

Simulating scenarios of feature distribution

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Introduction

There are a few big questions that underlie the assessment of the effects of anthropogenic infrastructure on wildlife. When performing environmental impact assessments, one aims not only to find which factors affect wildlife and how strongly, but also (i) at which spatial scale there are effects and (ii) how these effects sum and interact when (as it is often the case) there are multiple infrastructures and vectors of landscape modification. The first question is also known as the Zone of Influence (ZoI) problem. The second is generally tackled in the context of cumulative impact assessment.

In the main text of this manuscript, we argue that the density of infrastructure features might better represent the cumulative effect of multiple features in the landscape than considering only the distance to the feature nearest to a given location. This, however, might vary depending on how the number of features in the landscape, how these features are distributed in space, and what is the ZoI of each of those features.

Here we simulate some landscapes with point-type infrastructure spread following different patterns, calculate the distance to the nearest feature and the density of features, at multiple scales, and assess when and how these variables might represent different sources of spatial variation.

Simulate landscapes

First we simulate some landscape using point-type infrastructure as an example. They could represent the spatial location of houses, cabins, or wind turbines, for example. To do this we'll use a few functions designed within the package `oneimpact`.

```
# Load packages
library(dplyr)
library(purrr)
library(ggplot2)

library(mobsim)
library(raster)
library(landscapetools)
library(sf)
library(spatstat)

library(oneimpact)

library(reshape2)
library(lsr)
```

We set a 30x30 km² landscape and can create simulate points following different spatial patterns. Here is an example landscape where the points are spread close to a single center (e.g. houses in a village).

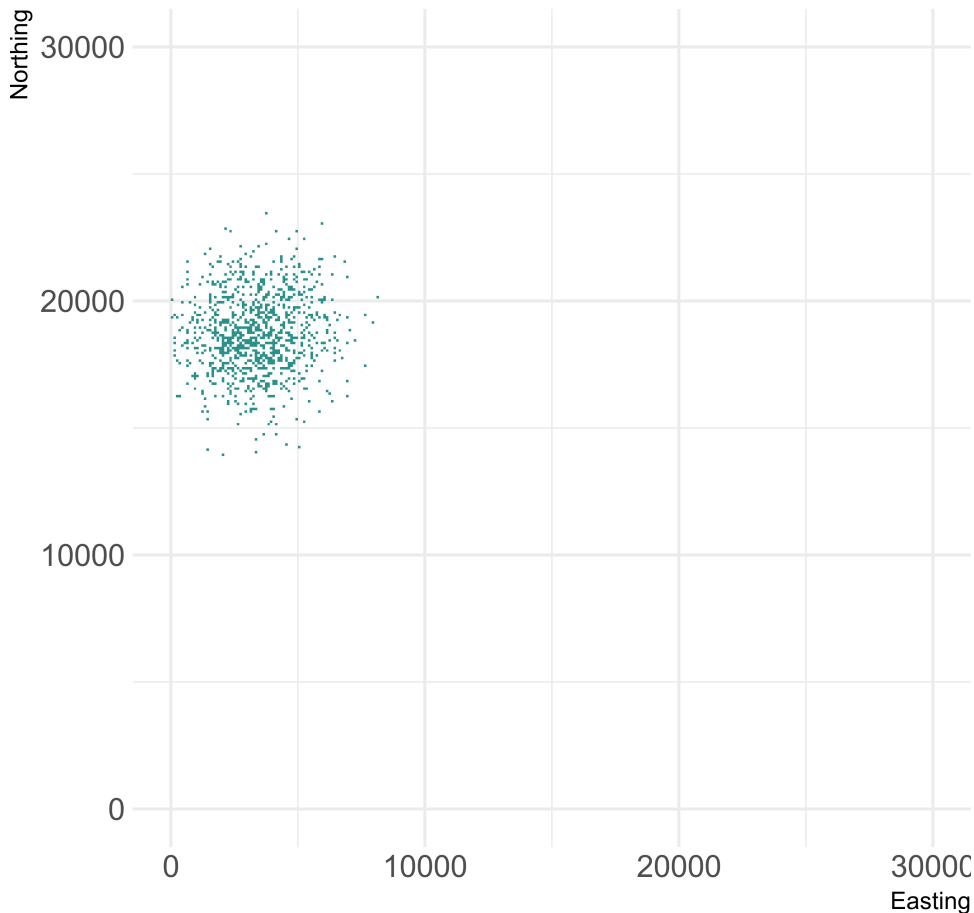
```

set.seed(1234)

# simulate a single patch
nfeat <- 1000 # number of features
ext <- 30000 # extension of the landscape
nc <- 1 # number of centers or patches
wd <- ext / 20 # width of the patch
pts <- set_points(
  n_features = 1000, centers = nc,
  width = wd, res = 100,
  extent_x = c(0, ext), extent_y = c(0, ext)
)

landscapetools::show_landscape(pts$rast, legend.position = "none")

```



We can now simulate landscapes with different patterns. We start with 3 scenarios: regular, random, and clumped distribution of points. Each scenario is simulated with 10, 100, and 1000 points, so that we have 12 scenarios in total.

```

set.seed(123)

# random, gradient, single center, multiple centers
methods <- c("regular", "random", "mobsim")
name <- c("regular", "random", "clumped5", "clumped1")
nfeat <- c(10, 100, 1000) # number of features
res <- 100 # resolution

```

```

ext <- 30000 # extent of the landscape
nc <- c(5, 1) # number of centers for clumped
wd <- c(0.05) * ext # width of the "patches"

# parameters
parms_df1 <- expand.grid(
  method = methods, n_features = nfeat,
  centers = nc[1], width = wd
)
parms_df2 <- expand.grid(
  method = methods[3], n_features = nfeat,
  centers = nc[2], width = wd
)
parms_df <- dplyr::bind_rows(parms_df1, parms_df2) %>%
  dplyr::arrange(n_features, method)
scenarios <- paste0(rep(name, 3), "_", rep(nfeat, each = 4))

# simulate points
pts <- parms_df %>% purrr::pmap(set_points,
  res = res,
  extent_x = c(0, ext),
  extent_y = c(0, ext)
)

landscapes <- purrr::map(pts, ~ .[[2]])
names(landscapes) <- scenarios

landscapetools::show_landscape(landscapes)

```

```

## Warning: `guides(<scale> = FALSE)` is deprecated. Please use `guides(<scale> =
## "none")` instead.

```

```

## Warning: Removed 7188 rows containing missing values (geom_raster).

```



```

# plot(landscapes, col = "black", nc = 4, legend = F)
# rasterVis::levelplot(stack(landscapes), layout = c(4,3), names.attr = scenarios,
#   par.settings = GrTheme, colorkey = FALSE)

```

We also store some variables related the distance between points for each of these scenarios. These variables are: (i) the average distance between points, the average nearest neighbor distance, and the average isolation.

```

# isolation to random points
isolation <- function(x, n_rand = 100, ext = c(0, 1), lonlat = FALSE) {
  # create random points
  rand <- data.frame(x = runif(n_rand, ext[1], ext[2]), y = runif(n_rand, ext[1], ext[2]))
  # calc dist
  dists <- pointDistance(x, rand, lonlat = lonlat)
}

```

```

# min dist
apply(dists, 2, min)
}

# mean isolation
mean_isolation <- function(x, n_rand = 100, ext = c(0, 1), lonlat = FALSE) {
  mean(isolation(x, n_rand = n_rand, ext = ext, lonlat = lonlat))
}

# points
pts_coords <- purrr::map(pts, first)
names(pts_coords) <- scenarios

# calculate distances
dist_scenarios <- data.frame(
  scenario = scenarios,
  mean_dist = map_dbl(pts_coords, function(x) mean(dist(x))),
  mean_nndist = map_dbl(pts_coords, function(x) mean(spatstat.geom::nndist(x))),
  mean_isolation = map_dbl(pts_coords, mean_isolation, n_rand = 150, ext = c(0, ext))
)

dist_scenarios

```

	scenario	mean_dist	mean_nndist	mean_isolation
##	regular_10	regular_10 15510.736	9486.8330	3910.7559
##	random_10	random_10 17450.119	5238.7641	5320.3333
##	clumped5_10	clumped5_10 15197.147	1155.7217	8224.4716
##	clumped1_10	clumped1_10 3029.684	1132.5368	11103.3612
##	regular_100	regular_100 15717.795	3000.0000	1182.9837
##	random_100	random_100 15252.246	1597.3085	1585.0777
##	clumped5_100	clumped5_100 9454.507	668.2567	5048.0218
##	clumped1_100	clumped1_100 2454.158	319.6537	13629.1097
##	regular_1000	regular_1000 15590.481	948.6833	371.6643
##	random_1000	random_1000 15765.710	486.1710	479.7201
##	clumped5_1000	clumped5_1000 14179.990	208.8593	4900.2276
##	clumped1_1000	clumped1_1000 2662.354	115.1179	10947.7893