

Package ‘PieGlyph’

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Title Axis Invariant Scatter Pie Plots

Version 0.1.0

Description

Extends 'ggplot2' to help replace points in a scatter plot with pie-chart glyphs showing the relative proportions of different categories. The pie glyphs are independent of the axes and plot dimensions, to prevent distortions when the plot dimensions are changed.

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Encoding UTF-8

Imports ggplot2, dplyr, tidyr, rlang, ggforce, forcats, plyr, grid, scales, cli, utils

Suggests ranger, maps, mapproj, knitr, tidyverse, rmarkdown

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VignetteBuilder knitr

URL <https://github.com/rishvish/PieGlyph>

BugReports <https://github.com/rishvish/PieGlyph/issues>

NeedsCompilation no

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draw_key_pie	<i>Legend key for the pie glyphs</i>
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Description

Controls the aesthetics of the legend entries for the pie glyphs

Usage

```
draw_key_pie(data, params, size)
```

Arguments

data	A single row data frame containing the scaled aesthetics to display in this key
params	A list of additional parameters supplied to the geom.
size	Width and height of key in mm.

Value

A grid grob

See Also

[draw_key](#)

geom_pie_glyph	<i>Scatter plot with points replaced by axis-invariant pie-chart glyphs</i>
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Description

This geom replaces the points in a scatter plot with pie-chart glyphs showing the relative proportions of different categories. The pie-chart glyphs are independent of the plot dimensions, so won't distort when the plot is scaled. The ideal dataset for this geom would contain columns with non-negative values showing the magnitude of the different categories to be shown in the pie glyphs (The proportions of the different categories within the pie glyph will be calculated automatically). The different categories can also be stacked together into a single column according to the rules of tidy-data (see vignette('tidy-data') or vignette('pivot') for more information).

Usage

```
geom_pie_glyph(
  mapping = NULL,
  data = NULL,
  slices,
  values = NA,
  stat = "identity",
  position = "identity",
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE,
  ...
)
```

Arguments

mapping	Set of aesthetic (see Aesthetics below) mappings to be created by <code>aes()</code> or <code>aes_()</code> . If specified and <code>inherit.aes = TRUE</code> (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.
data	The data to be displayed in this layer of the plot. The default, <code>NULL</code> , inherits the plot data specified in the <code>ggplot()</code> call. A <code>data.frame</code> , or other object, will override the plot data. All objects will be fortified to produce a data frame. See <code>fortify()</code> for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a <code>data.frame</code> , and will be used as the layer data. A function can be created from a formula (e.g. <code>~ head(.x, 10)</code>).
slices	Each pie glyph in the plot shows the relative abundances of a set of categories; those categories are specified by this argument and should contain numeric and non-negative values. The names of the categories can be the names of individual columns (wide format) or can be stacked and contained in a single column (long format using <code>pivot_longer()</code>). The categories can also be specified as the numeric indices of the columns.
values	This parameter is not needed if the data is in wide format. The default is <code>NA</code> assuming that the categories are in separate columns. However, if the pie categories are stacked in one column, this parameter describes the column for the values of the categories shown in the pie glyphs. The values should be numeric and non-negative and the proportions of the different slices within each observation will be calculated automatically.
stat	The statistical transformation to use on the data for this layer, either as a ggproto <code>Geom</code> subclass or as a string naming the stat stripped of the <code>stat_</code> prefix (e.g. "count" rather than "stat_count")
position	Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use <code>position_jitter</code>), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.


```

C = round(runif(10, 3, 7), 2),
D = round(runif(10, 1, 9), 2))

head(plot_data)

## Basic plot
ggplot(data = plot_data, aes(x = system, y = response))+
  geom_pie_glyph(slices = c('A', 'B', 'C', 'D'),
                data = plot_data)+
  theme_classic()

## Change pie radius and border colour
ggplot(data = plot_data, aes(x = system, y = response))+
  # Can also specify slices as column indices
  geom_pie_glyph(slices = 4:7, data = plot_data,
                colour = 'black', radius = 0.5)+
  theme_classic()

## Map radius to a variable
p <- ggplot(data = plot_data, aes(x = system, y = response))+
  geom_pie_glyph(aes(radius = group),
                slices = c('A', 'B', 'C', 'D'),
                data = plot_data, colour = 'black')+
  theme_classic()

p

## Add custom labels
p <- p + labs(x = 'System', y = 'Response',
             fill = 'Attributes', radius = 'Group')

p

## Change slice colours
p + scale_fill_manual(values = c('#56B4E9', '#CC79A7',
                                '#F0E442', '#D55E00'))

##### Stack the attributes in one column
# The attributes can also be stacked into one column to generate
# the plot. This variant of the function is useful for situations
# when the data is in tidy format. See vignette('tidy-data') and
# vignette('pivot') for more information.

plot_data_stacked <- plot_data %>%
  pivot_longer(cols = c('A', 'B', 'C', 'D'),
              names_to = 'Attributes',
              values_to = 'values')

head(plot_data_stacked, 8)

```

```
ggplot(data = plot_data_stacked, aes(x = system, y = response))+
  # Along with slices column, values column is also needed now
  geom_pie_glyph(slices = 'Attributes', values = 'values')+
  theme_classic()
```

pieGrob

Create pie-chart glyph

Description

This function creates a pie-chart glyph. The proportions of the different slices are calculated automatically using the numbers in the values parameter.

Usage

```
pieGrob(
  x = 0.5,
  y = 0.5,
  values,
  radius = 1,
  radius_unit = "cm",
  edges = 360,
  col = "black",
  fill = NA,
  lwd = 1,
  lty = 1,
  alpha = 1,
  default.units = "npc"
)
```

Arguments

x	A number or unit object specifying x-location of pie chart
y	A number or unit object specifying y-location of pie chart
values	A numeric vector specifying the values of the different slices of the pie chart
radius	A number specifying the radius of the pie-chart
radius_unit	Character string specifying the unit for the radius of the pie-chart
edges	Number of edges which make up the circumference of the pie-chart (Increase for higher resolution)
col	Character specifying the colour of the border between the pie slices
fill	A character vector specifying the colour of the individual slices
lwd	Line width of the pie borders
lty	Linetype of the pie borders
alpha	Number between 0 and 1 specifying the opacity of the pie-charts
default.units	Change the default units for the position and radius of the pie-glyphs

Value

A grob object

Examples

```
library(grid)
grid.newpage()
p1 <- pieGrob(x = 0.2, y= 0.2,
             values = c(.7,.1,.1,.1), radius = 1,
             fill = c('purple','red','green','orange'))
grid.draw(p1)

## Change unit of radius
grid.newpage()
p2 <- pieGrob(x = 0.5, y= 0.75,
             values = c(1,2,3,4,5), radius = 1,
             radius_unit = 'in',
             fill = c('purple','yellow','green','orange','blue'))
grid.draw(p2)

## Change border attributes
grid.newpage()
p3 <- pieGrob(x = 0.5, y= 0.5,
             values = c(10, 40, 50), radius = 20,
             radius_unit = 'mm',
             col = 'red', lwd = 5, lty = 3,
             fill = c('purple','yellow','blue'))
grid.draw(p3)
```

scale_radius_discrete *Scales for the pie glyph radius*

Description

scale_radius_*() is useful for adjusting the radius of the pie glyphs.

Usage

```
scale_radius_discrete(..., range = c(0.25, 0.6), unit = "cm")

scale_radius_manual(..., values, unit = "cm", breaks = waiver(), na.value = NA)

scale_radius_continuous(..., range = c(0.25, 0.6), unit = "cm")

scale_radius(..., range = c(0.25, 0.6), unit = "cm")
```

Arguments

...	Arguments passed on to continuous_scale
minor_breaks	One of: <ul style="list-style-type: none"> • NULL for no minor breaks • <code>waiver()</code> for the default breaks (one minor break between each major break) • A numeric vector of positions • A function that given the limits returns a vector of minor breaks. Also accepts rlang lambda function notation.
oob	One of: <ul style="list-style-type: none"> • Function that handles limits outside of the scale limits (out of bounds). Also accepts rlang lambda function notation. • The default (<code>scales:::censor()</code>) replaces out of bounds values with NA. • <code>scales:::squish()</code> for squishing out of bounds values into range. • <code>scales:::squish_infinite()</code> for squishing infinite values into range.
na.value	Missing values will be replaced with this value.
expand	For position scales, a vector of range expansion constants used to add some padding around the data to ensure that they are placed some distance away from the axes. Use the convenience function expansion() to generate the values for the expand argument. The defaults are to expand the scale by 5% on each side for continuous variables, and by 0.6 units on each side for discrete variables.
position	For position scales, The position of the axis. left or right for y axes, top or bottom for x axes.
super	The super class to use for the constructed scale
range	a numeric vector of length 2 that specifies the minimum and maximum size of the plotting symbol after transformation.
unit	Unit for the radius of the pie glyphs. Default is "cm", but other units like "in", "mm", etc. can be used.
values	a set of aesthetic values to map data values to. The values will be matched in order (usually alphabetical) with the limits of the scale, or with breaks if provided. If this is a named vector, then the values will be matched based on the names instead. Data values that don't match will be given <code>na.value</code> .
breaks	One of: <ul style="list-style-type: none"> • NULL for no breaks • <code>waiver()</code> for the default breaks computed by the transformation object • A numeric vector of positions • A function that takes the limits as input and returns breaks as output (e.g., a function returned by <code>scales:::extended_breaks()</code>). Also accepts rlang lambda function notation.
na.value	The aesthetic value to use for missing (NA) values

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