

## Norwegian spruces dataset

```
library(ppjsdm)
#> Registered S3 method overwritten by 'spatstat':
#>   method      from
#> print.boxx cli
library(spatstat)
#> Loading required package: spatstat.data
#> Loading required package: nlme
#> Loading required package: rpart
#>
#> spatstat 1.63-0      (nickname: 'Space camouflage')
#> For an introduction to spatstat, type 'beginner'
remove(list = ls())

set.seed(1)
```

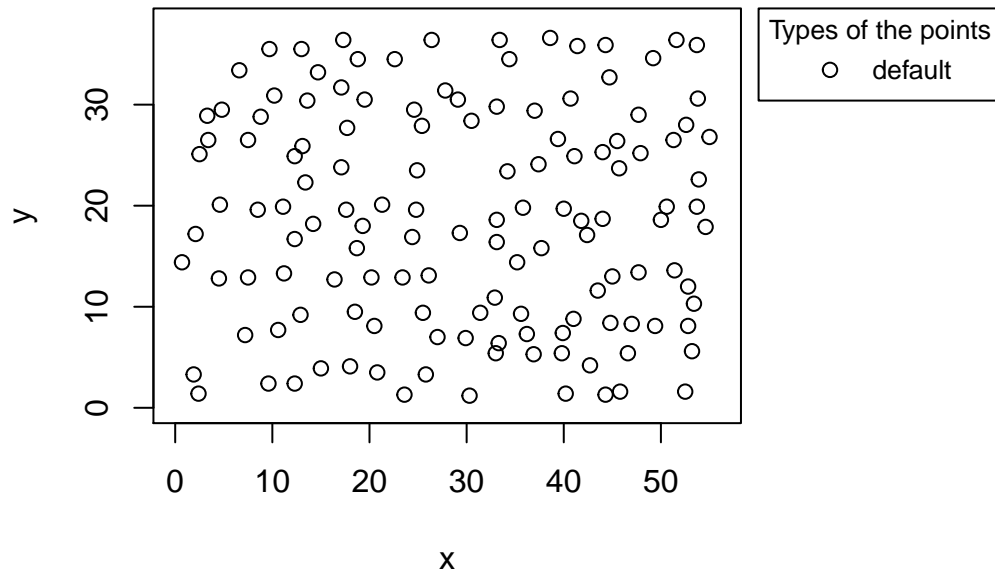
This vignette explains how to use the `ppjsdm` package with the `spruces` dataset from `spatstat`. We begin by loading the data with all species.

```
truncated_spruces <- spruces
# Window(truncated_spruces) <- owin(c(3, 56), c(0, 35))
configuration <- Configuration(truncated_spruces$x, truncated_spruces$y, marks = truncated_spruces$marks)
window <- Rectangle_window(c(0, 56), c(0, 38))
```

The point configuration is plotted below.

```
print(configuration)
#> An S3 object representing a configuration.
#>
#> Number of points: 134.
par(mar = c(5, 4, 4, 13) + 0.1)
plot(configuration, window = window)
```

## Points in the configuration

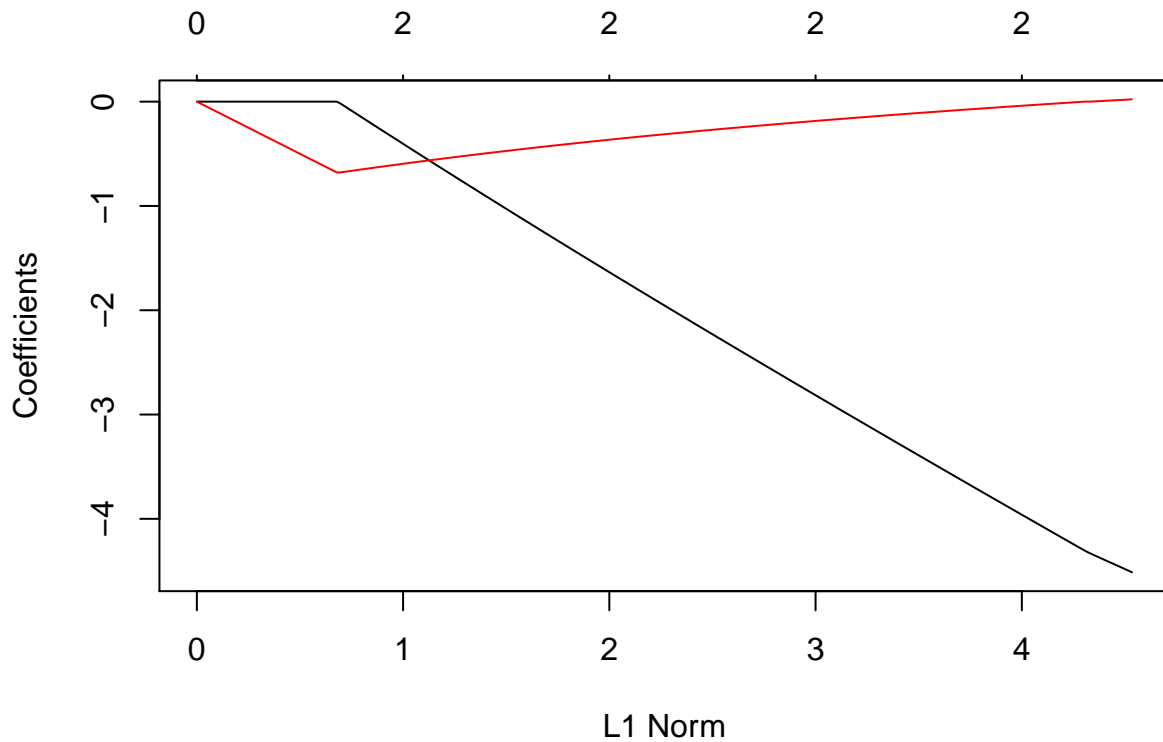


We provide a series of ranges for the interaction radii, and let the fitting function calibrate the model.

```
short_range <- c(0, 20)
medium_range <- c(0, 20)
long_range <- c(0, 20)
model <- "square_exponential"
medium_range_model <- "square_exponential"
```

We can now call the fitting function.

```
fit <- ppjsdm::gibbsm(configuration,
  window = window,
  model = model,
  medium_range_model = medium_range_model,
  short_range = short_range,
  medium_range = medium_range,
  long_range = long_range,
  use_glmnet = TRUE,
  use_aic = TRUE,
  saturation = 2)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.000000 -1.378804 -4.320294 0.000000
plot(fit$complete)
```



```
print(fit$coefficients)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.000000 -1.378804 -4.320294 0.000000
print(fit$best_short)
#> [1,]
#> [1,] 5.531309
print(fit$best_medium)
#> [1,]
#> [1,] 13.14443
print(fit$best_long)
#> [1,]
#> [1,] 21.62295
print(fit$aic)
#> [1] -388.9813
print(fit$bic)
#> [1] -384.48
```

We may then plot the corresponding Papangelou conditional intensity.

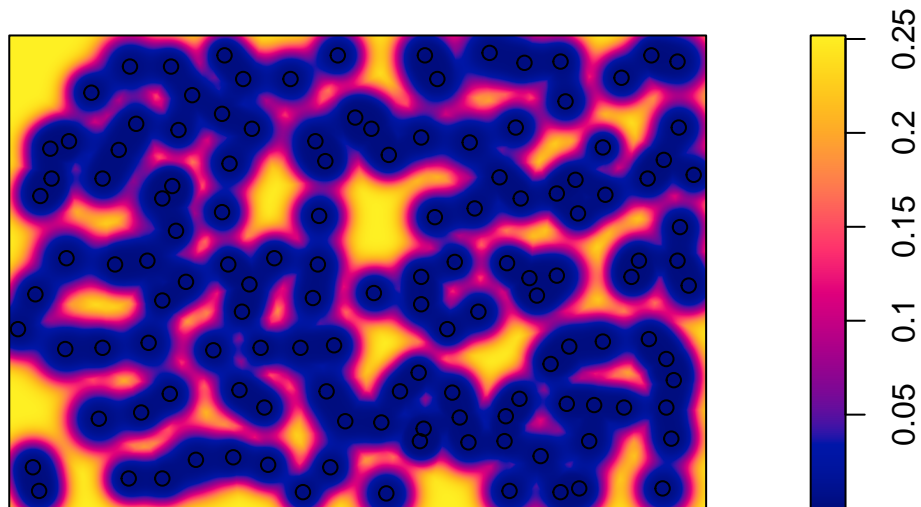
```
parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma
plot_papangelou(window = window,
                 configuration = configuration,
                 type = 1,
                 mark = mean(spruces$marks),
```

```

model = model,
medium_range_model = medium_range_model,
alpha = alpha,
lambda = lambda,
beta = matrix(0, 1, 0),
gamma = gamma,
covariates = list(),
short_range = fit$best_short,
medium_range = fit$best_medium,
long_range = fit$best_long,
saturation = 2)

```

**as.im(t(z), W = window)**



It is also possible to draw from the model.

```

parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma
draw <- ppjsdm::rgibbs(window = window,
                        alpha = alpha,
                        lambda = lambda,
                        gamma = gamma,
                        model = model,
                        medium_range_model = medium_range_model,
                        short_range = fit$best_short,
                        medium_range = fit$best_medium,
                        long_range = fit$best_long,

```

```

      types = levels(types(configuration)),
      mark_range = c(min(get_marks(configuration)), max(get_marks(configuration))),
      steps = 10000000)

print(draw)
#> An S3 object representing a configuration.
#>
#> Number of points: 132.

par(mar = c(5, 4, 4, 13) + 0.1)
plot(draw, window = window)

```

## Points in the configuration

