Package 'ggpolar'

October 13, 2022

Type Package

Title Dots and Their Connections in Polar Coordinate System

Version 0.2.2

Maintainer Shixiang Wang <w_shixiang@163.com>

Description Provides basic graphing functions to fully demonstrate
point-to-point connections in a polar coordinate space.

License GPL (>= 3)

URL https://github.com/ShixiangWang/polar

BugReports https://github.com/ShixiangWang/polar/issues

Depends ggplot2

Imports rlang

Suggests ezcox, ggnewscale, knitr, rmarkdown, survival, cli

VignetteBuilder knitr

Encoding UTF-8

RoxygenNote 7.2.1

NeedsCompilation no

Author Shixiang Wang [aut, cre] (<https://orcid.org/0000-0001-9855-7357>)

Repository CRAN

Date/Publication 2022-10-10 10:50:02 UTC

R topics documented:

polar_connect	*Connects dots*

Description

Check polar_init() for examples.

Usage

```
polar_connect(data, x1, x2, ...)
```

Arguments

data	a data.frame contains connections of all events.
x1, x2	the column names (without quote) storing connected events.
...	parameters passing to ggplot2::geom_segment, expect c(x, xend, y, yend) these 4 mapping parameters.

Value

a ggplot object.

polar_init	*Init a dot plot in polar system*

Description

Init a dot plot in polar system

Usage

```
polar_init(data, x, ...)
```

Arguments

data	a data.frame contains all events, e.g., genes.
x	the column name (without quote) storing event list.
...	parameters passing to ggplot2::geom_point.

Value

a ggplot object.

Examples

```
# -------------------
#  Init a polar plot
# -------------------

data <- data.frame(x = LETTERS[1:7])

p1 <- polar_init(data, x = x)
p1

# Set aes value
p2 <- polar_init(data, x = x, size = 3, color = "red", alpha = 0.5)
p2

# Set aes mapping
set.seed(123L)
data1 <- data.frame(
  x = LETTERS[1:7],
  shape = c("r", "r", "r", "b", "b", "b", "b"),
  color = c("r", "r", "r", "b", "b", "b", "b"),
  size = abs(rnorm(7))
)
# Check https://ggplot2.tidyverse.org/reference/geom_point.html
# for how to use both stroke and color
p3 <- polar_init(data1, x = x, aes(size = size, color = color, shape = shape), alpha = 0.5)
p3

# -------------------
#  Connect polar dots
# -------------------
data2 <- data.frame(
  x1 = LETTERS[1:7],
  x2 = c("B", "C", "D", "E", "C", "A", "C"),
  color = c("r", "r", "r", "b", "b", "b", "b")
)
p4 <- p3 + polar_connect(data2, x1, x2)
p4

# Unlike polar_init, mappings don't need to be included in aes()
p5 <- p3 + polar_connect(data2, x1, x2, color = color, alpha = 0.8, linetype = 2)
p5

# Use two different color scales
if (requireNamespace("ggnewscale")) {
  library(ggnewscale)
  p6 = p3 +
    new_scale("color") +
    polar_connect(data2, x1, x2, color = color, alpha = 0.8, linetype = 2)
  p6 + scale_color_brewer()
  p6 + scale_color_manual(values = c("darkgreen", "magenta"))
}
```

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