

# Package: spatgeom (via r-universe)

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**Type** Package

**Title** Geometric Spatial Point Analysis

**Version** 0.3.0

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**Description** The implementation to perform the geometric spatial point analysis developed in Hernández & Solís (2022) <doi:10.1007/s00180-022-01244-1>. It estimates the geometric goodness-of-fit index for a set of variables against a response one based on the 'sf' package. The package has methods to print and plot the results.

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**URL** <https://github.com/maikol-solis/spatgeom>

**BugReports** <https://github.com/maikol-solis/spatgeom/issues>

**Encoding** UTF-8

**LazyData** true

**Imports** ggplot2, scales, sf, dplyr, lwgeom, cowplot, purrr

**RoxygenNote** 7.3.1

**Roxygen** list(markdown=TRUE)

**Depends** R (>= 3.6.0)

**Suggests** rmarkdown, knitr, testthat (>= 2.1.0)

**Repository** <https://maikol-solis.r-universe.dev>

**RemoteUrl** <https://github.com/maikol-solis/spatgeom>

**RemoteRef** HEAD

**RemoteSha** c470f2d816016f577983c25cffdd2e1b76353450

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donut_data	<i>Donut example</i>
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### Description

Generate data points with the shape of a donut.

### Usage

```
donut_data(n, a, b, theta)
```

### Arguments

n	Number of data points.
a	Lower bound of the second variable.
b	Upper bound of the second variable.
theta	Angle of the donut.

### Value

A data frame with three variables. Variable 'y' is the response, variable 'x1' makes the donut shape with 'y', and 'x2' is a uniform random variable between a and b. '

### Examples

```
xy <- donut_data(n = 30, a = -1, b = 1, theta = 2 * pi)
```

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linear_data	<i>Linear example</i>
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### Description

Generate data points with a linear relationship.

### Usage

```
linear_data(n = 100, a = -3, b = 3)
```

**Arguments**

n                      Number of data points.  
 a, b                    Lower and upper bound of the uniform distribution.

**Value**

A data frame with three variables. Variable 'y = 0.6 \* x1 + 0.3 \* x2

- 0.1 \* x3' is the response, and 'x1', 'x2' and 'x3' are uniform random variables between a and b.

**Examples**

```
xy <- linear_data(n = 30, a = -1, b = 1)
```

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plot\_alpha\_shape            *Plot alpha-shape for spatgeom objects*

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**Description**

Plot alpha-shape for spatgeom objects.

**Usage**

```
plot_alpha_shape(x, alpha, font_size = 12)
```

**Arguments**

x                      an object of class spatgeom.  
 alpha                 value of alpha determining the maximum length between points to build the alpha-shape.  
 font\_size            a integer that increases the font size in the plot.

**Value**

a [ggplot](#) object with the raw alpha-shape for the original data at resolution alpha

**Examples**

```
xy <- donut_data(n = 30, a = -1, b = 1, theta = 2 * pi)
estimation <- spatgeom(y = xy[, 1], x = xy[, -1])
plot_alpha_shape(estimation, alpha = c(0.9, 1.2))
```

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plot_curve	<i>plot spatgeom objects</i>
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### Description

Plot method for objects of class spatgeom.

### Usage

```
plot_curve(x, type = "curve", font_size = 12)
```

### Arguments

x	an object of class spatgeom
type	a string that could be curve or deriv. The option curve plots the curve of alpha against geom_corr from the function <code>spatgeom()</code> . The deriv option plots the numerical derivative.
font_size	a integer that increases the font size in the plot.

### Value

a `ggplot` object with the geometric indices (or its derivative). The plot is generated with the nalphas point of alpha and geom\_corr from the function `spatgeom`.

In each panel, the theoretical CSR process is drawn using  $\exp(-\text{intensity} * \pi * x^2)$ . where the intensity depends on each panel.

### Examples

```
xy <- donut_data(n = 30, a = -1, b = 1, theta = 2 * pi)
estimation <- spatgeom(y = xy[, 1], x = xy[, -1])
plot_curve(estimation, type = "curve")
plot_curve(estimation, type = "deriv")
```

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print.spatgeom	<i>print a spatgeom object</i>
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### Description

Print method for objects of class spatgeom.

**Usage**

```
## S3 method for class 'spatgeom'  
print(x, return_table = FALSE, ...)
```

**Arguments**

`x` an object of class `spatgeom`

`return_table` if TRUE, returns a data frame with the estimated values. Otherwise, print the data frame in console. Defaults to FALSE

`...` further arguments passed to the `plot` function

**Value**

Print the estimate given by `spatgeom`.

**Examples**

```
xy <- donut_data(n = 30, a = -1, b = 1, theta = 2 * pi)  
  
estimation <- spatgeom(y = xy[, 1], x = xy[, -1])  
  
print(estimation)
```

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spatgeom

*Geometric Spatial Point Pattern Analysis*

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**Description**

Function to estimate the geometric correlation between variables.

**Usage**

```
spatgeom(  
  x,  
  y,  
  scale_pts = FALSE,  
  nalphas = 100,  
  envelope = FALSE,  
  domain_type = c("bounding-box", "convex-hull"),  
  mc_cores = 1  
)
```

**Arguments**

<code>x</code>	numeric matrix or data.frame of covariables.
<code>y</code>	numeric vector of responses in a model.
<code>scale_pts</code>	boolean to make the estimations with scaled variables. Default FALSE.
<code>nalphas</code>	number of alphas generated for creating the geometric measure of fit index. Default 100.
<code>envelope</code>	boolean to determine if the Monte-Carlo should be estimated. Default FALSE.
<code>domain_type</code>	character with the type of domain to use. It can be either "bounding-box" or "convex-hull". Default "bounding-box".
<code>mc_cores</code>	integer with the number of parallel process to run (if available). Default 1.

**Value**

A list of class `spatgeom` with the following elements:

**call** The function call.

**x** `x` input.

**y** `y` output.

**results** A list of size `ncol(x)` corresponding to each column of `x`. Each element of the list has:

**triangles** a data frame of class `sfc` (see `sf::st_sf()`) with columns `geometry`, `segments`, `max_length` and `alpha`. The data.frame contains the whole Delanauy triangulation for the corresponding column of `x` and `y`. The `segments` column are the segments of each individual triangle and `max_length` is the maximum length of them.

**geom\_indices** a data frame with columns `alpha` and `geom_survival`. The `alpha` column is a numeric vector of size `nalphas` from the minimum to the maximum distance between points estimated in the data. The `geom_survival` column is the value  $1 - (\text{alpha shape Area}) / (\text{containing box Area})$ .

**intensity** the intensity estimated for the corresponding column of `x` and `y`.

**mean\_n** the mean number of points in the point process.

**envelope\_data** a data frame in tidy format with 40 runs of a CSR process, if `envelope=TRUE`, The CSR is created by generating  $n$  uniform points in the plane, where  $n$  is drawn from Poisson distribution with parameter `mean_n`.

**References**

Hernández, A.J., Solís, M. Geometric goodness of fit measure to detect patterns in data point clouds. *Comput Stat* (2022). <https://doi.org/10.1007/s00180-022-01244-1>

**Examples**

```
xy <- donut_data(n = 30, a = -1, b = 1, theta = 2 * pi)
estimation <- spatgeom(y = xy[, 1], x = xy[, -1])

# If you want to estimate the envelope, you can use the envelope argument to
# TRUE. This will take a while to run.
## Not run:
```

```
estimation_with_envelope <- spatgeom(  
  y = xy[, 1], x = xy[, -1],  
  envelope = TRUE  
)  
  
## End(Not run)
```

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