# Step 1.1 Creating field-based plots

# 1.1.1. Creating plots based on original samples

### Diane ESPEL

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# 1 Objectives

This R script aims to generate square quadrat plots centered on ecological sampling points collected in the field from both recent and historical sources. The workflow includes:

- Loading reference field sample files for different sources (recent "FIELD" and historical "HFI").
- Projecting geographic coordinates to a local metric system (EPSG:32739)
- Computing square buffer areas around the points
- Exporting both sampling centroids and polygon quadrats as shapefiles for further spatial analysis

# 2 Script explanation

## 2.1 Clean environment and graphics

```
rm(list = ls()) # Clear all objects from the R environment to start fresh
graphics.off() # Close all graphics devices (if any plots are open)
```

## 2.2 Load required packages

```
library(sp) # For SpatialPolygons and Spatial data structures.
library(sf) # For modern handling of spatial vector data.
library(dplyr) # For data manipulation (filtering, joining, etc.).
library(data.table) # For fast file reading and merging.
```

## 2.3 Define Global Variables

Note: A global variable is a variable defined outside of any function. This means the variable is accessible from any part of the code, including inside functions. A global variable retains its value throughout the execution of the R script unless it is explicitly modified in the code.

It is important to define at a minimum:

- the "District": the archipelago of interest (e.g. "CRO" for Crozet archipelago)
- the "Island": the island within the archipelago of interest (e.g. "POS" for Possession island)

```
District='CRO' # 3-letter code for the archipelago
Island='POS' # 3-letter code for the island
```

## 2.4 Set working directory

You must define a general root directory ("localHOME") that serves as the base path for your input and output data. This directory should point to the local environment where:

- input sample files are located under "/data/samples"
- outputs (e.g., plots or shapefiles) will be saved under "/data/vector/Plots/PrimaryTypo"

You can create additional subfolders within this structure to organize your outputs by data type, source, or year of analysis.

```
# Define local root directory
#localHOME = paste0("your_local_path/")
localHOME=paste0("/home/genouest/cnrs_umr6553/despel/CARTOVEGE/")

# Define path to load input sample data
open_samples_path=paste0(localHOME, "data/samples")

# Define path to save generated outputs (plots)
save_plots_path=paste0(localHOME, "data/vector/Plots/PrimaryTypo")
```

# 2.5 Create field reference samples plots

The script processes two sources: "FIELD" (recent field samples) and "HFI" (historical field samples) in 7 main steps:

### 1) Load and merge Sample Files

- List files corresponding to each source.
- Load and concatenate the sample datasets.
- Remove duplicates and assign source information.

## 2) Apply Coordinate Projection

- Convert longitude and latitude from EPSG:4326 to EPSG:32739 (UTM zone).
- Add new projected coordinates (X and Y in meters).

### 3) Prepare Final Table

• Select and rename key columns: observation ID, coordinates, surface, habitat levels, etc.

### 4) Compute Quadrat Corner Coordinates

- Define quadrat size (e.g., 4 meters side).
- Compute coordinates for corners of each square (N, S, E, W).
- Ensure no duplicated observation IDs.

## 5) Save Sampling Centroids

• Export updated sampling point data (with projected coordinates) to CSV.

## 6) Draw and Save Quadrats

- Generate square polygons clockwise from NW corner.
- Build SpatialPolygons assigning each an ID and then a SpatialPolygonsDataFrame.
- Join back sample attributes to each polygon from the original observations (coordinates, date, source, habitat levels, etc.).
- Convert the quadrats to sf object and export to shapefile.

#### 7) Export Results

• Export quadrat polygons and centroid points as .shp files using st\_write() from the sf package.

```
file_list <- list()</pre>
 for (i in 1:length(all_files)) {
  file_list[[i]] <- read.csv(all_files[i], sep = ";", dec = ".", stringsAsFactors = FALSE) # Read

→ each file and store in list

  assign(paste0(source,i), read.csv(all_files[i], sep = ";", dec = ".", stringsAsFactors =
   \hookrightarrow FALSE)) # Assign file to dynamic variable
 Nb_database = length(all_files) # Count number of files imported
 print(paste0("Number of imported databases: ", Nb_database))
 # Merge all dataframes into a single one (even if only one input file)
 df = rbindlist(file_list)
 # Remove potential duplicates
 New_df = unique(df)
 New_df = as.data.frame(New_df)
 # Add source identifier
 New df$Source = source
 print(paste0("Number of unique observations: ", length(unique(New_df$N_observation))))
 # Convert XY coordinates columns -----
 filt_points <- New_df</pre>
 # Convert coordinates from EPSG 4326 (decimal degrees) to EPSG 32739 (UTM Zone 39S)
 print("Columns Longitude_ddd and Latitude_ddd are in EPSG 4326 (decimal degrees) and must be
 ⇔ converted to EPSG 32739 (meters)")
 filt_points <- st_as_sf(filt_points, coords = c("Longitude_ddd", "Latitude_ddd"), crs = 4326) %>%
   st_transform(32739)
 # Extract converted coordinates (X and Y in meters)
 filt_points_m <- st_coordinates(filt_points)</pre>
 # Re-add longitude and latitude columns
 filt_points$xcoord_m <- filt_points_m[, "X"]</pre>
 filt_points$ycoord_m <- filt_points_m[, "Y"]</pre>
 filt_points$Longitude_ddd <- New_df[, "Longitude_ddd"]</pre>
 filt_points$Latitude_ddd <- New_df[, "Latitude_ddd"]</pre>
 # Select columns of interest
final_points <- filt_points[, c("N_observation", "Date_manip", "New_protocole", "Source",</pre>
# Rename columns for clarity
 colnames(final_points) <- c("N_obs", "Date", "Protocole", "Source", "Longitude",</pre>
 → "Latitude", "xcoord_m", "ycoord_m", "Surface", "Longueur_m", "Largeur_m", "Hab_L1", "Hab_L2",
 Hab_L3", "Hab_L4", "geometry")
 # Create spatial polygons ------
 # Compute corners of quadrat based on center (coordinates) and desired size
 print("Defining quadrat side length (m) and calculating corner coordinates")
 length_quadrat <- 4</pre>
 radius <- length_quadrat / 2
 poly_points <- data.frame(final_points)</pre>
```

```
poly_points$yPlus <- poly_points$ycoord_m + radius # North</pre>
 poly_points$xPlus <- poly_points$xcoord_m + radius # East</pre>
 poly_points$yMinus <- poly_points$ycoord_m - radius # South</pre>
 poly_points$xMinus <- poly_points$xcoord_m - radius # West</pre>
 print("Check for duplicate observation IDs (should be unique)")
 indices_doublons <- which(duplicated(poly_points$N_obs) | duplicated(poly_points$N_obs, fromLast
\rightarrow = TRUE))
 print(indices_doublons)
 # Construct quadrat polygons (5 points: NW to NW clockwise to close the shape)
 quadrats <- cbind(poly_points$xMinus, poly_points$yPlus, # NW
                   poly_points$xPlus, poly_points$yPlus,
                   poly_points$xPlus, poly_points$yMinus, # SE
                   poly_points$xMinus, poly_points$yMinus, # SW
                   poly_points$xMinus, poly_points$yPlus) # back to NW
 # Create spatial polygons from coordinates
 ID <- poly_points$N_obs # Extract observation IDs</pre>
 print("Creating spatial polygons (quadrat plots)")
 polysHabitat <- SpatialPolygons(</pre>
   mapply(function(poly, id) {
     xy <- matrix(poly, ncol=2, byrow=TRUE)</pre>
                                                  # Create coordinate matrix
     Polygons(list(Polygon(xy)), ID = id)
                                                # Create Polygon object with ID
   },
   split(quadrats, row(quadrats)), ID),
                                                # Split coordinates by row (one polygon per row)
   proj4string = CRS("+init=EPSG:32739")
                                                # Define projection
 plot(polysHabitat) # Plot polygons for visual check
 # Create SpatialPolygonsDataFrame to store attributes
 # This object links each quadrat polygon with its corresponding metadata (attributes)
 polys.df <- SpatialPolygonsDataFrame(polysHabitat, # The SpatialPolygons object containing
\rightarrow geometry
                                       data.frame(N obs = ID, row.names = ID) # Create a minimal
                                       → data.frame with IDs as row names
polys.df\$xcoord_m = poly_points\$xcoord_m # Add additional attribute fields to the
\hookrightarrow SpatialPolygonsDataFrame
polys.df$ycoord_m = poly_points$ycoord_m
 polys.df$Longitude = poly_points$Longitude
 polys.df$Latitude = poly_points$Latitude
 polys.df$Date = poly_points$Date
 polys.df$Protocole = poly_points$Protocole
 polys.df$Source = poly_points$Source
 polys.df$Surface = poly_points$Surface
 polys.df$Longueur_m = poly_points$Longueur_m
 polys.df$Largeur_m = poly_points$Largeur_m
 polys.df$Hab_L1 = poly_points$Hab_L1
 polys.df$Hab_L2 = poly_points$Hab_L2
 polys.df$Hab_L3 = poly_points$Hab_L3
 polys.df$Hab_L4 = poly_points$Hab_L4
 # Save the final shapefiles ---
 # Export polygons as shapefile
```

This script enables the spatial representation of ecological sampling points and prepares geospatial layers suitable for GIS analysis or habitat modeling.