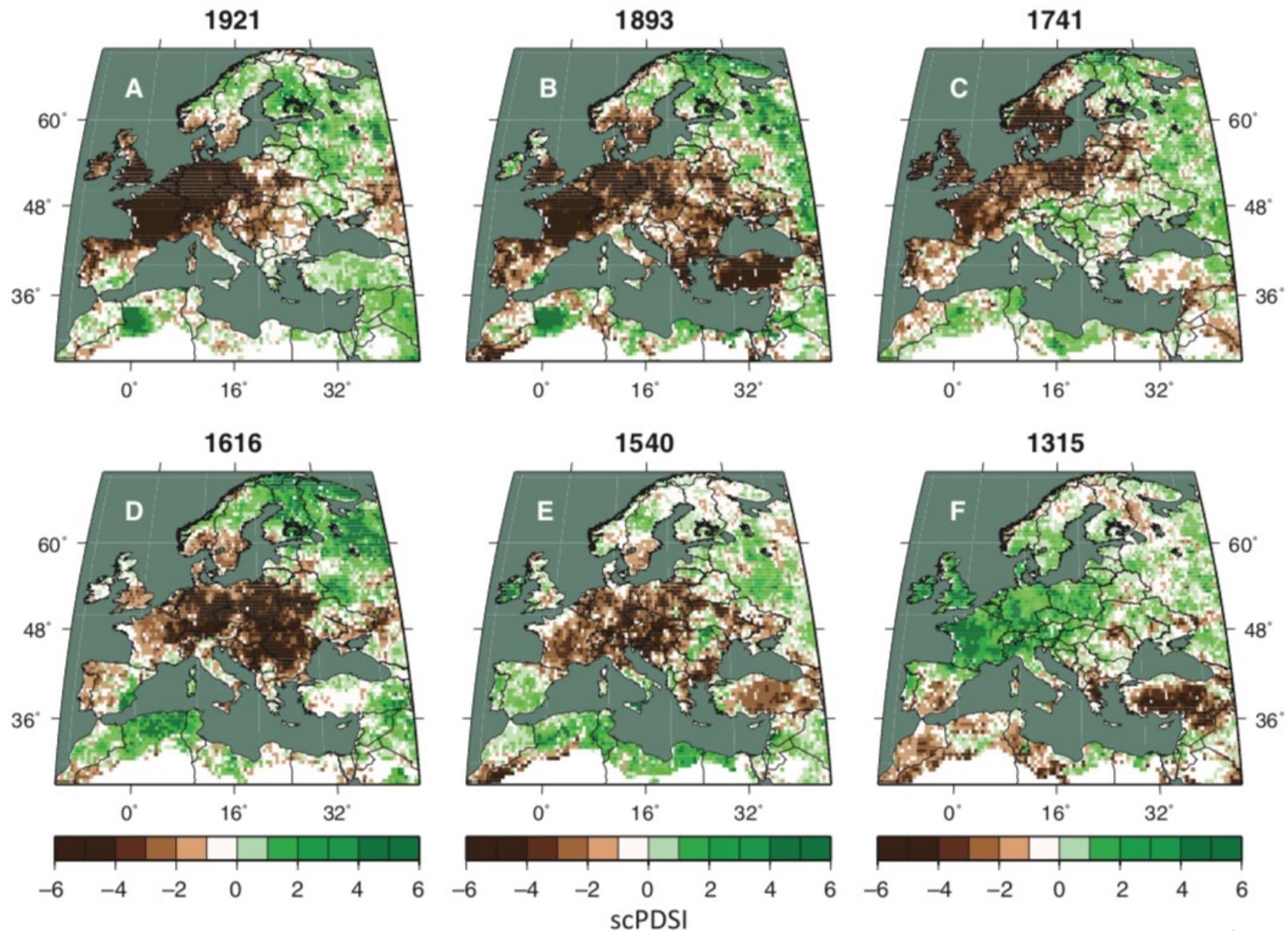
Species	Age	Type	ID	Location	Collector(s), Dater(s), Reference
4000+ years					
<i>Pinus longaeva</i> , Great Basin bristlecone pine	4900**	XD	WPN-114, Prometheus	Wheeler Peak, Nevada, USA	Currey 1965; Salzer and Baisan 2018
<i>Pinus longaeva</i> , Great Basin bristlecone pine	4850*	XD	Methuselah	White Mountains, California, USA	Ed Schulman, Tom Harlan
3000+ years					
<i>Fitzroya cupressoides</i> , alerce	3622	XD		Chile	Lara and Villalba 1993
<i>Sequoiadendron giganteum</i> , giant sequoia	3266**	XD	CBR26	Sierra Nevada, California, USA	Malcolm Hughes, Ramzi Touchan, Ed Wright
<i>Sequoiadendron giganteum</i> , giant sequoia	3220**	XD	D-21	Sierra Nevada, California, USA	Douglass 1919
<i>Sequoiadendron giganteum</i> , giant sequoia	3075**	XD	D-23	Sierra Nevada, California, USA	Douglass 1919
<i>Sequoiadendron giganteum</i> , giant sequoia	3033**	XD	CMC 3	Sierra Nevada, California, USA	Tom Swetnam, Chris Baisan
2000+ years					
<i>Juniperus occidentalis</i> , western juniper	2675**	XD	Scofield Juniper	Sierra Nevada, California, USA	Miles and Worthington 1998

<http://www.rmtrr.org/>

Temperature anomalies through space from 1880



Spatial distribution and severity of past drought events



Cook et al. 2015

Outline

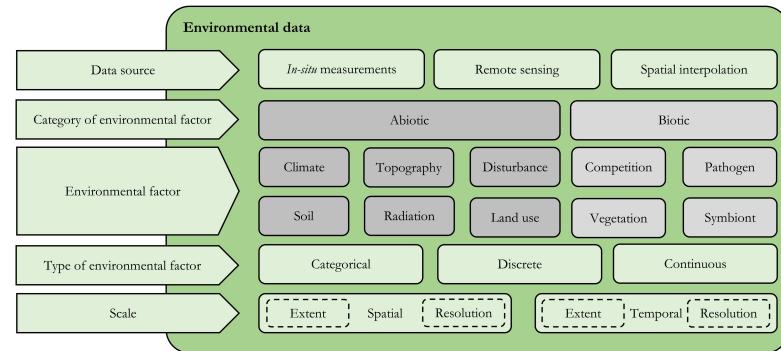
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Source of environmental data



Measures on the field....



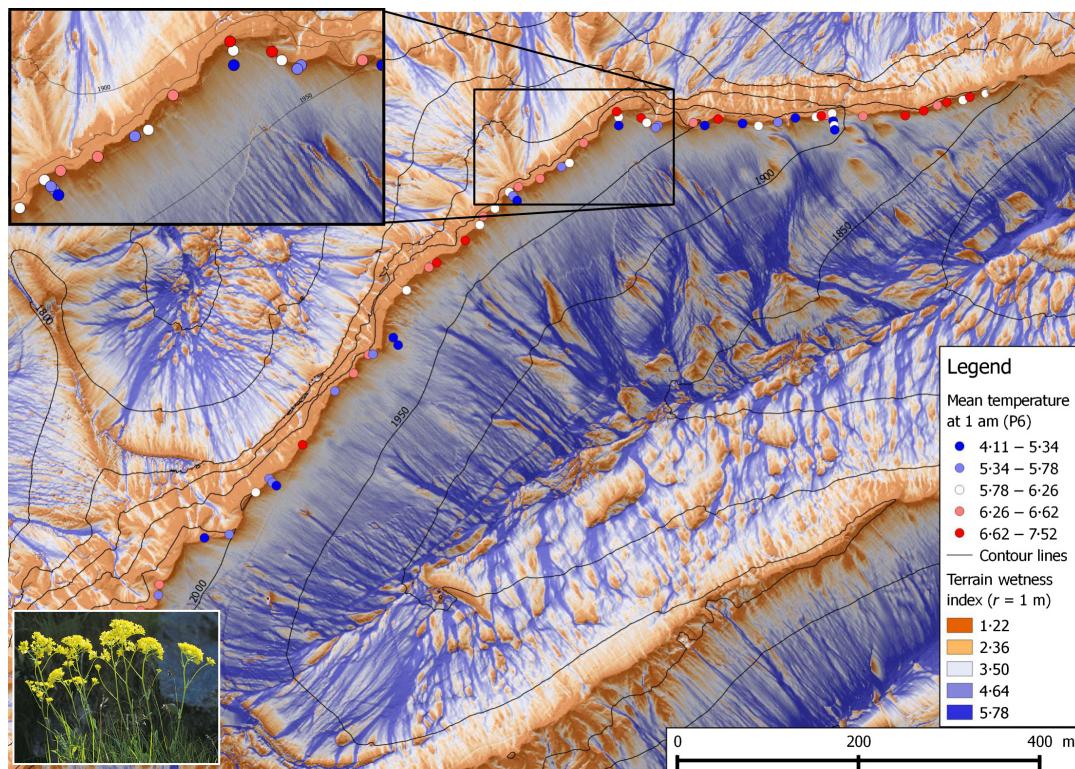
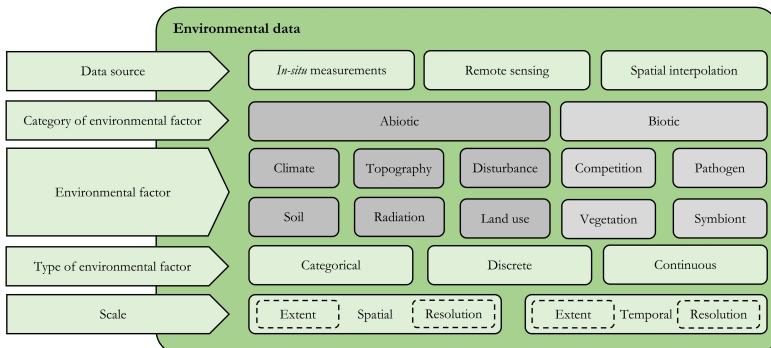
or modelling from observations



Categories of environmental data

- Abiotic:
 - Soil
 - Climate
 - Nutriments
 - Topography
 - ...

- Biotic:
 - Pathogens
 - Symbionts
 - Herbivory
 - Pollinators
 - Vegetation
 - Competition
 - ...



Type of environmental factors

Categorical ...

- A variable that contains a **finite number of categories** or distinct groups

Discrete ...

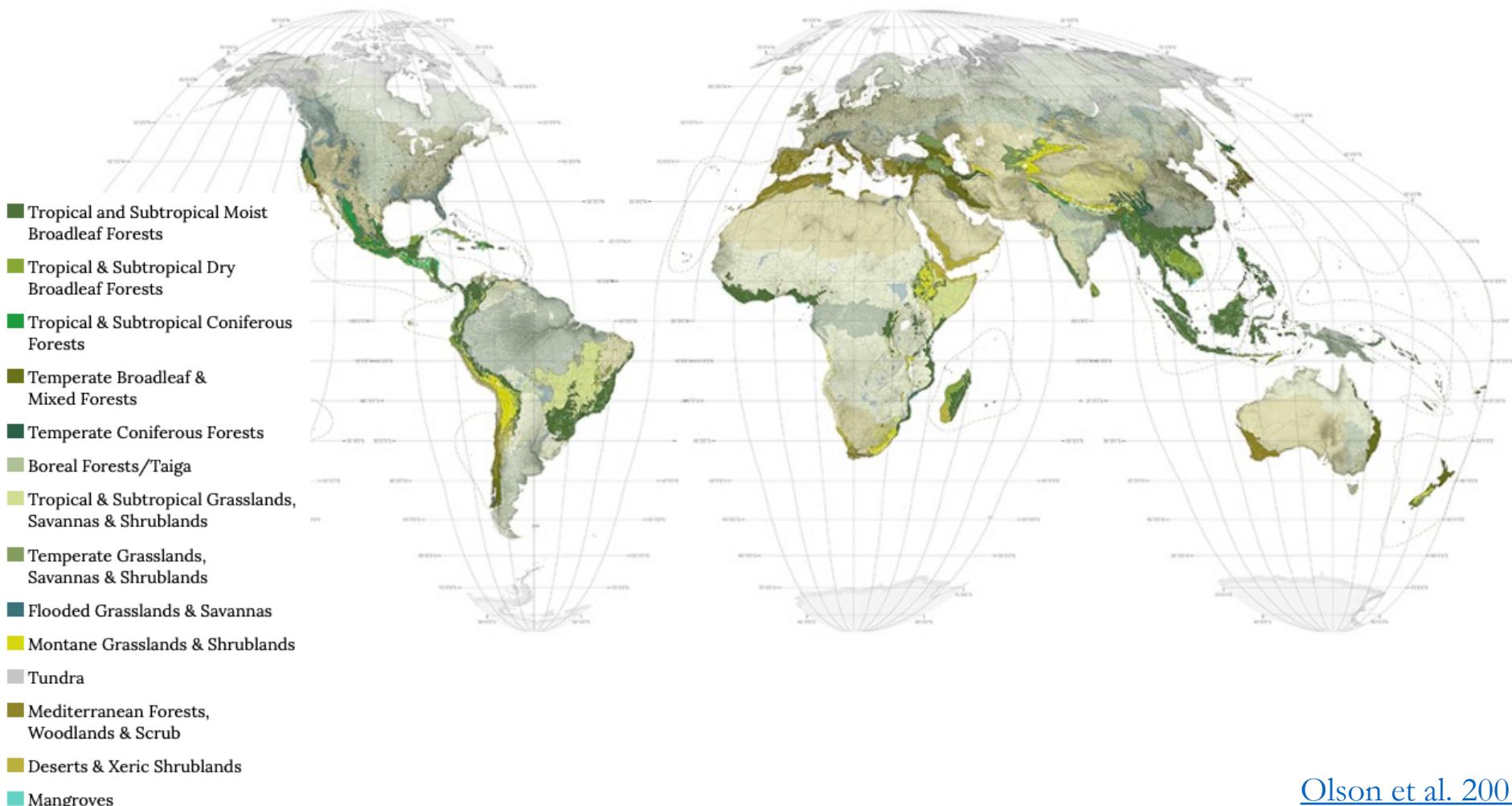
- A variable that can take **only certain values** along an interval

... versus Continuous environmental data

- A variable that can take **any value** at any point along an interval

Example of categorical environmental variable

- Geographic (e.g., biogeographical areas, **ecoregions**, among others...)



Example of discrete environmental variable

- Floristic bioindicators (e.g., Landolt indicators among others...)

Pinus cembra L.

Species 304900

Search Previous Next

Pinaceae / Pinus / *Pinus cembra* L.

Synthesis Map Ecology Nomenclature Status

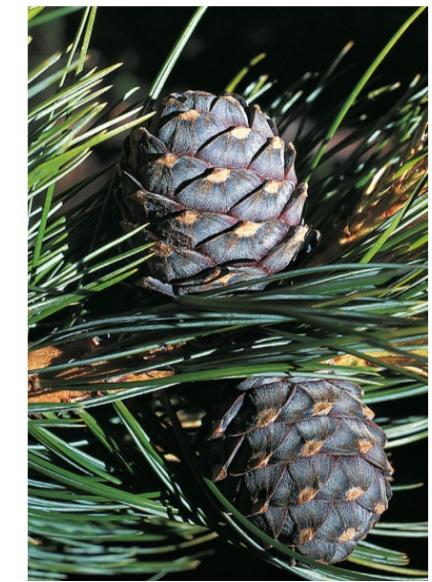
1034060

Ecological indicator values by Landolt & al. (2010)

Soil factors		Climatic factors		Salinity tolerance	
Humidity Value H	3	Light Value L	3	Salinity Index	--
Reaction Value R	2	Temperature factor T	2		
Nutrients value N	2	Continentality K	4		

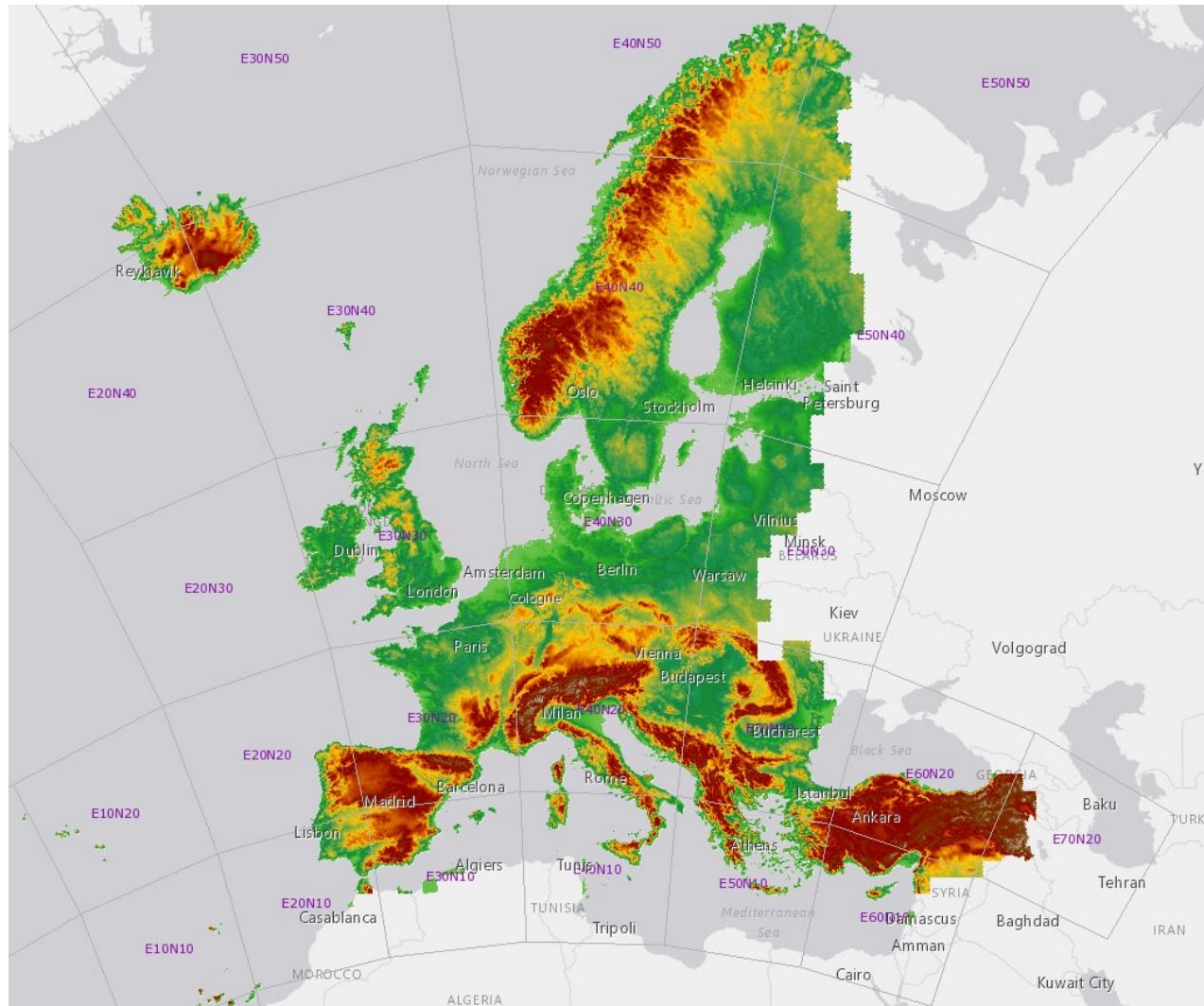
Nutrients value N	
1	very low in nutrients
2	low in nutrients
3	medium-poor to medium-rich in nutrients
4	rich in nutrients
5	very rich in nutrients

Light Value L	
1	very shady
2	shady
3	lighted areas
4	luminous
5	highly luminous



Example of continuous environmental variable

- Topography on a European scale with Copernicus



Some environmental databases

Soil

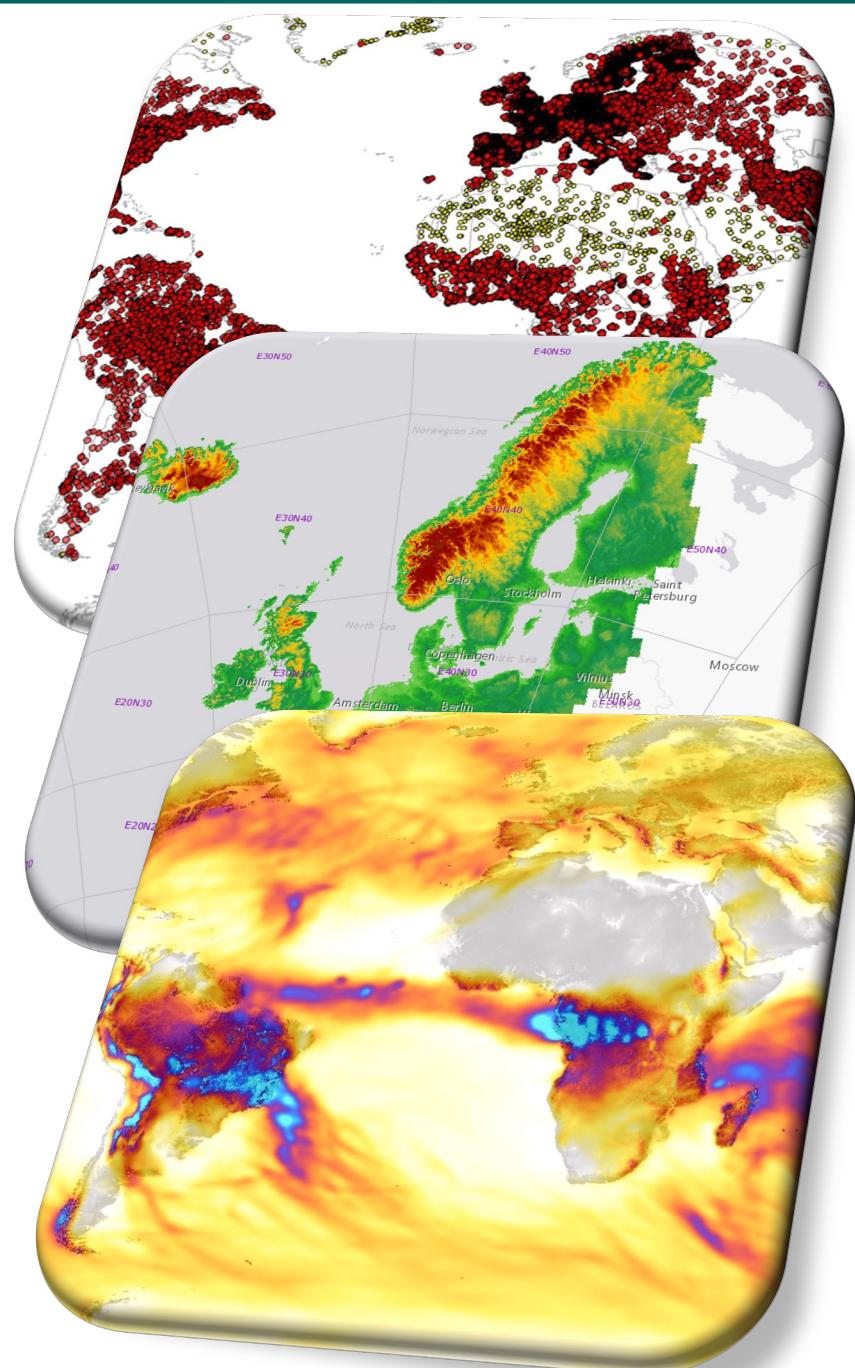
- [SoilGrids](#)
- [CORINE Land Cover](#)
- [US Land Cover Database](#)
- [Global soil temperature](#)
- ...

Topography

- [Copernicus EU-DEM](#)
- [GMTED2010](#)
- [SRTM](#)
- ...

Climate

- [CHELSA](#)
- [WorldClim](#)
- [World Ocean Database](#)
- ...



Why using CHELSA?



Climatologies at high resolution for the earth's land surface areas

Comparison to other high resolution climate products

CHELSA, as most other climate products is a ‘model’ that aims at representing reality as precise as possible. Nevertheless, it is just a model, and has its strength and weaknesses. Here, we want to provide a simple comparison to some other products such as [PRISM](#) or [WorldClim](#), to illustrate some of the differences between these models.

The comparison of the predictions with existing products and station data indicates an improvement in the spatio-temporal performance of the precipitation data based on cloud cover-informed downscaling. For this newly developed semi-mechanistic downscaling approach for daily precipitation, high resolution (30 arc sec) satellite-derived cloud frequency was incorporated.

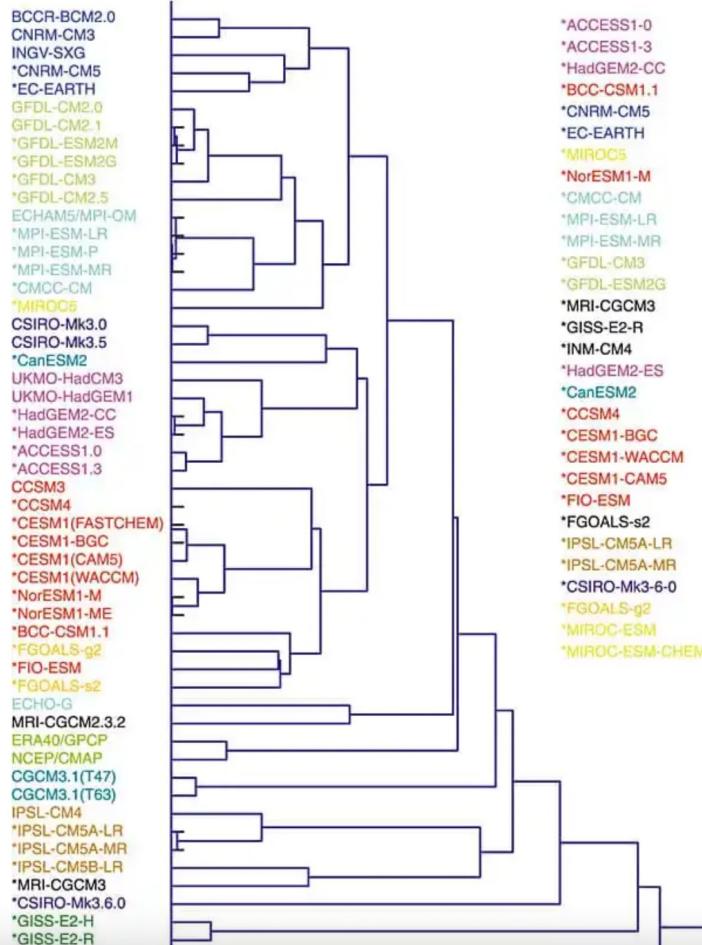
For more information see:

Karger, D.N., Wilson, A.M., Mahony, C., Zimmermann, N.E., Jetz, W. (2021): Global daily 1km land surface precipitation based on cloud cover-informed downscaling. **Scientific Data.** doi.org/10.1038/s41597-021-01084-6

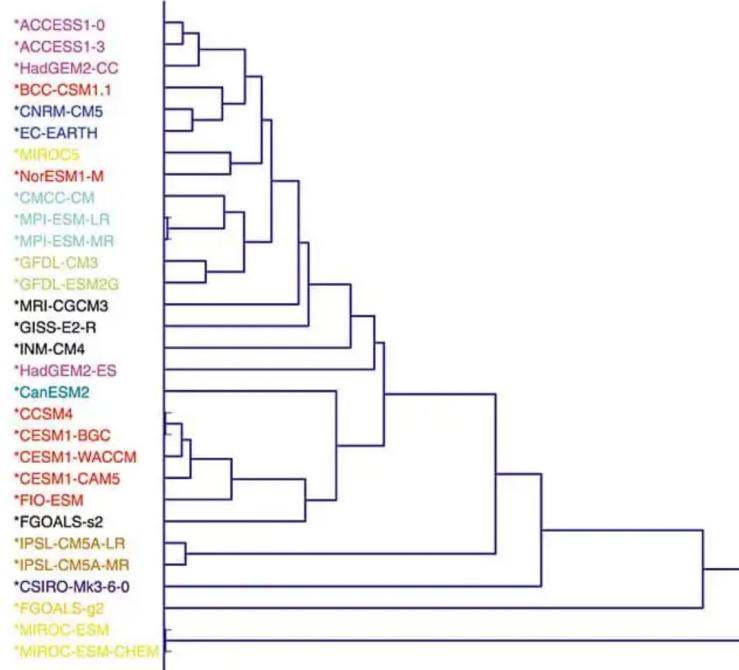
Selection of climate models

Consider those that are less interdependent under current and/or future conditions

a) Control state



b) Projected change RCP8.5



File formats of environmental data

Vector data:

- Points: e.g., occurrence data
- Lines: e.g., barrier to migration
- Polygons: e.g., species range

“shapefile” is the most common vector file type

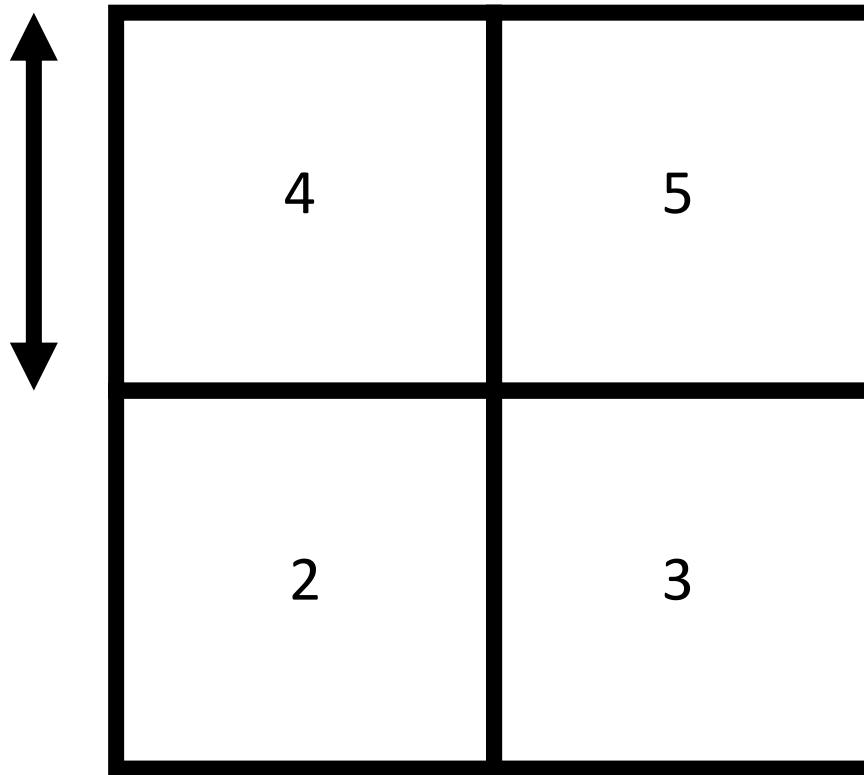
Raster data:

- Made of pixels regularly spaced (usually!)
- Pixels represent values (continuous) or classes (discrete)
- Possible to work at multiple spatial resolutions
(e.g. Leempoel et al 2015, Methods Ecol. Evol.)

“GeoTIFF”, “Esri Grid” and “ASCII Grid” are the most popular raster file types

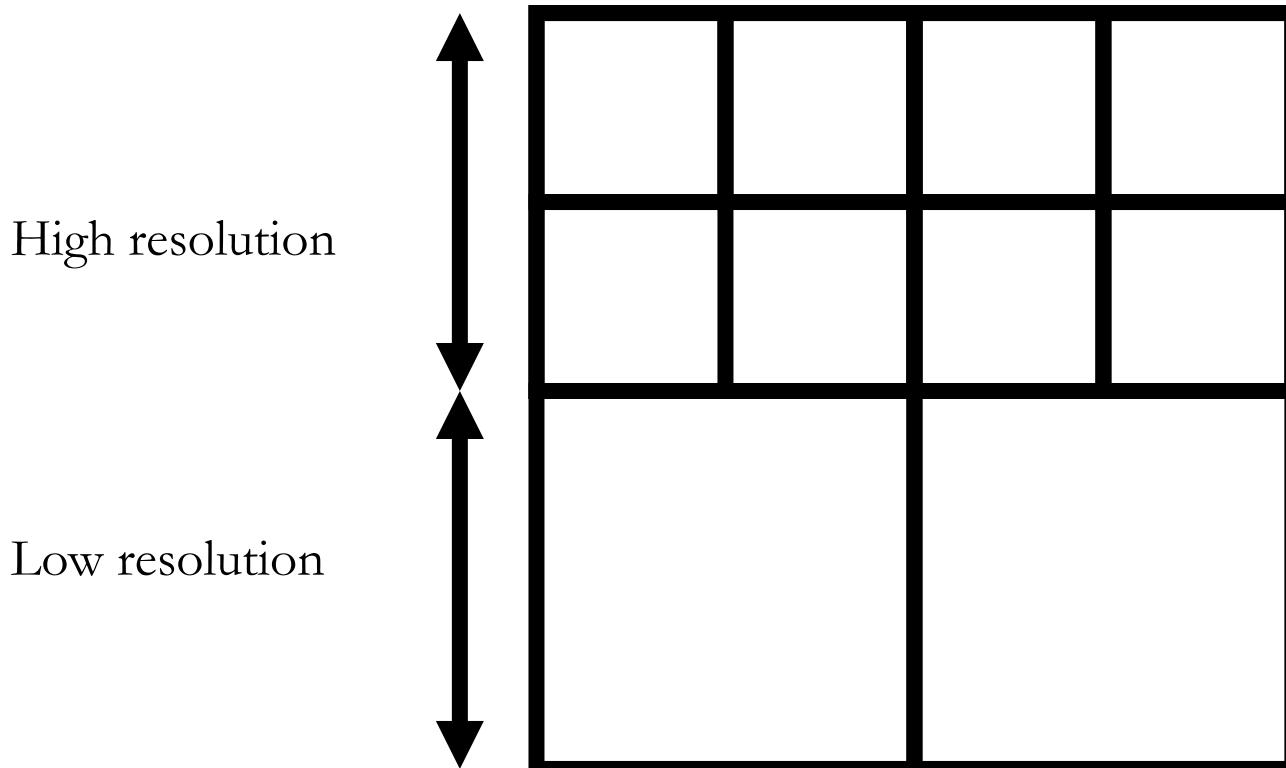
<https://gisgeography.com/gis-formats/>

Spatial resolution of interpolated environmental data



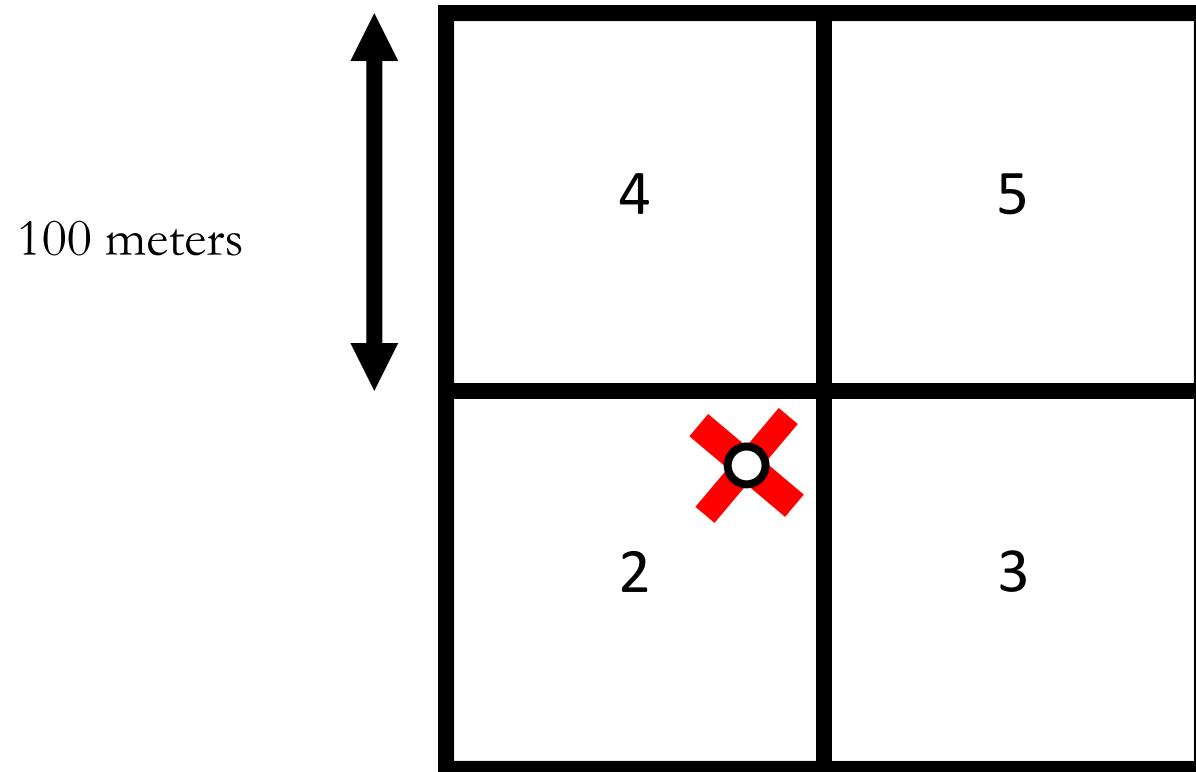
“Spatial resolution refers to the size of the smallest object that can be resolved on the ground. In a digital image, the resolution is limited by the pixel size, i.e. the smallest resolvable object cannot be smaller than the pixel size.”

Spatial resolution of interpolated environmental data

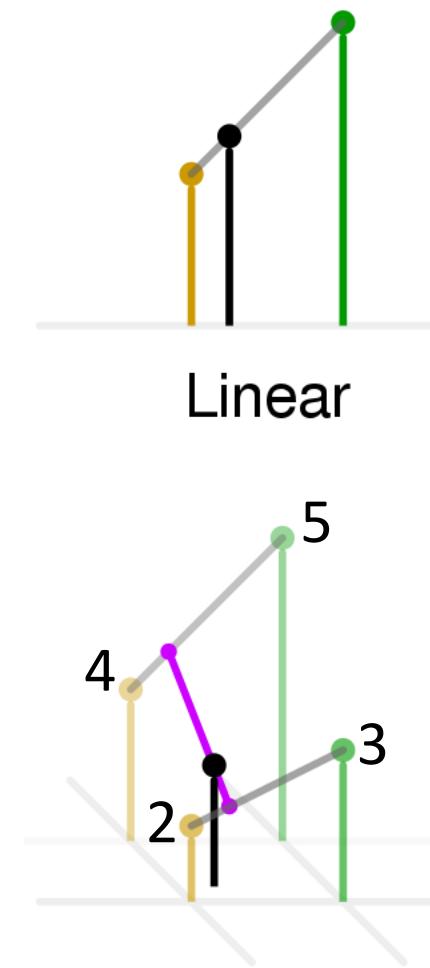


But the resolution does not inform about the accuracy of data !

The trick of data extraction with bilinear interpolation



Your geographic coordinates



Bilinear

Outline

I. General introduction

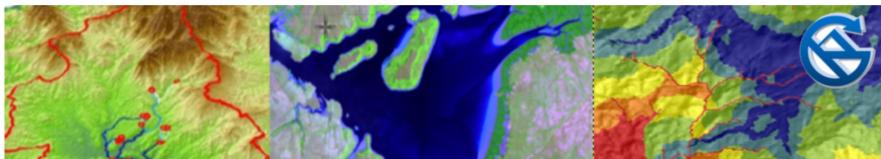
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Software used to handle environmental data

- R packages: e.g. raster, sf, rgdal, maps...
- QGIS: <https://www.qgis.org/en/site/>
- SAGA: <https://saga-gis.sourceforge.io/en/index.html>
- ArcGIS: <https://www.arcgis.com/index.html>



Saga GIS tutorials

About ▾ Working with Satellite imagery. ▾ Terrain Analysis ▾ Processing tools ▾ Applications ▾
Advanced Topics ▾ Downloads Trouble shooting Training Manual Videos ▾

Nick Eubank
Assistant Research Professor, Duke University SSRI

Home Research Educational Resources CV

R has a full library of tools for working with spatial data. This includes tools for both vector and raster data, as well as interfacing with data from other sources (like ArcGIS) and making maps.

<https://www.nickeubank.com/gis-in-r/>

Ben Gurion University of the Negev
Preface
0.1 Welcome
0.2 What is R?
0.3 R and analysis of spatial data
0.4 Other materials

Introduction to Spatial Data Programming with R

Michael Dorman

<http://132.72.155.230:3838/r/>

GenTree EU project – *Fagus sylvatica* populations

<https://www.gentree-h2020.eu>



Optimizing the management and sustainable
use of forest genetic resources in Europe

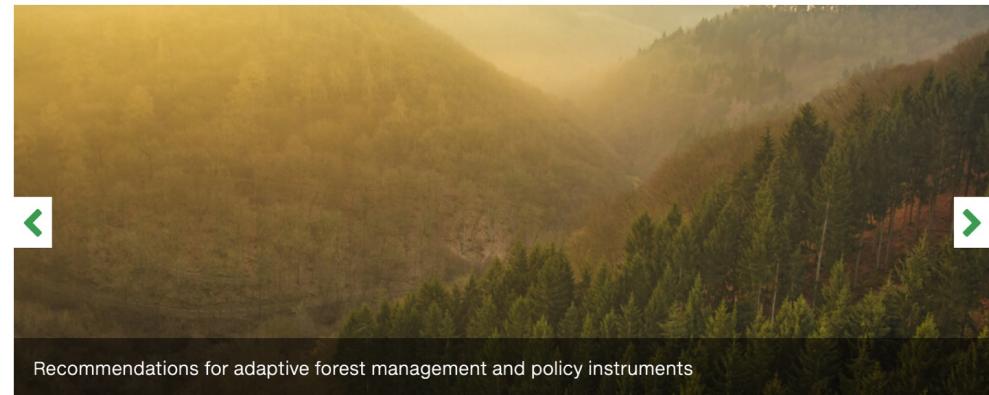


Horizon 2020
European Union Funding
for Research & Innovation

HOME ABOUT RESOURCES EVENTS NEWS



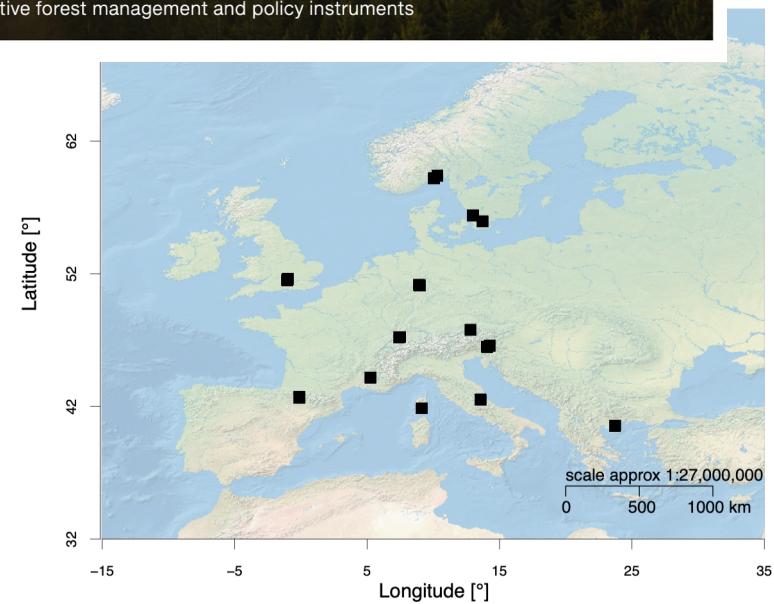
Quantify key processes affecting genomic diversity



Recommendations for adaptive forest management and policy instruments



GENTREE 2020 Avignon



Genetic, environmental, phenotypic, dendrochronological, and leaf trait data of trees species across Europe

Exome capture sequencing:

- *Betula pendula* (silver birch)
 - *Fagus sylvatica* (European beech)
 - *Picea abies* (Norway spruce)
 - *Pinus pinaster* (maritime pine)
 - *Pinus sylvestris* (Scots pine)
 - *Populus nigra* (European black poplar)
 - *Quercus petraea*. (sessile oak)
- 18-24 populations; ~25 indiv./pop.

Single Primer Enrichment Technology genotyping:

- *Abies alba* (silver fir)
 - *Pinus cembra* (Swiss stone pine)
 - *Pinus halepensis* (Aleppo pine)
 - *Pinus nigra* (European black pine)
 - *Taxus baccata* (English yew)
- 10-14 populations; ~25 indiv./pop.



GigaScience, 10, 2021, 1–13

doi: 10.1093/gigascience/giab010
DATA NOTE

DATA NOTE

The GenTree Platform: growth traits and tree-level environmental data in 12 European forest tree species

Lars Opgenoorth ^{1,2,*}, Benjamin Dauphin ², Raquel Benavides ³, Katrin Heer ¹, Paraskevi Alizoti ⁴, Elisabet Martínez-Sancho², ...

...

SCIENTIFIC DATA

OPEN

DATA DESCRIPTOR

The GenTree Dendroecological Collection, tree-ring and wood density data from seven tree species across Europe

Elisabet Martínez-Sancho et al.^{*}

...

DATA PAPER

Global Ecology and Biogeography

A journal of
WILEY
Biodiversity

The GenTree Leaf Collection: Inter- and intraspecific leaf variation in seven forest tree species in Europe

Raquel Benavides¹ | Bárbara Carvalho¹ | Cristina C. Bastias^{1,2} | David López-Quiroga¹ | Antonio Mas¹ | Stephen Cavers³ | Alan Gray³ |

...

Tutorial and online resources – GitHub project

1. Get a background map and project it

```
# Download manually the raster layer and paste the zip file into the dat/ folder
browseURL("http://www.naturalearthdata.com/downloads/10m-raster-data/10m-natural-earth-2/",
           browser=getOption("browser"), encodeIfNeeded=F)
unzip(zipfile=paste(path2,"NE2_LR_LC.zip",sep=""), exdir=paste(path2,"NE2_LR_LC",sep=""))

# Load it
back.map <- stack(paste(path2,"NE2_LR_LC/NE2_LR_LC.tif",sep=""))

# Define geographic coordinate system
browseURL("https://epsg.io", browser=getOption("browser"), encodeIfNeeded=F)
crs(back.map) <- CRS("+init=EPSG:4326")

# Global map
plotRGB(back.map, axes=F)

# Europe map
plotRGB(back.map, ext=eu.extent, axes=F)
```

The screenshot shows the Natural Earth website's main page. At the top, there's a navigation bar with links for Home, Features, Downloads, Blog, Issues, Corrections, and About. Below the navigation bar, a banner for 'Natural Earth' features a globe icon and text about free vector and raster map data at 1:10m, 1:50m, and 1:110m scales. A search bar is also present. The main content area is titled '1:10m Natural Earth II'. It includes a brief description of the dataset, mentioning its idealized nature and historical significance. Below the description is a small thumbnail image of a world map from the dataset. To the right of the thumbnail, there are download links for 'Download large size' (118.92 MB) and 'Download medium size' (69.15 MB), both labeled as version 3.2.0. At the bottom of the page, there are links for 'About | Issues | Version History »'.

Tutorial and online resources – GitHub project

2. Get occurrence data and species range

```
# Download manually tree occurrences and extent (shapefile) and paste it into the dat/ folder
browseURL("https://figshare.com/collections/A_high-resolution_pan-European_tree_occurrence_dataset/3288407/1",
           browser=getOption("browser"), encodeIfNeeded=F)

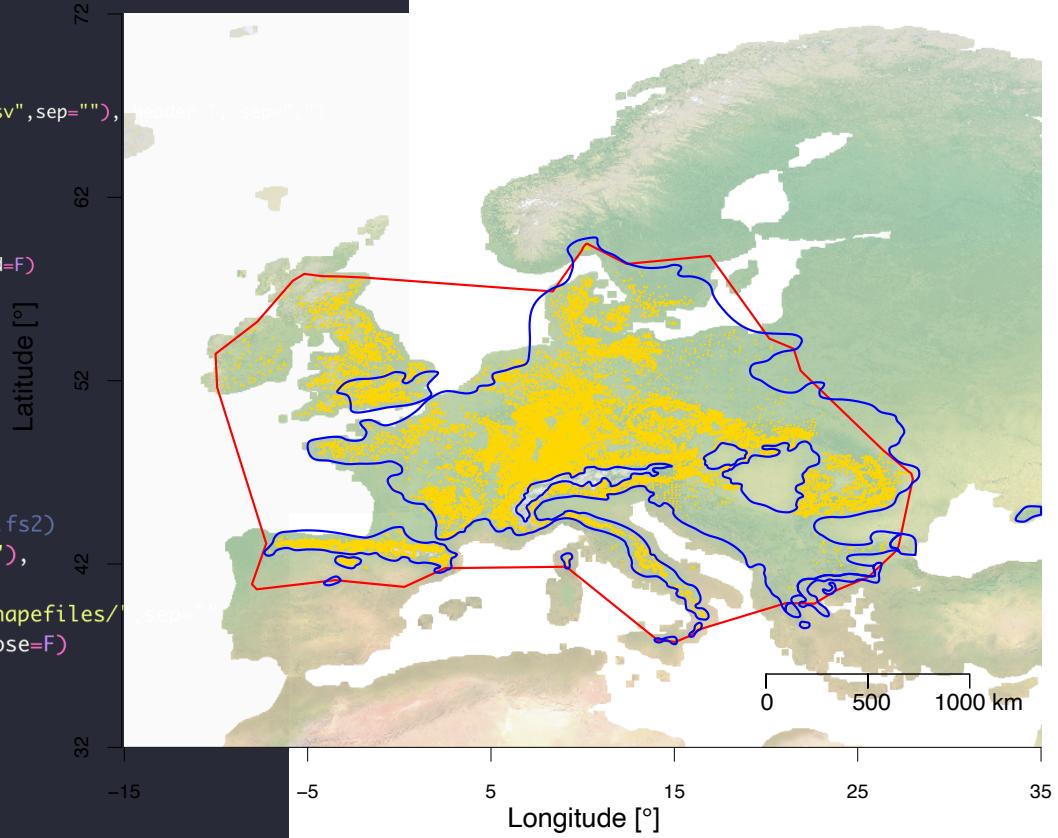
# Download manually tree species range (shapefile) and paste it into the dat/ folder
browseURL("https://www.euforegen.org/species/fagus-sylvatica/",
           browser=getOption("browser"), encodeIfNeeded=F)

# Load tree occurrence data
occ <- read.table(paste(path2, "EUForestTreeOccurrences/EUForestsspecies.csv", sep=""),
                  occ.fs <- data.frame(occ[occ$SPECIES.NAME=="Fagus sylvatica",])
head(occ.fs); dim(occ.fs)
plot(occ.fs$x, occ.fs$y, cex=0.2)

# Transform coordinates from ETRS89 to WGS84
browseURL("https://epsg.io", browser=getOption("browser"), encodeIfNeeded=F)
d <- data.frame(lon=occ.fs$x, lat=occ.fs$y)
coordinates(d) <- c("lon", "lat")
proj4string(d) <- CRS("+init=epsg:3035") # ETRS89 / ETRS-LAEA
CRS.new <- CRS("+init=epsg:4326") # WGS84
d.wgs84 <- spTransform(d, CRS.new)
str(d.wgs84)
cd <- data.frame(d.wgs84@coords)
occ.fs$lonwgs84 <- cd$lon
occ.fs$latwgs84 <- cd$lat

# Load the species' occurrence (rg.fs1) and natural range (rg.fs2)
rg.fs1 <- readOGR(dsn= paste(path2, "EUForestTreeRanges/", sep=""),
                   layer="Fagus_sylvatica", verbose=F)
rg.fs2 <- readOGR(dsn= paste(path2, "FsylSpeciesRangeEuforegen/shapefiles/",
                   layer="Fagus_sylvatica_sylvatica_plg", verbose=F)
crs(rg.fs1) <- CRS('+init=EPSG:4326')
crs(rg.fs2) <- CRS('+init=EPSG:4326')
rg.fs1 <- raster::crop(rg.fs1, eu.extent)
rg.fs2 <- raster::crop(rg.fs2, eu.extent)

# Visualise occurrence data and species' natural range
pdf(paste(path1, "FS_OccurrencesAndRange.pdf", sep=""), width=7.5, height=6.15)
par(mfrow=c(1,1), oma=c(4, 4, 0.5, 0.5), mai=c(0, 0, 0, 0), mar=c(0, 0, 0, 0))
```



Tutorial and online resources – GitHub project

3. Download environmental data

```
# Climate -- download rasters from CHELSA
browseURL("https://chelsa-climate.org/downloads/",
           browser=getOption("browser"), encodeIfNeeded=F)
getOption("timeout") # check the time out setting
options(timeout=1000) # set a longer time if needed
for(i in 1:length(bio.nb1)){# download climatic layers

download.file("https://chelsa-climate.org/wp-admin/download-page/CHELSA_tech_specification_V2.pdf",
              paste(path3,"CHELSA/CHELSA_tech_specification_V2.pdf",sep="")) # download the documentation too

# Topography -- download raster from GMTED2010
browseURL("https://topotools.cr.usgs.gov/gmted_viewer/gmted2010_global_grids.php",
           browser=getOption("browser"), encodeIfNeeded=F)
download.file("https://edcintl.cr.usgs.gov/downloads/sciweb1/shared/topo/downloads/GMTED/Grid_ZipFiles/mn30_grd.zip",
              paste(path2,"mn30_grd.zip",sep=""))
unzip(zipfile=paste(path2,"mn30_grd.zip",sep=""), exdir=paste(path2,"mn30_grd",sep=""))

# Soil -- download rasters from ISRIC
browseURL("https://data.isric.org/geonetwork/srv/eng/catalog.search#/metadata/2fdf1a33-487f-4aa7-a6b6-b2b49483ad48",
           browser=getOption("browser"), encodeIfNeeded=F)
download.file("https://files.isric.org/soilgrids/latest/data_aggregated/1000m/phh2o/phh2o_5-15cm_mean_1000.tif",
              paste(path2,"SoilGrid/phh2o_5-15cm_mean_1000.tif",sep=""))
download.file("https://files.isric.org/soilgrids/latest/data_aggregated/1000m/phh2o/phh2o_60-100cm_mean_1000.tif",
              paste(path2,"SoilGrid/phh2o_60-100cm_mean_1000.tif",sep=""))
```

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4. Extract environmental data from occurrences

```
##### Extract environmental data from occurrences #####
head(occ.fs); dim(occ.fs)
env.fs <- occ.fs
for(i in 1:length(bio.nb2)){}

gmted <- raster::raster(paste(path2, "mn30_grd/mn30_grd/w001000.adf", sep=""))
env.fs$gmted <- raster::extract(gmted, env.fs[, c("lonwgs84","latwgs84")], "bilinear")

pH.top <- raster::raster(paste(path2, "SoilGrid/phh2o_5-15cm_mean_1000.tif", sep="")) # ISRIC layers
pH.top <- projectRaster(pH.top, crs=CRS("+init=EPSG:4326"))
raster::plot(pH.top)
env.fs$pH.top <- raster::extract(pH.top, env.fs[, c("lonwgs84","latwgs84")], "bilinear") / 10
pH.dee <- raster::raster(paste(path2, "SoilGrid/phh2o_60-100cm_mean_1000.tif", sep="")) # ISRIC layers
pH.dee <- projectRaster(pH.dee, crs=CRS("+init=EPSG:4326"))
raster::plot(pH.dee)
env.fs$pH.dee <- raster::extract(pH.dee, env.fs[, c("lonwgs84","latwgs84")], "bilinear") / 10

head(env.fs)
env.fs <- env.fs[,-c(3:10)]
write.table(env.fs, paste(path1, "FS_OccurrencesEnvDatExtracted.csv", sep=""),
            row.names=F, col.names=T, quote=F, sep=";") # save table with extracted environmental data
```

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5. Crop layers for EU extent

6. Visualise environmental data / explore ecological gradients

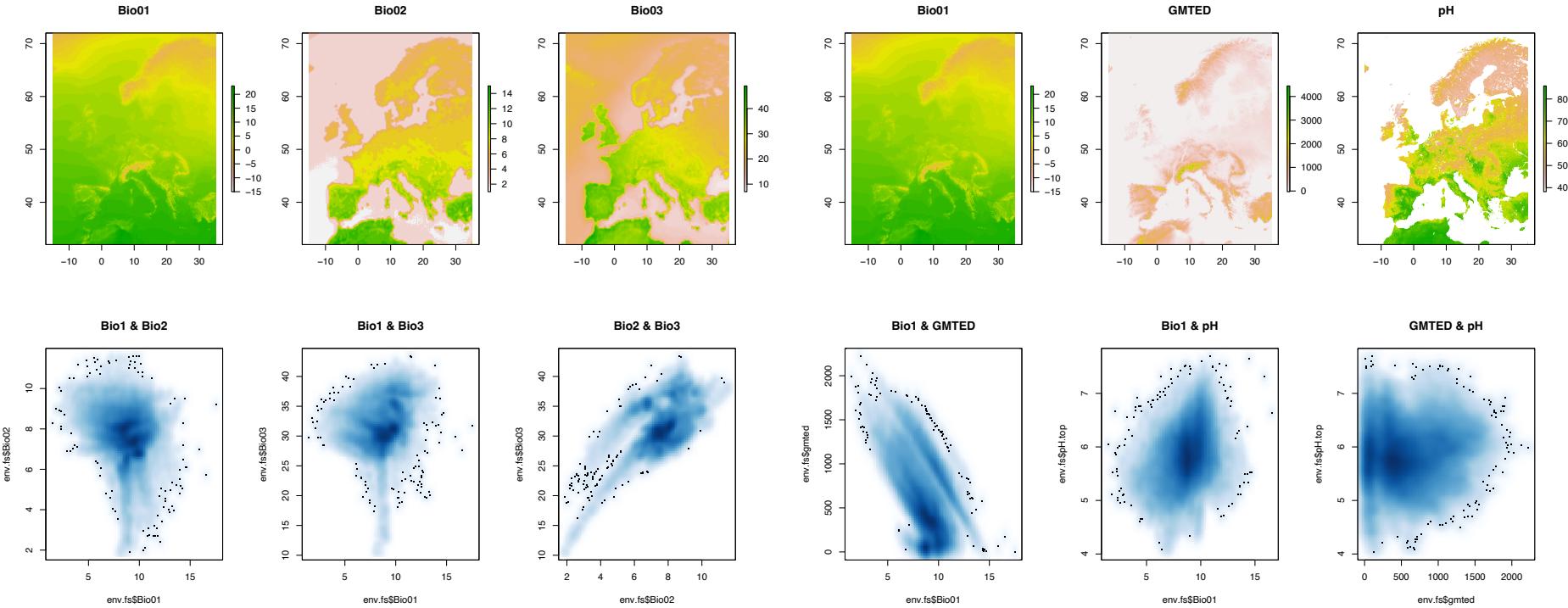
```
#### Crop layers for EU extent ####
bio01.c <- crop(bio01, eu.extent)
bio02.c <- crop(bio02, eu.extent)
bio03.c <- crop(bio03, eu.extent)
gmted.c <- crop(gmted, eu.extent)
pH.top.c <- crop(pH.top, eu.extent)
pH.dee.c <- crop(pH.dee, eu.extent)

#### Visualise environmental data and explore ecological gradients ####
pdf(paste(path1,"FS_OccurrencesEnvDatExtracted1.pdf",sep=""), width=7.5*1.2, height=6.15*1.2)
par(mfrow=c(2,3))
plot(bio01.c, main=paste("Bio",bio.nb2[1],sep=""), alpha=150)
plot(bio02.c, main=paste("Bio",bio.nb2[2],sep=""), alpha=150)
plot(bio03.c, main=paste("Bio",bio.nb2[3],sep=""), alpha=150)
smoothScatter(env.fs$Bio01, env.fs$Bio02, main="Bio1 & Bio2")
smoothScatter(env.fs$Bio01, env.fs$Bio03, main="Bio1 & Bio3")
smoothScatter(env.fs$Bio02, env.fs$Bio03, main="Bio2 & Bio3")
dev.off()

pdf(paste(path1,"FS_OccurrencesEnvDatExtracted2.pdf",sep=""), width=7.5*1.2, height=6.15*1.2)
par(mfrow=c(2,3))
plot(bio01.c, main=paste("Bio",bio.nb2[1],sep=""), alpha=150)
plot(gmted.c, main=paste("GMTED",sep=""), alpha=150)
plot(pH.top.c, main=paste("pH",sep=""), alpha=150)
smoothScatter(env.fs$Bio01, env.fs$gmted, main="Bio1 & GMTED")
smoothScatter(env.fs$Bio01, env.fs$pH.top, main="Bio1 & pH")
smoothScatter(env.fs$gmted, env.fs$pH.top, main="GMTED & pH")
dev.off()
```

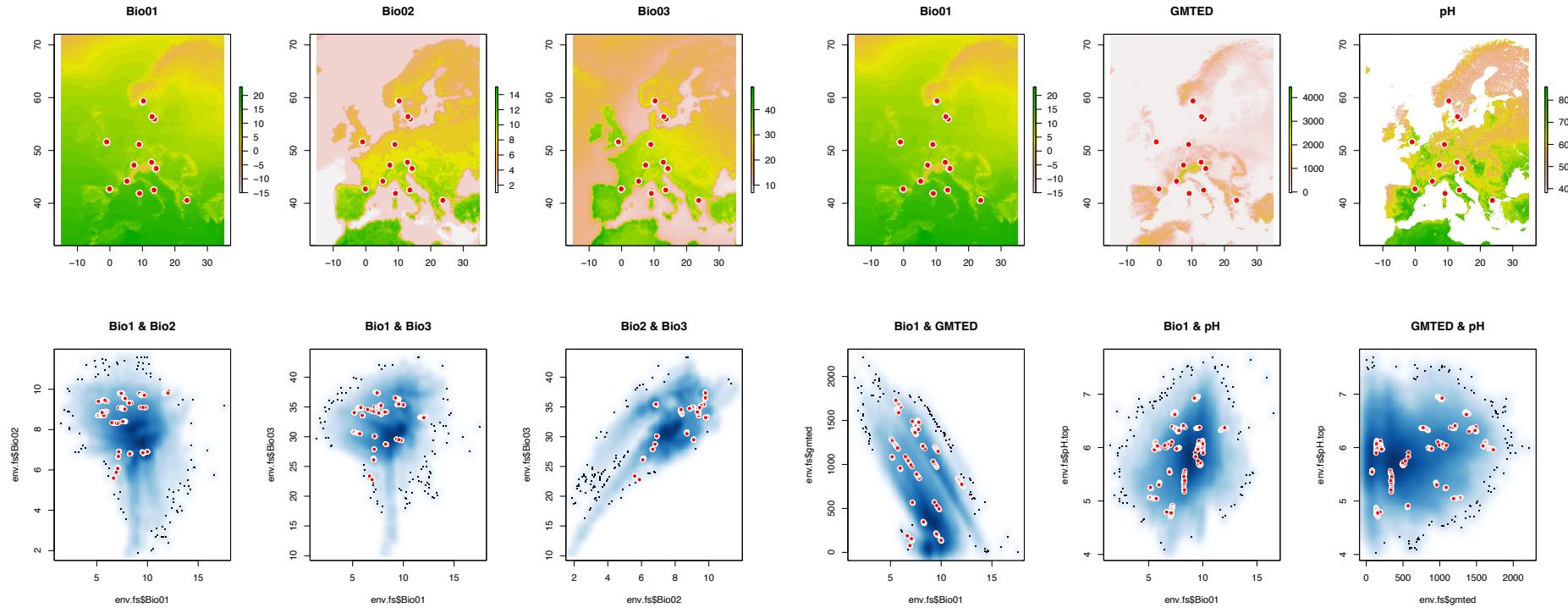
Tutorial and online resources – GitHub project

6. Visualise environmental data / explore ecological gradients



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7. Sampling locations, data extraction and visualisation



Outline

I. General introduction

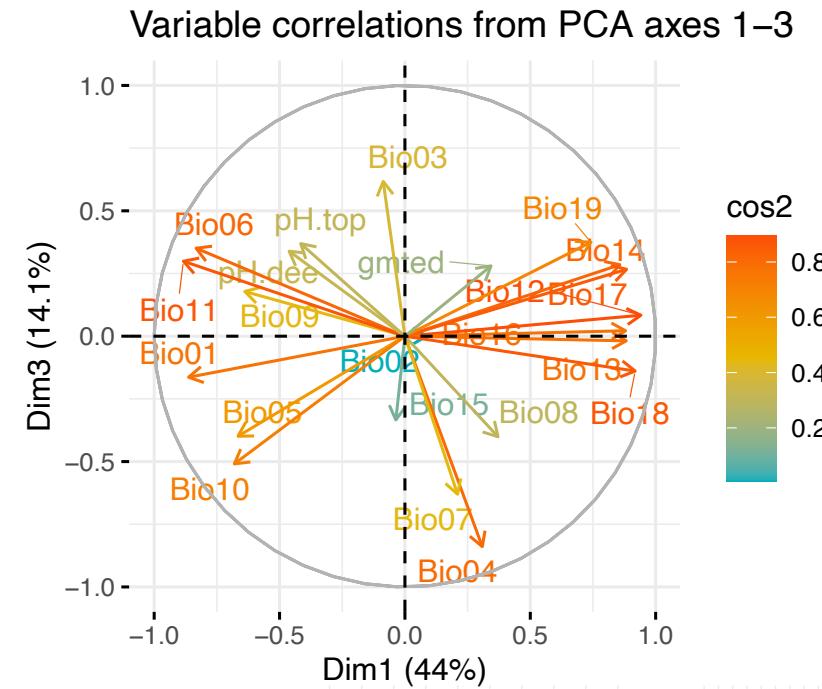
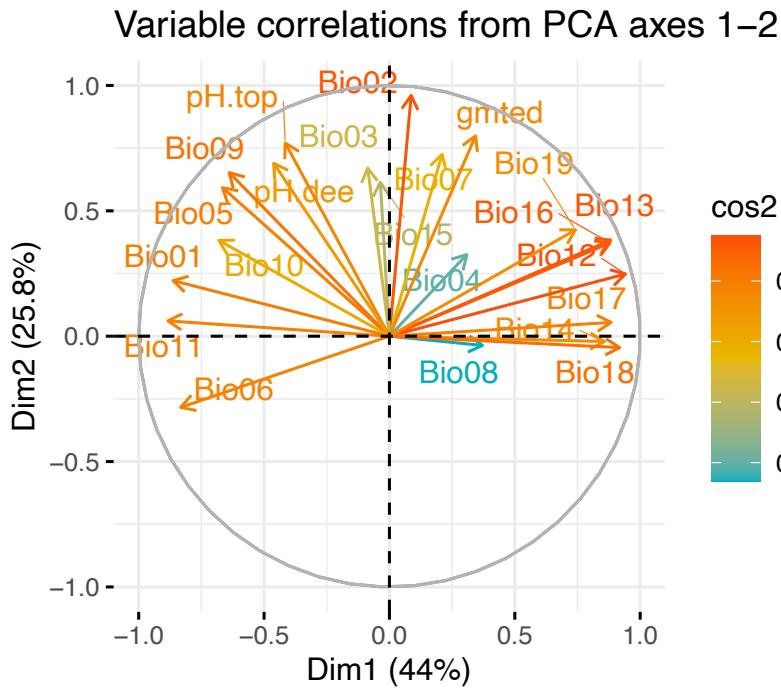
II. A brief portray of environmental data

III. A short tutorial on how to handle environmental data in R

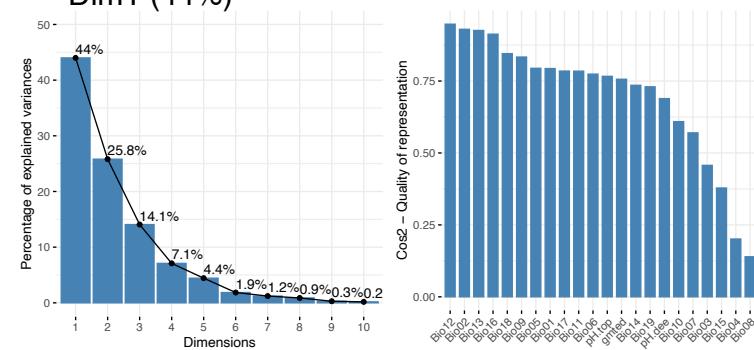
IV. Reduction of environmental variables

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8. PCA of environmental data at the sampling locations



Use principal components as predictors



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9. Correlation plot of environmental data

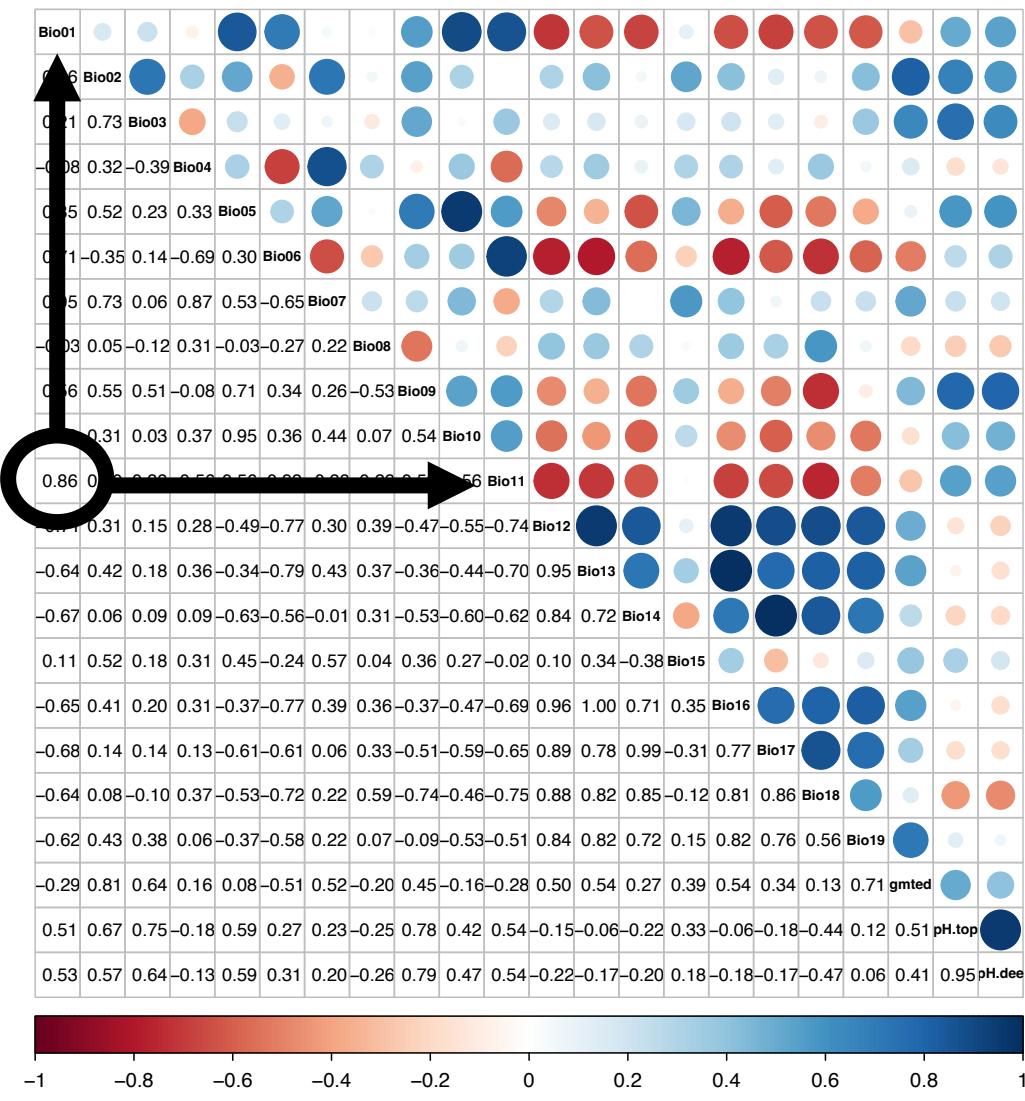
10. Pearson correlation matrix of

environmental data

9. Retain uncorrelated variables

based on Pearson's coefficient

	Bio01	Bio04	Bio08	Bio12	Bio15	gmted	pH.top
Bio01		-0.077	-0.025	-0.711	0.106	-0.291	0.507
Bio04	-0.077		0.308	0.276	0.309	0.160	-0.178
Bio08	-0.025	0.308		0.394	0.037	-0.199	-0.246
Bio12	-0.711	0.276	0.394		0.104	0.496	-0.149
Bio15	0.106	0.309	0.037	0.104		0.389	0.330
gmted	-0.291	0.160	-0.199	0.496	0.389		0.507
pH.top	0.507	-0.178	-0.246	-0.149	0.330	0.507	



Take home message

- Question which environmental data are relevant for your study; measured, interpolated, biotic, abiotic...
- Explore environmental gradients from occurrence data before starting sampling
- Be careful with spatial resolution \neq accuracy
- An environment variable depends on spatial and temporal scale
- Define the reference period fitting the selective pressures
- Always inspect variable interdependence
- Variable selection is also a way to improve predictive power of env. variables

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One and a half year post-doctoral research position on the landscape genomics of forest trees and their ectomycorrhizal fungi





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Ecological Genetics

Head: **Dr. Martina Peter**



Identifying seed sources for highly adaptable oak forests in a changing climate

The international BiodivERsA project ACORN aims to identify oak populations capable to cope with future drought-stress due to climate change.



OAKID2

Transferability and phylogenetic extension of species-diagnostic SNP markers in European white oak species



GenWood

We link genomic and dendro-phenotypic data in Norway Spruce (*Picea abies*) to study the genomic basis of wood anatomy and assess the adaptive potential of forest trees under climate change.



BLACKDYNAMITE

This Swiss-Spanish project investigates the dynamics and competition of the black Périgord and Burgundy truffles under the influence of climate change.



Root endophytic fungi and their contribution to arctic and alpine climate change feedback

Arctic and Alpine vegetation dynamics under climate change can exert a lasting impact on belowground processes and ultimately on world's largest carbon stocks. This long-term field manipulation investigates the functional significance of plant interactions with species-specific endophytic fungi.



Adaptation of trees to extreme climate events

Extreme climate events are expected to become more frequent with climate warming. We test genetic differentiation in resistance to late frost and drought in silver fir populations planted in the 1980s in Switzerland and abroad.



Tool kit conservation genetics

Partners from the private and academic sector joined to set up a ready-to-use tool kit, which allows authorities and consultancies to apply conservation genetic methods in a standardized way in their everyday work.



Metalink

Soil contamination with heavy metals is a widespread problem affecting natural, cultivated, urban and industrial sites throughout Europe. The main objective in this Swiss/Polish collaboration project is to characterize the links between environmental, genotypic, and phenotypic features in the metal-tolerant plant model *Arabidopsis halleri*.



GenTree

GenTree aims at providing the European forestry sector with better knowledge, methods and tools for optimizing the management and sustainable use of forest genetic resources in Europe in the context of climate change and continuously evolving demands for forest products and services.



PiCadapt - Adaptive genetic variation of Swiss stone pine in response to environmental gradients across the Alpine timberline ecotone

PiCadapt aims to study gene-encoded nucleotide variation in the bird-dispersed *P. cembra*, a keystone tree species of the Alpine timberline ecotone, in response to environmental gradients.



GeneScale

GeneScale describes genomic variation in the Alpine rock-cress (*Arabis alpina*) and seeks indications of local adaptation to small-scale environmental conditions.