## Fin pines dataset

This vignette explains how to use the ppjsdm package with the finpines dataset from spatstat.

#### Taking marks into account

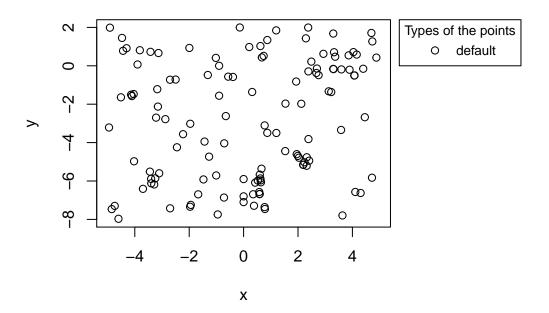
If marks are provided, the interaction radii are proportional to the marks. We begin with that setting.

```
configuration <- Configuration(finpines$x, finpines$y, marks = finpines$marks$height)
window <- Rectangle_window(c(-5, 5), c(-8, 2))</pre>
```

The point configuration is plotted below.

```
print(configuration)
#> An S3 object representing a configuration.
#>
#> Number of points: 126.
par(mar = c(5, 4, 4, 13) + 0.1)
plot(configuration, window = window)
```

### Points in the configuration

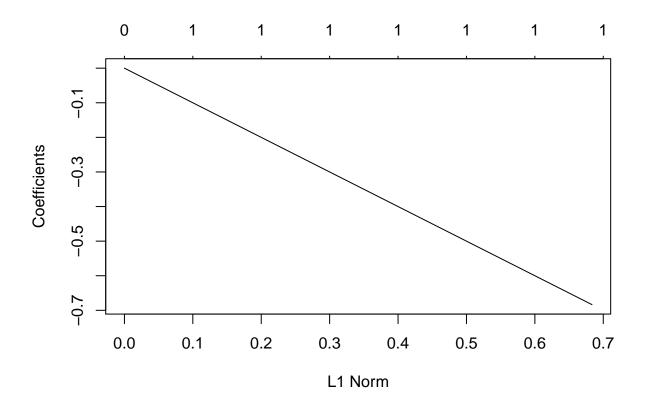


We provide a series of ranges for the interaction radii, and let the fitting function calibrate the model.

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

We can now call the fitting function.

```
fit <- ppjsdm::gibbsm(configuration,</pre>
                      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 1.6174061 -0.6834957
                                         0.0000000
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
```

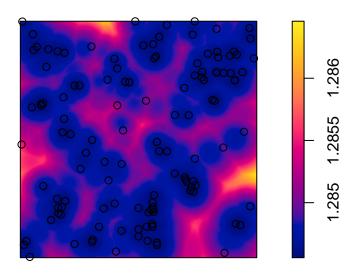


```
print(fit$coefficients)
#> (Intercept) log_lambda1
                             alpha_1_1
                                          gamma_1_1
     0.0000000 1.6174061 -0.6834957
                                         0.0000000
print(fit$best_short)
           [,1]
#> [1,] 15.2557
print(fit$best_medium)
#>
            [,1]
#> [1,] 26.50541
print(fit$best_long)
            [,1]
#> [1,] 35.95118
print(fit$aic)
#> [1] -240.8092
print(fit$bic)
#> [1] -236.3699
```

We may then plot the corresponding Papangelou conditional intensity.

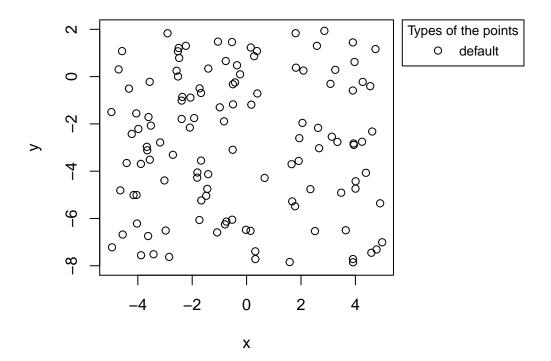
```
model = model,
medium_range_model = medium_range_model,
alpha = alpha,
lambda = lambda,
beta = matrix(0, 1, 0),
gamma = gamma,
covariates = list(),
short_range = fit$best_short,
medium_range = fit$best_medium,
long_range = fit$best_long,
saturation = 2)
```

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



It is also possible to draw from the model.

### Points in the configuration

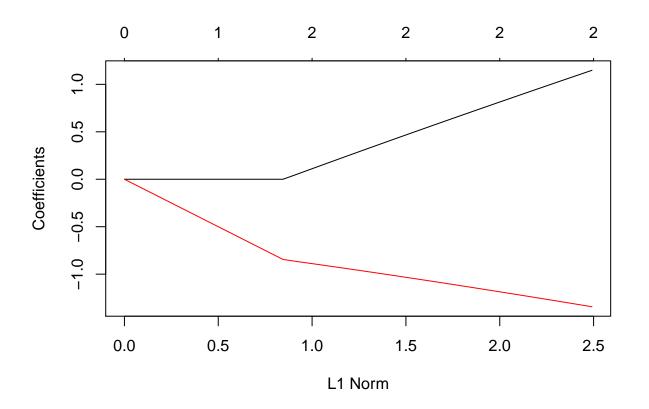


# Interaction radii not proportional to marks

```
In this section, we do not account for marks.
```

```
configuration <- Configuration(finpines$x, finpines$y)</pre>
```

We call the fitting function on this unmarked point process.

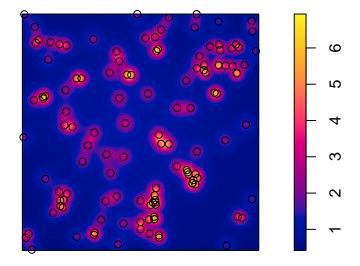


```
print(fit$coefficients)
#> (Intercept) log_lambda1
                             alpha_1_1
                                          gamma_1_1
      0.000000
                  1.617406
                              1.147943
                                         -1.343462
print(fit$best_short)
             [,1]
#>
#> [1,] 0.3419659
print(fit$best_medium)
            [,1]
#>
#> [1,] 11.41415
print(fit$best_long)
            [,1]
#> [1,] 25.13677
print(fit$aic)
#> [1] -274.2492
print(fit$bic)
#> [1] -265.3769
```

We may then plot the corresponding Papangelou conditional intensity.

```
parameters <- get_parameters_from_fit(fit)</pre>
lambda <- parameters$lambda</pre>
alpha <- parameters$alpha</pre>
gamma <- parameters$gamma</pre>
plot_papangelou(window = window,
                 configuration = configuration,
                 type = 1,
                 mark = mean(get_marks(configuration)),
                 model = model,
                 medium_range_model = medium_range_model,
                 alpha = alpha,
                 lambda = lambda,
                 beta = matrix(0, 1, 0),
                 gamma = gamma,
                 covariates = list(),
                 short_range = fit$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</pre>
```

```
draw <- ppjsdm::rgibbs(window = window,</pre>
                       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: 176.
par(mar = c(5, 4, 4, 13) + 0.1)
plot(draw, window = window)
```

## Points in the configuration

