Practical presence-only data

Biodiversity modelling

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Introduction

This practical document presents R code for the section of the course regarding presence-only data.

Different examples will be given using the course data for each of the approach discuss during the course.

In the present document, we will focus on the distribution of Passerella iliaca.

Load R packages

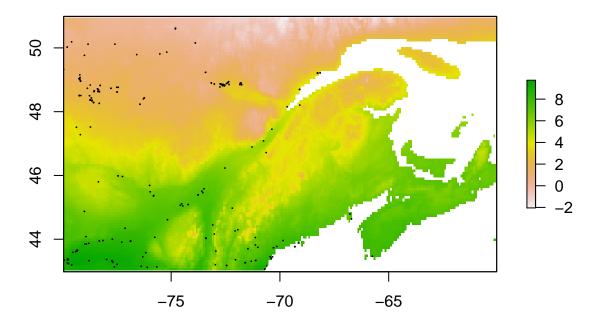
```
library(sp)
library(raster)
```

Load the data

```
bird <- readRDS("bird5.RDS")
climatePresent <- readRDS("climate_Present.RDS")
road <- readRDS("road_Distance.RDS")</pre>
```

Let's look at data we have for Passerella iliaca

```
sp <- bird$Oporornis.agilis
plot(climatePresent[[1]])
points(sp, pch = 19, cex = 0.1)</pre>
```



Survey region

library(maptools)

For point process models, it is important to gather some information about the survey region. Specifically, knowing the area of the survey area is important to know

Focus only on the occurences that fall into the survey regions

```
spRegion <- intersect(sp, landPoly)</pre>
```

Define quadrature points

```
# Define random samples
set.seed(12)
xSmpl <- runif(20000, xmin(landPoly), xmax(landPoly))</pre>
```

Extracting the climate data for the region of interest

This involves extracting the data for the presence points but also for the quadrature points.

Using weighted GLMs

Downweighted Poisson regression

```
# Build response variable
spQuad <- c(rep(1, length(spRegion)), rep(0, nquad))

# Build weight
weight <- rep(1/nquad, length(spQuad))
weight[spQuad == 0] <- areaRegion / nquad

# Point process model
sel <- sample(length(spQuad), 1000)

spModel <- glm(spQuad/weight ~.,</pre>
```

```
data = climateSpQuad,
family = poisson(),
weights = weight)
```

Are there enough quadrature points?

Redefine quadrature points

Extracting the climate data for the region of interest

This involves extracting the data for the presence points but also for the quadrature points.

Recalculate the downweighted Poisson regression

Check if enough quadrature points were used

[1] 1.36245

Perfect!

Draw a map of the model

```
climateDat <- as.data.frame(scale(values(climatePresent)))</pre>
# Calculate the estimated intensity for the survey region
intensityMap <- predict(spModel,</pre>
                          newdata = climateDat,
                          type = "response")
# Build the raster object
pred <- raster(climatePresent)</pre>
values(pred) <- intensityMap</pre>
# Draw the map
plot(pred)
50
                                                                           0.008
                                                                           0.006
                                                                           0.004
46
                                                                           0.002
4
                                  -70
                  -75
                                                 -65
```

Because this model assumes all occurrence points are independent, let's take this into account.

Warning

The code in the following section is very (!) long to run. It was included for you to know how to do these types of models in R, I recommend not to run it during the course.

Area-interaction model

To estimate an area-interaction model we need to use the spatstat R package.

library(spatstat)

To fit an area-interaction model, we first need to define the best radius to use. For illustration purposes, let's assume a radius of $10~\rm km$

```
# Build owin object
landWin <-as.owin(landPoly)</pre>
# Define point process object
spPPP <- ppp(coordinates(sp)[,1],</pre>
              coordinates(sp)[,2],
              window = landWin,
              check = FALSE)
# Define quadrature point object
quadPoint <- ppp(quadSel[,1], quadSel[,2],window = landWin)</pre>
Q <- quadscheme(data = spPPP, dummy = quadPoint, method = "grid")
# Formula
formu <- as.formula(paste0("~", paste0("bio",1:19, collapse = "+")))</pre>
# Format climate
climateList <- vector("list", length = nlayers(climatePresent))</pre>
for(i in 1:nlayers(climatePresent)){
  climateList[[i]] <- as.im(climatePresent[[i]])</pre>
}
names(climateList) <- paste0("bio", 1:19)</pre>
AImodel <- ppm(Q,
                trend = formu,
                covariates = climateList,
                interaction = AreaInter(0.1))
```