

Package: OddsPlotty (via r-universe)

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Type Package

Title Odds Plot to Visualise a Logistic Regression Model

Version 1.0.2

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Description Uses the outputs of a logistic regression model, from `caret` <<https://CRAN.R-project.org/package=caret>>, to build an odds plot. This allows for the rapid visualisation of odds plot ratios and works best with the outputs of CARET's GLM model class, by returning the final trained model.

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URL <https://github.com/StatsGary/OddsPlotty>

LazyData FALSE

Imports caret, mlbench, magrittr, ggplot2, tibble, ggthemes, e1071, tidymodels, rmarkdown

Suggests knitr, covr, testthat, markdown

VignetteBuilder knitr

RoxygenNote 7.1.13

Encoding UTF-8

Repository <https://statsgary.r-universe.dev>

RemoteUrl <https://github.com/statsgary/oddsplotty>

RemoteRef HEAD

RemoteSha 018e7c1a4176c9f4b64b8054abcc117bb49349d7

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`odds_plot`*odds_plot - a function to create Odds Plots*

Description

This has been created to generate odds plots on the back of results from a generalised linear model.

Usage

```
odds_plot(  
  x,  
  x_label = "Variables",  
  y_label = "Odds Ratio",  
  title = NULL,  
  subtitle = NULL,  
  point_col = "blue",  
  error_bar_colour = "black",  
  point_size = 5,  
  error_bar_width = 0.3,  
  h_line_color = "black"  
)
```

Arguments

<code>x</code>	The trained caret GLM logistic regression model
<code>x_label</code>	The label name for the <code>x_label</code>
<code>y_label</code>	The label name for the <code>y_label</code>
<code>title</code>	Title for the Odds Plot
<code>subtitle</code>	Subtitle for the Odds Plot
<code>point_col</code>	Defaults to blues, but R colour codes can be passed
<code>error_bar_colour</code>	the colour of the error bar
<code>point_size</code>	the point size of the plot
<code>error_bar_width</code>	the width of the displayed error bar
<code>h_line_color</code>	the colour of the horizontal line

Value

A list of the odds returned from logistic regression and a plot showing the odds

Examples

```
#We will use the cancer dataset to build a GLM model to predict cancer status
#this will detail whether the patient has a benign or malignant
library(mlbench)
library(caret)
library(tibble)
library(ggplot2)
library(OddsPlotty)
library(e1071)
library(ggthemes)

#Bring in the data
data("BreastCancer", package = "mlbench")
breast <- BreastCancer[complete.cases(BreastCancer), ]
breast <- breast[, -1]
head(breast, 10)
breast$class <- factor(breast$class)
for(i in 1:9) {
  breast[, i] <- as.numeric(as.character(breast[, i]))
}

#Train GLM model
glm_model <- train(Class ~ ., data = breast, method = "glm", family = "binomial")

#Visualise the data with OddsPlotty
plotty <- OddsPlotty::odds_plot(glm_model$finalModel, title = "Odds Plot")
plotty$odds_plot

#Extract underlying odds ratios
plotty$odds_data
```

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