Package 'pacviz'

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Title Pac-Man Residual Function		
Version 1.0.0.0 Description License MIT + file LICENSE Depends R (>= 3.3.3)		
		Imports circlize, graphics, plotrix, stats, utils
		Suggests knitr, rmarkdown
		VignetteBuilder knitr
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LazyData true		
RoxygenNote 7.1.1		
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pacman Pac-Man Residual Function		

Description

A visualization technique in R for regression analysis results, specifically residual values, based on a restricted radial coordinate system. It provides a broad view perspective on the performance of regression models, and supports most model inputs. See the pacviz documentation page for more information: https://pharaohcola13.github.io/pacviz/book/

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Usage

```
pacman(
    x,
    y,
    title,
    unit,
    axis_label,
    model = lm(y ~ x, data = data.frame(x, y)),
    color1 = "gold",
    standardize = FALSE
)
```

Arguments

x, y	Numeric data
title	Figure title
unit	String to define units on the angular axis (For temperature measurements use 'degC' or 'degF')
axis_label	Angular axis label
model	An object for which the extraction of model residuals is meaningful.
color1	Color value as string or rgb
standardize	Boolean to standardize the residual value

Value

Pac-Man residual plot

Examples

```
# Generic Pac-Man residual
x <- rnorm(20, mean=0, sd=10)
y <- log(rnorm(20, mean=0, sd=10), base=exp(1))
pacman(x,y,'Example 1','units', 'Axis Label')

# Pac-Man residual using alternate color, residual standardization, and temperature units
x <- rnorm(20, mean=0, sd=10)
y <- log(rnorm(20, mean=0, sd=10), base=exp(1))
pacman(x,y, 'Example 2', 'degC', "Temperature", color1="lightblue", standardize=TRUE)</pre>
```

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```