

Lansing dataset

```
library(ppjsdm)
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())

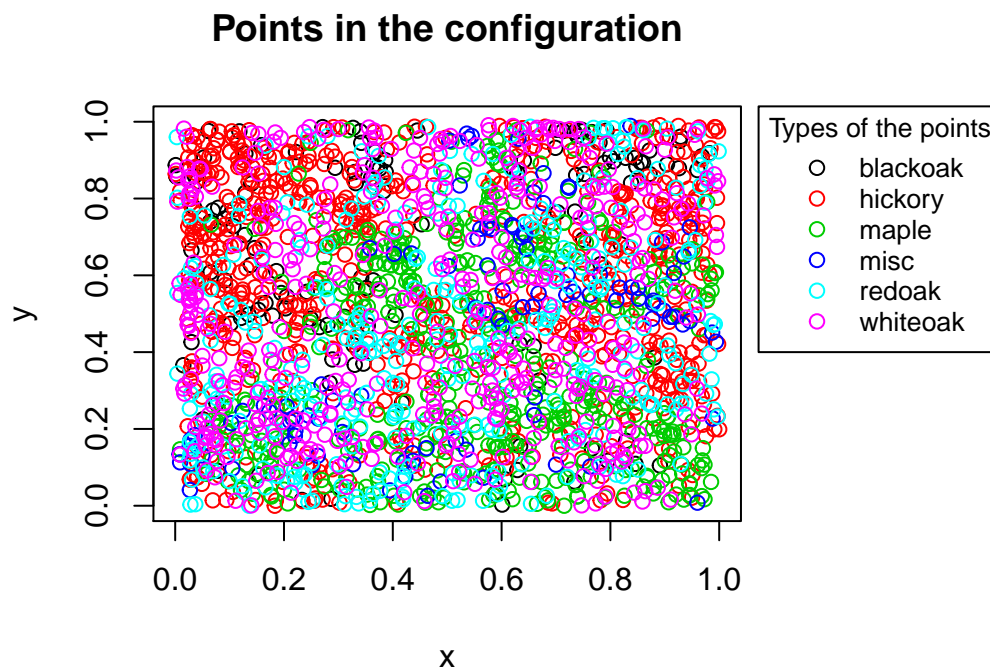
set.seed(1)
```

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

```
data(lansing)
configuration <- as.Configuration(lansing)
window <- Rectangle_window(c(0, 1), c(0, 1))
```

The point configuration is plotted below.

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



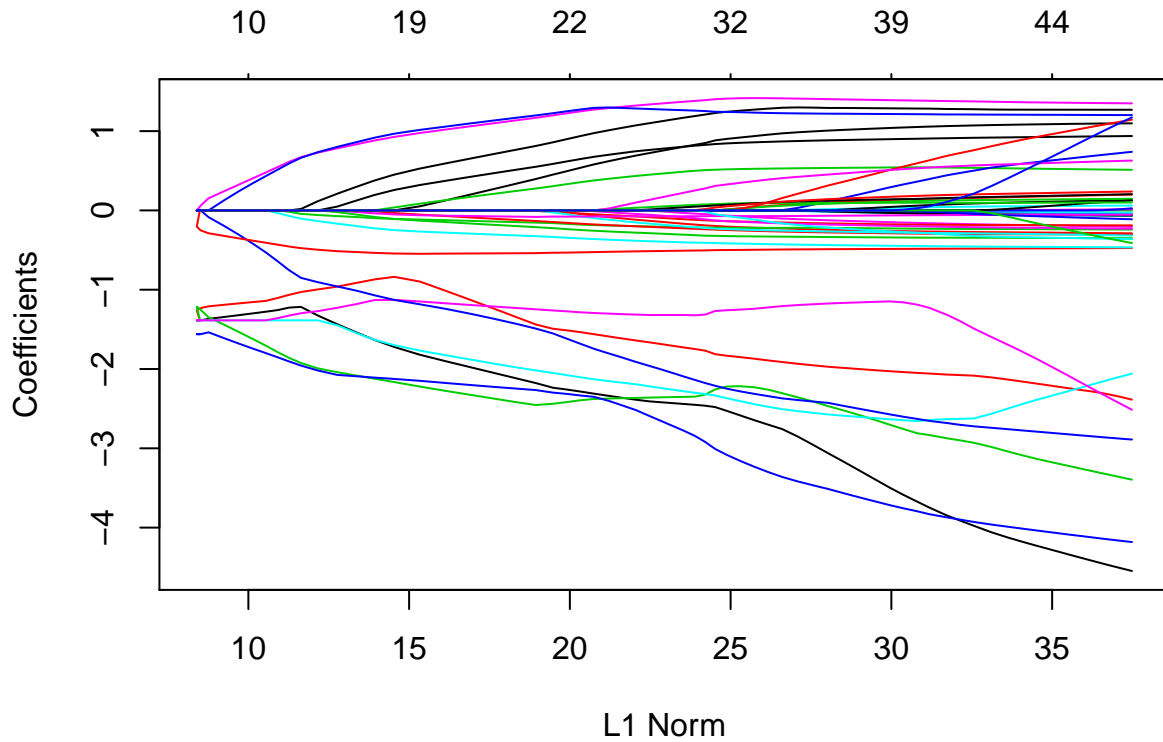
We fit the data with the Geyer model.

```
short_range <- c(0, 0.1)
medium_range <- c(0, 0.1)
long_range <- c(0, 0.1)
model <- "square_bump"
medium_range_model <- "square_exponential"
```

The matrix radii models interaction radii within a species, and between species.

```
library(GA)
#> Loading required package: foreach
#> Loading required package: iterators
#> Registered S3 method overwritten by 'cli':
#>   method      from
#>   print.boxx spatstat
#> Package 'GA' version 3.2
#> Type 'citation("GA")' for citing this R package in publications.
#>
#> Attaching package: 'GA'
#> The following object is masked from 'package:utils':
#>
#>   de
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)
#>      (Intercept) shifted_log_lambda1 shifted_log_lambda2 shifted_log_lambda3
#>      0.000000000      1.951228461      5.698058134      4.404886250
#> shifted_log_lambda4 shifted_log_lambda5 shifted_log_lambda6      alpha_1_1
#>      2.126328707      4.956884463      5.401294264      1.269946516
#>      alpha_1_2      alpha_1_3      alpha_1_4      alpha_1_5
#>     -0.282293460     -0.238050263     -2.824272293     -0.062633135
#>      alpha_1_6      alpha_2_2      alpha_2_3      alpha_2_4
#>     -0.193683296      1.090959696     -0.477229588      0.000000000
#>      alpha_2_5      alpha_2_6      alpha_3_3      alpha_3_4
#>     -0.029437068     -0.461679620      1.357647230     -0.059112559
#>      alpha_3_5      alpha_3_6      alpha_4_4      alpha_4_5
#>     -0.186497441     -0.348321440      1.204809229     -0.022587330
#>      alpha_4_6      alpha_5_5      alpha_5_6      alpha_6_6
#>     -0.222835173      0.929707515     -0.302092571      0.521135149
#>      gamma_1_1      gamma_1_2      gamma_1_3      gamma_1_4
#>      0.654463290      0.057139477      0.008525546      0.115030672
#>      gamma_1_5      gamma_1_6      gamma_2_2      gamma_2_3
#>      0.183628321      0.140219636      0.000000000     -0.304917935
#>      gamma_2_4      gamma_2_5      gamma_2_6      gamma_3_3
#>     -0.055647074      0.074849670      0.000000000      1.001921642
#>      gamma_3_4      gamma_3_5      gamma_3_6      gamma_4_4
#>      0.098435822     -0.086637654     -0.348059414      0.605807220
#>      gamma_4_5      gamma_4_6      gamma_5_5      gamma_5_6
#>      0.189267056      0.224372856      0.000000000     -0.249581615
```

```
#>      gamma_6_6
#>      0.785722135
plot(fit$complete)
```



```
print(fit$coefficients)
#>      (Intercept) shifted_log_lambda1 shifted_log_lambda2 shifted_log_lambda3
#>      0.000000000      1.951228461      5.698058134      4.404886250
#> shifted_log_lambda4 shifted_log_lambda5 shifted_log_lambda6      alpha_1_1
#>      2.126328707      4.956884463      5.401294264      1.269946516
#>      alpha_1_2      alpha_1_3      alpha_1_4      alpha_1_5
#>      -0.282293460      -0.238050263      -2.824272293      -0.062633135
#>      alpha_1_6      alpha_2_2      alpha_2_3      alpha_2_4
#>      -0.193683296      1.090959696      -0.477229588      0.000000000
#>      alpha_2_5      alpha_2_6      alpha_3_3      alpha_3_4
#>      -0.029437068      -0.461679620      1.357647230      -0.059112559
#>      alpha_3_5      alpha_3_6      alpha_4_4      alpha_4_5
#>      -0.186497441      -0.348321440      1.204809229      -0.022587330
#>      alpha_4_6      alpha_5_5      alpha_5_6      alpha_6_6
#>      -0.222835173      0.929707515      -0.302092571      0.521135149
#>      gamma_1_1      gamma_1_2      gamma_1_3      gamma_1_4
#>      0.654463290      0.057139477      0.008525546      0.115030672
#>      gamma_1_5      gamma_1_6      gamma_2_2      gamma_2_3
#>      0.183628321      0.140219636      0.000000000      -0.304917935
#>      gamma_2_4      gamma_2_5      gamma_2_6      gamma_3_3
#>      -0.055647074      0.074849670      0.000000000      1.001921642
#>      gamma_3_4      gamma_3_5      gamma_3_6      gamma_4_4
```

```

#>      0.098435822      -0.086637654      -0.348059414      0.605807220
#>      gamma_4_5      gamma_4_6      gamma_5_5      gamma_5_6
#>      0.189267056      0.224372856      0.000000000      -0.249581615
#>      gamma_6_6
#>      0.785722135
print(fit$best_short)
#>      [,1]      [,2]      [,3]      [,4]      [,5]      [,6]
#> [1,] 0.07438044 0.02074919 0.02074919 0.02074919 0.02074919 0.02074919
#> [2,] 0.02074919 0.05015042 0.02074919 0.02074919 0.02074919 0.02074919
#> [3,] 0.02074919 0.02074919 0.04098474 0.02074919 0.02074919 0.02074919
#> [4,] 0.02074919 0.02074919 0.02074919 0.04907959 0.02074919 0.02074919
#> [5,] 0.02074919 0.02074919 0.02074919 0.02074919 0.04214092 0.02074919
#> [6,] 0.02074919 0.02074919 0.02074919 0.02074919 0.02074919 0.03608095
print(fit$best_medium)
#>      [,1]      [,2]      [,3]      [,4]      [,5]      [,6]
#> [1,] 0.05557372 0.06441114 0.06441114 0.06441114 0.06441114 0.06441114
#> [2,] 0.06441114 0.04227391 0.06441114 0.06441114 0.06441114 0.06441114
#> [3,] 0.06441114 0.06441114 0.06181586 0.06441114 0.06441114 0.06441114
#> [4,] 0.06441114 0.06441114 0.06441114 0.05579612 0.06441114 0.06441114
#> [5,] 0.06441114 0.06441114 0.06441114 0.06441114 0.06670669 0.06441114
#> [6,] 0.06441114 0.06441114 0.06441114 0.06441114 0.06441114 0.03620342
print(fit$best_long)
#>      [,1]      [,2]      [,3]      [,4]      [,5]      [,6]
#> [1,] 0.1078154 0.10137172 0.10137172 0.10137172 0.1013717 0.10137172
#> [2,] 0.1013717 0.07480855 0.10137172 0.10137172 0.1013717 0.10137172
#> [3,] 0.1013717 0.10137172 0.09867876 0.10137172 0.1013717 0.10137172
#> [4,] 0.1013717 0.10137172 0.10137172 0.09012743 0.1013717 0.10137172
#> [5,] 0.1013717 0.10137172 0.10137172 0.10137172 0.1156353 0.10137172
#> [6,] 0.1013717 0.10137172 0.10137172 0.10137172 0.1013717 0.08114627
print(fit$aic)
#> [1] -5368.356

```

We may then plot the corresponding Papangelou conditional intensity.

```

parameters <- get_parameters_from_fit(fit)
lambda <- parameters$lambda
alpha <- parameters$alpha
gamma <- parameters$gamma
plot_papangelou(window = window,
  configuration = configuration,
  type = 1,
  model = model,
  medium_range_model = medium_range_model,
  alpha = alpha,
  lambda = lambda,
  beta = matrix(0, 6, 0),
  gamma = gamma,
  covariates = list(),
  short_range = fit$best_short,
  medium_range = fit$best_medium,
  long_range = fit$best_long,
  saturation = 2)
#> Warning: data contain duplicated points

```

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

