

Package ‘predictoR’

April 30, 2022

Title Predictive Data Analysis System

Version 2.0.7

Description Perform a supervised data analysis on a database through a 'shiny' graphical interface. It includes methods such as K-Nearest Neighbors, Decision Trees, ADA Boosting, Extreme Gradient Boosting, Random Forest, Neural Networks, Deep Learning, Support Vector Machines and Bayesian Methods.

License GPL (>= 2)

Imports DT (>= 0.19), golem (>= 0.3.1), shiny (>= 1.7.1), rlang (>= 0.4.11), config (>= 0.3.1), xtable (>= 1.8-4), glmnet (>= 4.1-2), rpart (>= 4.1-15), colourpicker (>= 1.1.1), traineR (>= 1.6.2), shinyjs (>= 2.0.0), xgboost (>= 1.4.1.1), rpart.plot (>= 3.0.9), echarts4r (>= 0.4.2), shinyAce (>= 0.4.1), htmltools (>= 0.5.2), shinydashboard (>= 0.7.2), shinycustomloader (>= 0.9.0), shinydashboardPlus (>= 2.0.3)

Depends R (>= 4.1)

Encoding UTF-8

URL <https://www.promidat.com>

RoxygenNote 7.1.1

NeedsCompilation no

Author Oldemar Rodriguez [aut, cre],
Andres Navarro [aut, prg],
Joseline Quirós [aut, prg],
Diego Jiménez [ctb, prg]

Maintainer Oldemar Rodriguez <oldemar.rodriguez@ucr.ac.cr>

Repository CRAN

Date/Publication 2022-04-29 23:50:03 UTC

R topics documented:

datos.disyuntivos	2
dfnormal	3

e_coeff_landa	3
e_cor	4
e_global_gauge	5
e_histboxplot	6
e_histnormal	7
e_JS	8
e_posib_lambda	8
e_qq	9
e_rf_error	10
predictoR	10
run_app	11

Index	12
--------------	-----------

datos.disyuntivos *Create disjunctive columns to a data.frame.*

Description

Create disjunctive columns to a data.frame.

Usage

```
datos.disyuntivos(data, var)
```

Arguments

data	a data.frame object.
var	the column name to apply disjunctive code.

Value

data.frame

Author(s)

Diego Jimenez <diego.jimenez@promidat.com>

Examples

```
datos.disyuntivos(iris, "Species")
```

dfnormal	<i>Data.frame with normal test</i>
----------	------------------------------------

Description

Data.frame with normal test

Usage

```
dfnormal(data)
```

Arguments

data a data.frame object only with the numeric columns.

Value

data.frame

Author(s)

Diego Jimenez <diego.jimenez@promidat.com>

Examples

```
dfnormal(iris[, -5])
```

e_coeff_landa	<i>Coefficients and lambda</i>
---------------	--------------------------------

Description

Plot the coefficients and selected lambda of a glmnet model.

Usage

```
e_coeff_landa(model, category, sel.lambda = NULL, label = "Log Lambda")
```

Arguments

model a glmnet model.
category a category of the variable to be predicted.
sel.lambda the selected lambda.
label a character specifying the title to use on selected lambda tooltip.

Value

echarts4r plot

Author(s)

Joseline Quiros <joseline.quiros@promidat.com>

Examples

```
modelo <- traineR::train.glmnet(Species~, iris)
e_coeff_landa(modelo, 'setosa', log(modelo$lambda[1]))
```

e_cor

Correlation plot

Description

Correlation plot

Usage

```
e_cor(x, colors = c("#FF5733", "#F8F5F5", "#2E86C1"))
```

Arguments

- x a data.frame with correlation values.
- colors a vector of lenght 3 with color values.

Value

echarts4r plot

Author(s)

Diego Jimenez <diego.jimenez@promidat.com>

Examples

```
p <- round(cor(iris[, -5]), 3)
e_cor(p)
```

e_global_gauge *Gauge Plot*

Description

Gauge Plot

Usage

```
e_global_gauge(  
  value = 100,  
  label = "Label",  
  color1 = "#B5E391",  
  color2 = "#90C468"  
)
```

Arguments

value	a number specifying the value of the graph.
label	a character specifying the title to use on legend.
color1	a color for the gauge.
color2	a shadowColor for the gauge.

Value

echarts4r plot

Author(s)

Joseline Quiros <joseline.quiros@promidat.com>

Examples

```
e_global_gauge(87, "Global Precision")
```

e_histboxplot *Histogram + boxplot*

Description

Histogram + boxplot

Usage

```
e_histboxplot(  
  data,  
  var.name,  
  colorBar = "steelblue",  
  colorPoint = "red",  
  titulos = c("Minimo", "Primer Cuartil", "Mediana", "Tercer Cuartil", "Maximo")  
)
```

Arguments

<code>data</code>	a numeric column of a data.frame.
<code>var.name</code>	a character value specifying the name of the variable.
<code>colorBar</code>	a color for the bars.
<code>colorPoint</code>	a color for the points.
<code>titulos</code>	a character vector of length 5 specifying the titles to use on legend.

Value

echarts4r plot

Author(s)

Diego Jimenez <diego.jimenez@promidat.com>

Examples

```
e_histboxplot(iris$Sepal.Width, "Sepal.Width")
```

e_histnormal	<i>Normal plot</i>
--------------	--------------------

Description

Normal plot

Usage

```
e_histnormal(  
  data,  
  colorbar = "steelblue",  
  colorline = "gray",  
  nombres = c("Histograma", "Curva Normal")  
)
```

Arguments

data	a numeric column of a data.frame.
colorbar	a color for the bars.
colorline	a color for the line.
nombres	a character vector of length 2 specifying the titles to use on legend.

Value

echarts4r plot

Author(s)

Diego Jimenez <diego.jimenez@promidat.com>

Examples

```
e_histnormal(iris$Sepal.Length)
```

e_JS*Eval character vectors to JS code*

Description

Eval character vectors to JS code

Usage

```
e_JS(...)
```

Arguments

...	character vectors to evaluate
-----	-------------------------------

Author(s)

Joseline Quiros <joseline.quiros@promidat.com>

Examples

```
e_JS('5 * 3')
```

e_posib_lambda*Possible lambda*

Description

Possible lambda

Usage

```
e_posib_lambda(
  cv.glm,
  labels = c("Valor Superior", "Valor Inferior", "lambda")
)
```

Arguments

cv.glm	a cv.glmnet model.
labels	a character vector of length 3 specifying the titles to use on legend.

Value

echarts4r plot

Author(s)

Joseline Quiros <joseline.quiros@promidat.com>

Examples

```
x      <- model.matrix(Species~, iris)[, -1]
y      <- iris[, 'Species']
cv.glm <- glmnet::cv.glmnet(x, y, standardize = TRUE, alpha = 1, family = 'multinomial')
e_posib_lambda(cv.glm)
```

e_qq

Qplot + Qline

Description

Qplot + Qline

Usage

```
e_qq(data, colorpoint = "steelblue", colorline = "gray")
```

Arguments

data	a numeric column of a data.frame.
colorpoint	a color for the points.
colorline	a color for the line.

Value

echarts4r plot

Author(s)

Diego Jimenez <diego.jimenez@promidat.com>

Examples

```
e_qq(iris$Sepal.Length)
```

e_rf_error*Error Evolution***Description**

Error Evolution

Usage

```
e_rf_error(model)
```

Arguments

model a random forest model.

Value

echarts4r plot

Author(s)

Joseline Quiros <joseline.quiros@promidat.com>

Examples

```
model <- traineR::train.randomForest(Species~, iris, mtry = 2, ntree = 20)
e_rf_error(model)
```

predictoR*Predictive Data Analysis System***Description**

Perform a supervised data analysis on a database through a 'shiny' graphical interface. It includes methods such as K-Nearest Neighbors, Decision Trees, ADA Boosting, Extreme Gradient Boosting, Random Forest, Neural Networks, Deep Learning, Support Vector Machines and Bayesian Methods.

Details

Package:	predictoR
Type:	Package
Version:	2.0.1
Date:	2021-06-11
License:	GPL (>=2)

Author(s)

Oldemar Rodriguez Rojas

Maintainer: Oldemar Rodriguez Rojas <oldemar.rodriguez@ucr.ac.cr>

`run_app`

Run the Shiny Application

Description

Run the Shiny Application

Usage

`run_app(...)`

Arguments

`...` A series of options to be used inside the app.

Index

```
* package
  predictoR, 10

  datos.disyuntivos, 2
  dfnormal, 3

  e_coeff_landa, 3
  e_cor, 4
  e_global_gauge, 5
  e_histboxplot, 6
  e_histnormal, 7
  e_JS, 8
  e_posib_lambda, 8
  e_qq, 9
  e_rf_error, 10

  predictoR, 10

  run_app, 11
```