

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`.

Taking marks into account

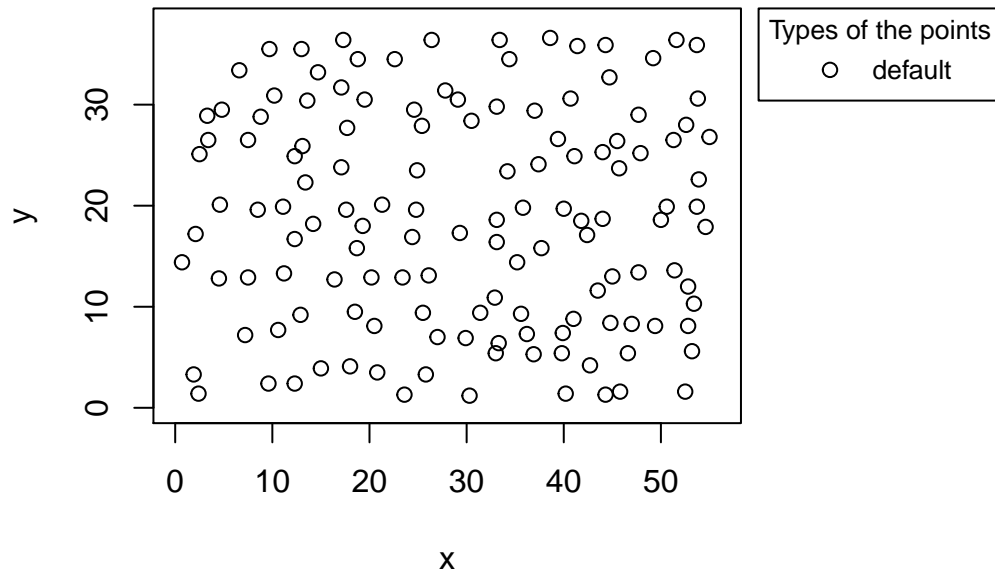
If marks are provided, the interaction radii are proportional to the marks. We begin by that setting.

```
configuration <- Configuration(spruces$x, spruces$y, marks = 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

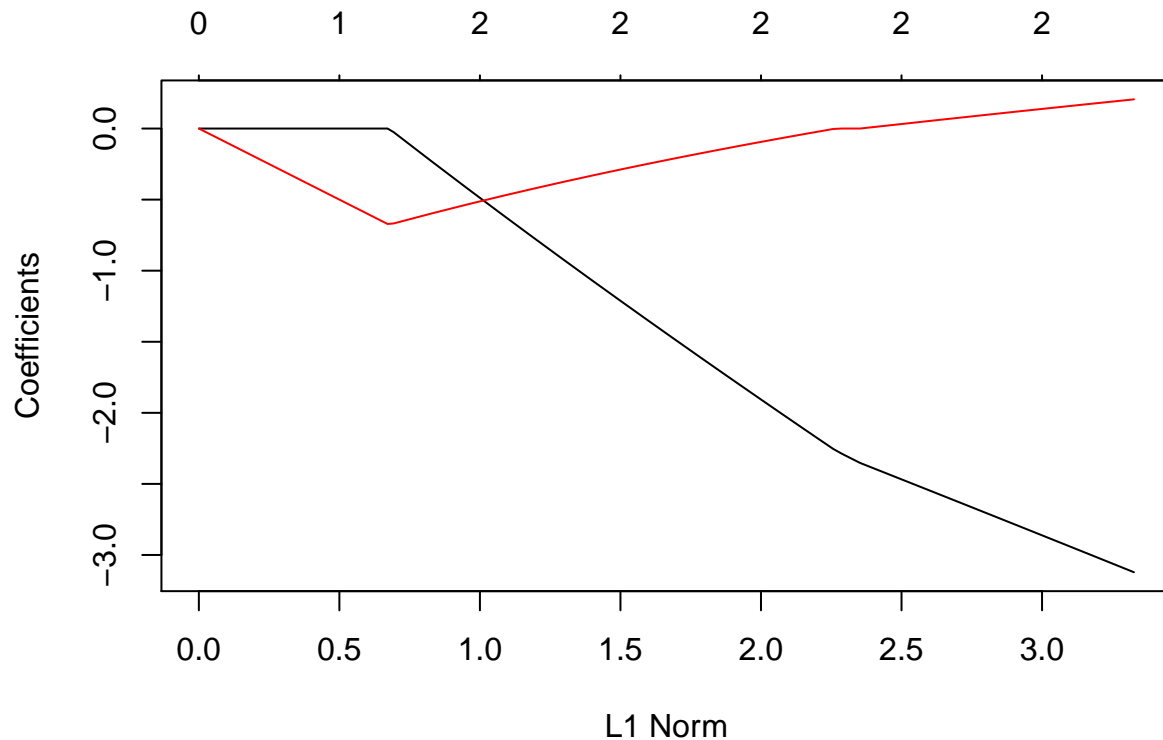


We provide a series of ranges for the interaction radii, and let the fitting function calibrate the model.

```
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 -3.121569 0.205435
plot(fit$complete)
```



```
print(fit$coefficients)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.000000 -1.378804 -3.121569 0.205435
print(fit$best_short)
#> [1]
#> [1,] 7.198859
print(fit$best_medium)
#> [1]
#> [1,] 16.98333
print(fit$best_long)
#> [1]
#> [1,] 23.77533
print(fit$aic)
#> [1] -372.2846
print(fit$bic)
#> [1] -363.288
```

We may then plot the corresponding Papangelou conditional intensity.

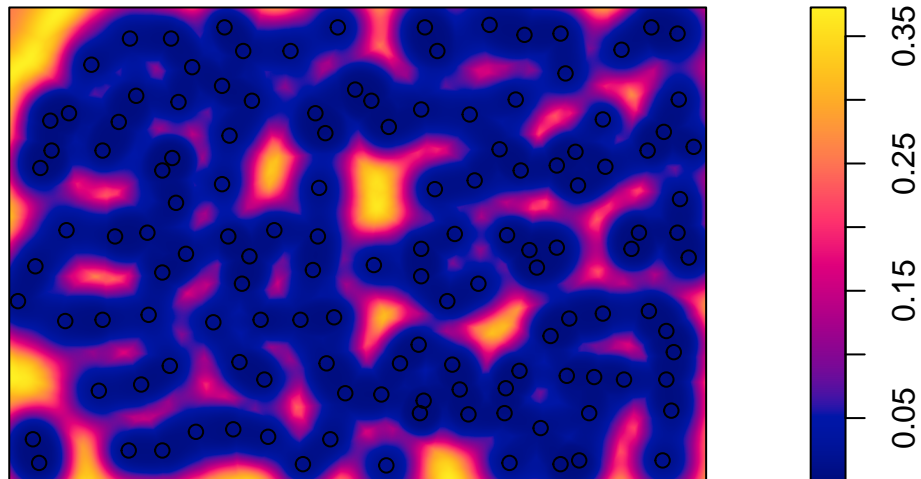
```
parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma
plot_papangelou(window = window,
                 configuration = configuration,
                 type = 1,
                 mark = mean(get_marks(configuration)),
```

```

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,

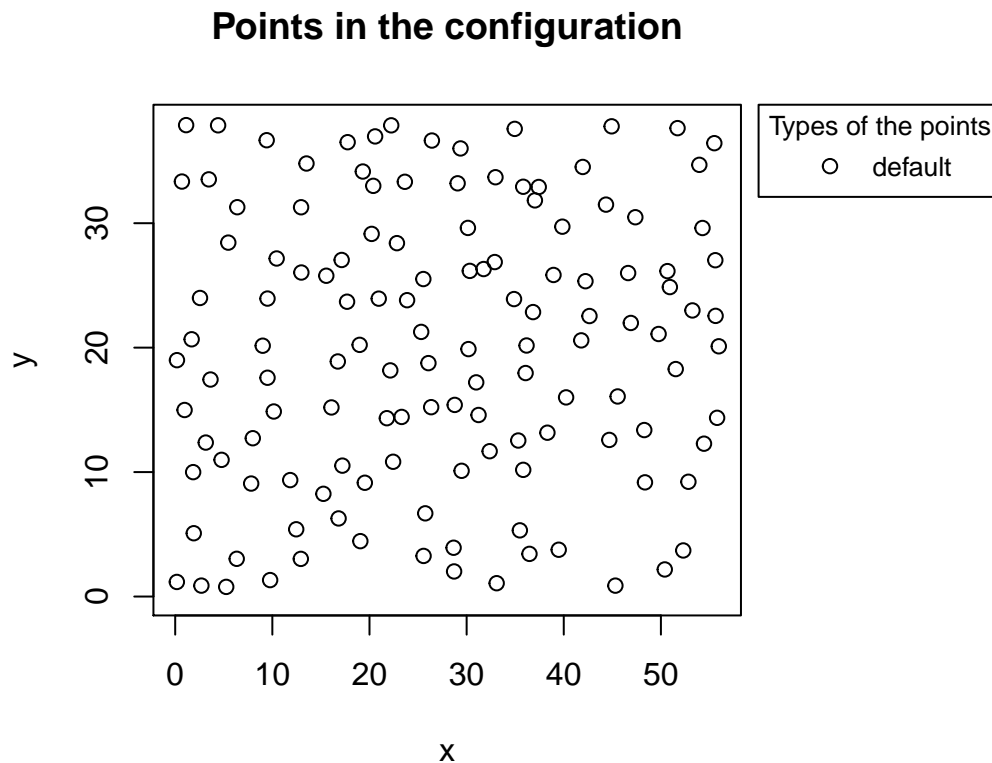
```

```

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: 130.

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

```



Interaction radii not proportional to marks

In this section, we do not account for marks.

```
configuration <- Configuration(spruces$x, spruces$y)
```

We call the fitting function on this unmarked point process.

```

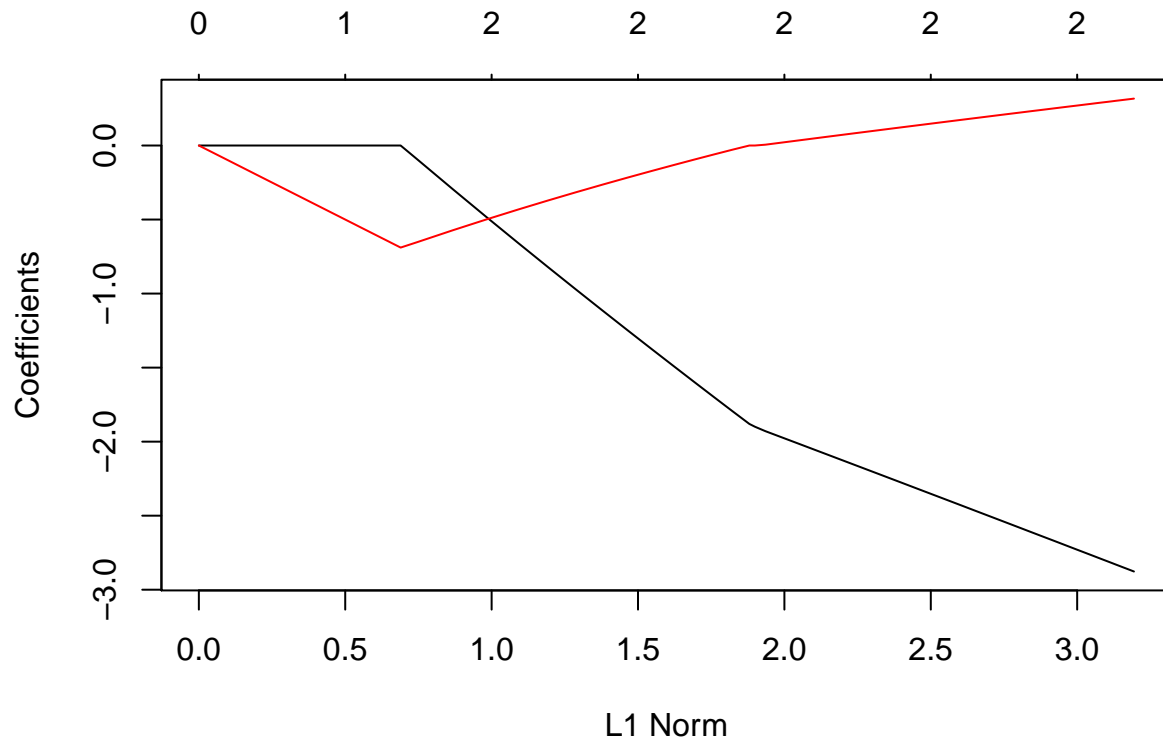
fit <- ppjsdm::gibbsm(configuration,
  window = window,
  model = model,
  medium_range_model = medium_range_model,
  short_range = short_range,
  medium_range = medium_range,

```

```

long_range = long_range,
use_glmnet = TRUE,
use_aic = TRUE,
saturation = 2)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.0000000 -1.3788037 -2.8771354 0.3167827
plot(fit$complete)

```



```

print(fit$coefficients)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.0000000 -1.3788037 -2.8771354 0.3167827
print(fit$best_short)
#> [1,] 2.048799
print(fit$best_medium)
#> [1,] 13.92174
print(fit$best_long)
#> [1,] 15.5096
print(fit$aic)
#> [1] -354.9212
print(fit$bic)
#> [1] -345.9246

```

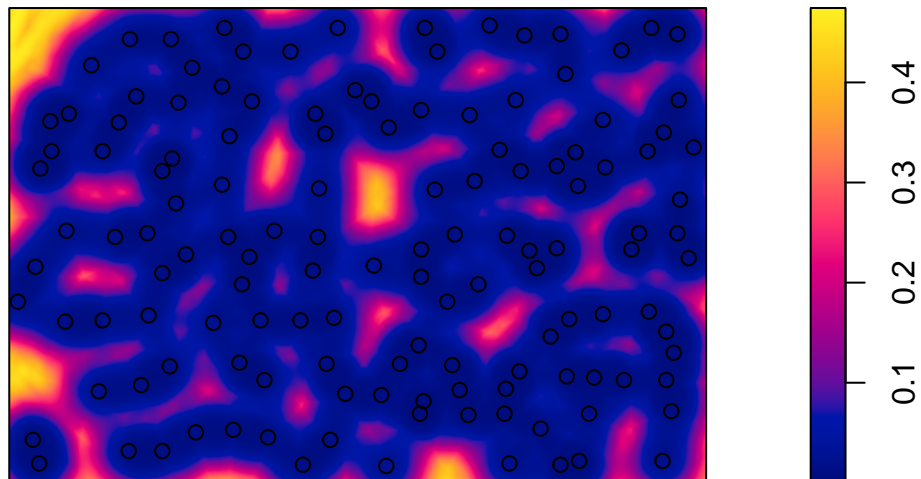
We may then plot the corresponding Papangelou conditional intensity.

```

parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma
plot_papangelou(window = window,
                 configuration = configuration,
                 type = 1,
                 mark = mean(get_marks(configuration)),
                 model = model,
                 medium_range_model = medium_range_model,
                 alpha = alpha,
                 lambda = lambda,
                 beta = matrix(0, 1, 0),
                 gamma = gamma,
                 covariates = list(),
                 short_range = fit$best_short,
                 medium_range = fit$best_medium,
                 long_range = fit$best_long,
                 saturation = 2)

```

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



And as previously, we draw from the model.

```

parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma

```

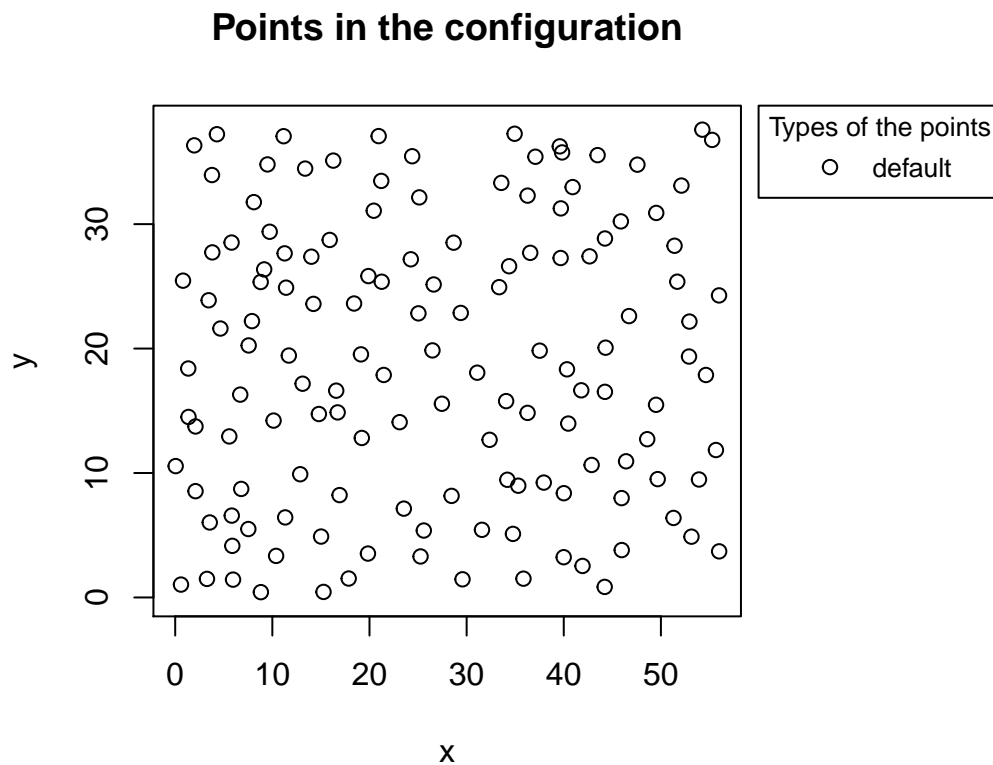
```

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: 137.

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

```

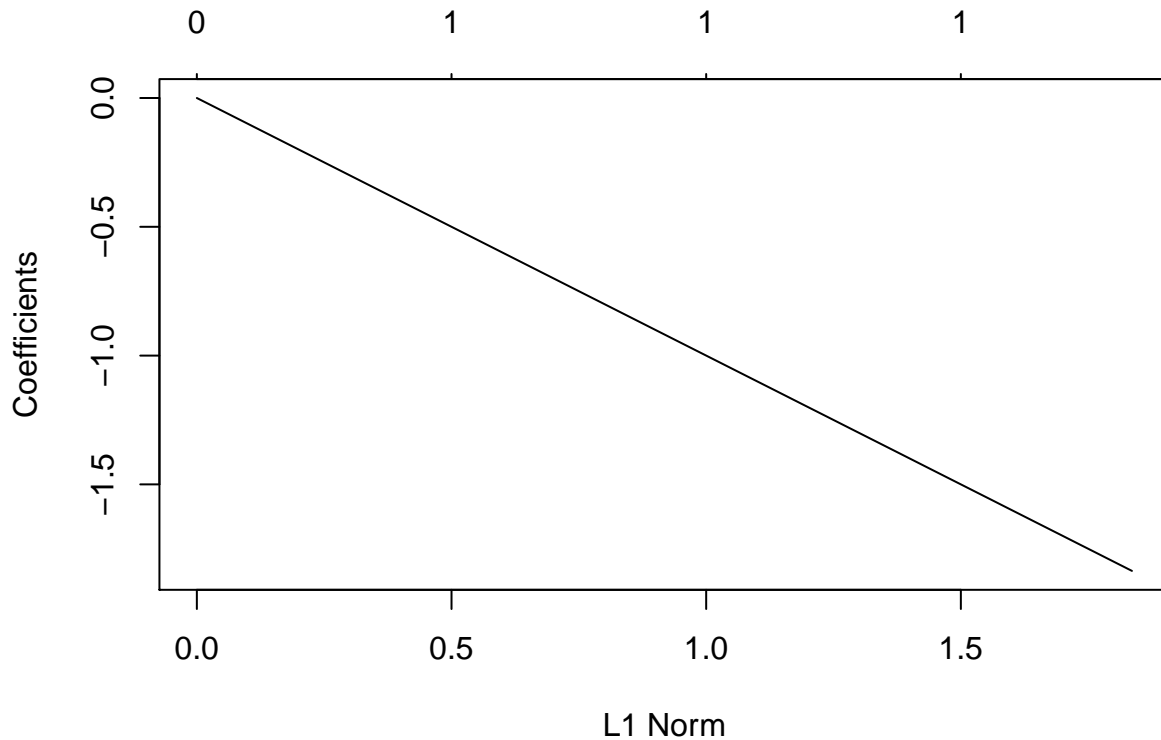


Comparison to classical Hard-core model.


```
configuration <- Configuration(spruces$x, spruces$y)
model <- "Geyer"
medium_range_model <- "Geyer"
```

We call the fitting function on this unmarked point process.

```
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 = 1)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.000000 -1.378804 -1.835898 0.000000
plot(fit$complete)
#> Warning in plotCoef(x$beta, lambda = x$lambda, df = x$df, dev = x$dev.ratio, : 1
#> or less nonzero coefficients; glmnet plot is not meaningful
```



```
print(fit$coefficients)
#> (Intercept) log_lambda1 alpha_1_1 gamma_1_1
#> 0.000000 -1.378804 -1.835898 0.000000
print(fit$best_short)
#> [ ,1]
```

```

#> [1,] 2.66237
print(fit$best_medium)
#>      [,1]
#> [1,] 12.44643
print(fit$best_long)
#>      [,1]
#> [1,] 22.93299
print(fit$aic)
#> [1] -309.05
print(fit$bic)
#> [1] -304.5487

```

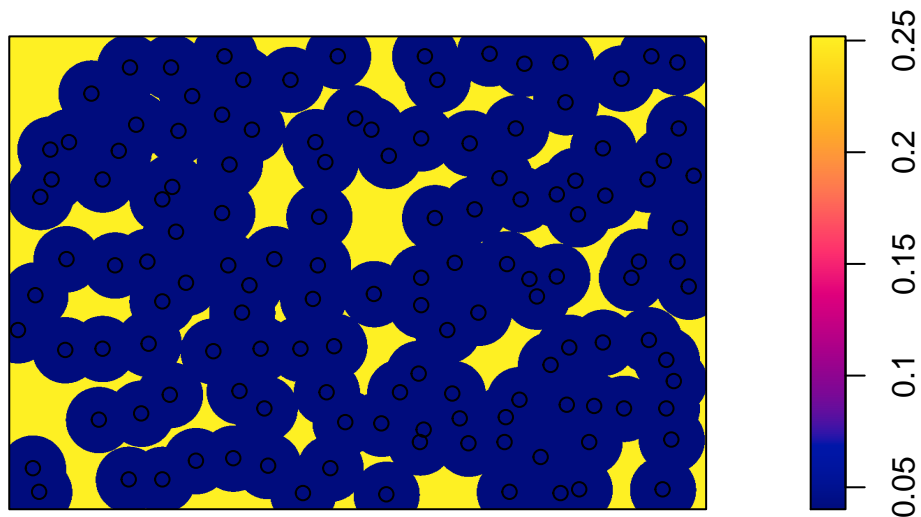
We may then plot the corresponding Papangelou conditional intensity.

```

parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma
plot_papangelou(window = window,
                 configuration = configuration,
                 type = 1,
                 mark = mean(get_marks(configuration)),
                 model = model,
                 medium_range_model = medium_range_model,
                 alpha = alpha,
                 lambda = lambda,
                 beta = matrix(0, 1, 0),
                 gamma = gamma,
                 covariates = list(),
                 short_range = fit$best_short,
                 medium_range = fit$best_medium,
                 long_range = fit$best_long,
                 saturation = 1)

```

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



And as previously, we 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 = 1000000)

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

