Package ‘iBreakDown’

October 13, 2022

Title Model Agnostic Instance Level Variable Attributions

Version 2.0.1

Description Model agnostic tool for decomposition of predictions from black boxes.
Supports additive attributions and attributions with interactions.
The Break Down Table shows contributions of every variable to a final prediction.
The Break Down Plot presents variable contributions in a concise graphical way.
This package works for classification and regression models.
It is an extension of the 'breakDown' package (Staniak and Biecek 2018) <doi:10.32614/RJ-2018-072>,
with new and faster strategies for orderings.
It supports interactions in explanations and has interactive visuals (implemented with 'D3.js' library).
The methodology behind is described in the 'iBreakDown' article (Gosiewska and Biecek 2019) <arXiv:1903.11420>
This package is a part of the 'DrWhy.AI' universe (Biecek 2018) <arXiv:1806.08915>.

Depends R (>= 3.5)

License GPL-3

Encoding UTF-8

RoxygenNote 7.1.1

Imports ggplot2

Suggests DALEX, knitr, rmarkdown, randomForest, e1071, ranger, nnet, testthat, r2d3, jsonlite, covr

VignetteBuilder knitr

URL https://ModelOriented.github.io/iBreakDown/,
https://github.com/ModelOriented/iBreakDown

BugReports https://github.com/ModelOriented/iBreakDown/issues

NeedsCompilation no

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Description

This function finds Variable Attributions via Sequential Variable Conditioning. It calls either `local_attributions` for additive attributions or `local_interactions` for attributions with interactions.

Usage

```r
break_down(x, ..., interactions = FALSE)
```

```r
## S3 method for class 'explainer'
break_down(x, new_observation, ..., interactions = FALSE)
```

```r
## Default S3 method:
break_down(
  x,
  data,
  predict_function = predict,
  new_observation,
  keep_distributions = FALSE,
  order = NULL,
  label = class(x)[1],
  ..., interactions = FALSE
)
```
Arguments

- `x`: an explainer created with function `explain` or a model.
- `...`: parameters passed to local_\* functions.
- `interactions`: shall interactions be included?
- `new_observation`: a new observation with columns that correspond to variables used in the model.
- `data`: validation dataset, will be extracted from x if it is an explainer.
- `predict_function`: predict function, will be extracted from x if it’s an explainer.
- `keep_distributions`: if TRUE, then distribution of partial predictions is stored and can be plotted with the generic plot().
- `order`: if not NULL, then it will be a fixed order of variables. It can be a numeric vector or vector with names of variables.
- `label`: name of the model. By default it is extracted from the ‘class’ attribute of the model.

Value

an object of the `break_down` class.

References


See Also

`localattributions`, `local_interactions`

Examples

```r
library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare,
                          data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
                               data = titanic_imputed,
                               y = titanic_imputed$survived,
                               label = "glm")

bd_glm <- break_down(explain_titanic_glm, titanic_imputed[1, ])
bd_glm
plot(bd_glm, max_features = 3)
```

## Not run:
```r
## Not run:
```
library("randomForest")
set.seed(1313)
# example with interaction
# classification for HR data
model <- randomForest(status ~ . , data = HR)
new_observation <- HR_test[1,]

explainer_rf <- explain(model,
data = HR[1:1000,1:5])

bd_rf <- break_down(explainer_rf,
new_observation)

head(bd_rf)
plot(bd_rf)

## End(Not run)

---

**break_down_uncertainty**

*Explanation Level Uncertainty of Sequential Variable Attribution*

---

**Description**

This function calculates the break down algorithm for \( B \) random orderings. Then it calculates the distribution of attributions for these different orderings. Note that the `shap()` function is just a simplified interface to the `break_down_uncertainty()` function with a default value set to \( B = 25 \).

**Usage**

```r
break_down_uncertainty(x, ..., keep_distributions = TRUE, B = 10)
```

### S3 method for class 'explainer'

```r
break_down_uncertainty(
x, new_observation, 
..., keep_distributions = TRUE, 
B = 10 
)
```

### Default S3 method:

```r
break_down_uncertainty(
x, data, 
predict_function = predict, 
new_observation, 
label = class(x)[1], 
..., 
```
break_down_uncertainty

    path = NULL,
    keep_distributions = TRUE,
    B = 10
)

shap(x, ..., B = 25)

Arguments

x an explainer created with function explain or a model.

... other parameters.

keep_distributions if TRUE then we will keep distribution for predicted values. It’s needed by the describe function.

B number of random paths

ew_observation a new observation with columns that correspond to variables used in the model.

data validation dataset, will be extracted from x if it is an explainer.

predict_function predict function, will be extracted from x if it is an explainer.

label name of the model. By default it’s extracted from the ‘class’ attribute of the model.

path if specified, then this path will be highlighted on the plot. Use average in order to show an average effect

Value

an object of the break_down_uncertainty class.

References


See Also

break_down, local_attributions

Examples

library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare,
                         data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
                               data = titanic_imputed,
                               y = titanic_imputed$survived,
                               B = 25)
label = "glm")

# there is no explanation level uncertainty linked with additive models
bd_glm <- break_down_uncertainty(explain_titanic_glm, titanic_imputed[1, ])
bd_glm
plot(bd_glm)

## Not run:
## Not run:
library("randomForest")
set.seed(1313)
model <- randomForest(status ~ ., data = HR)
new_observation <- HR_test[1,]
explainer_rf <- explain(model,
  data = HR[1:1000, 1:5])

bd_rf <- break_down_uncertainty(explainer_rf,
  new_observation)
bd_rf
plot(bd_rf)

# example for regression - apartment prices
# here we do not have interactions
model <- randomForest(m2.price ~ ., data = apartments)
explainer_rf <- explain(model,
  data = apartments_test[1:1000, 2:6],
  y = apartments_test$m2.price[1:1000])

bd_rf <- break_down_uncertainty(explainer_rf, apartments_test[1,])
bd_rf
plot(bd_rf)

bd_rf <- break_down_uncertainty(explainer_rf, apartments_test[1,], path = 1:5)
plot(bd_rf)

bd_rf <- break_down_uncertainty(explainer_rf, apartments_test[1,], path = c("floor", "no.rooms", "district",
  "construction.year", "surface"))
plot(bd_rf)

bd <- break_down(explainer_rf, apartments_test[1,])
plot(bd)

s <- shap(explainer_rf, apartments_test[1,])
plot(s)

## End(Not run)
describe Generates Textual Explanations for Predictive Models

Description

Generic function describe generates natural language explanations based on break_down and shap explanations, what enhances their interpretability.

Usage

describe(x, nonsignificance_treshold = 0.15, ...)

## S3 method for class 'break_down'
describe(
  x,
  nonsignificance_treshold = 0.15,
  ..., 
  label = NULL,
  short_description = FALSE,
  display_values = FALSE,
  display_numbers = FALSE,
  display_distribution_details = FALSE,
  display_shap = FALSE
)

## S3 method for class 'break_down_uncertainty'
describe(
  x,
  nonsignificance_treshold = 0.15,
  ..., 
  label = NULL,
  short_description = FALSE,
  display_values = FALSE,
  display_numbers = FALSE,
  display_distribution_details = FALSE,
  display_shap = FALSE
)

Arguments

x an explanation created with break_down or shap
nonsignificance_treshold a numeric specifying a threshold for variable importance
... other arguments
label a character string describing model’s prediction
short_description  
a boolean, returns a short description

display_values  
a boolean, displays variables’ values

display_numbers  
a boolean, displays a description containing numerical values

display_distribution_details  
a boolean, displays details about the distribution of model’s predictions

display_shap  
a boolean, adds information about variables’ average contribution. Use only with shap explanation.

Details

Function describe generates a textual explanations by extracting information from a break_down or shap explanation. It makes an argument justifying why the model’s prediction is lower or higher, than it’s average prediction. The description consists of an introduction, argumentation and summary making use from the claim, support, evidence argumentation structure, as recomended for the World Universities Debating style.

The function first selects one of four different scenarios, due to nonsignificance_treshold. The chosen scenario can be one of the following: 1. Model’s prediction for the selected instance is significantly higher than the average prediction. 2. Model’s prediction is significantly lower. 3. Model’s prediction is close to it’s average prediction, however there are significant variables counteracting with each other 4. Model’s prediction is close to it’s average prediction and all the variables are rather nonsignificant. Then an explanation due to the chosen scenario is generated.

Value

A character string of textual explanation

Examples

library("DALEX")
library("randomForest")
library("iBreakDown")

  titanic <- na.omit(titanic)
  model_titanic_rf <- randomForest(survived == "yes" ~ gender + age + class + embarked + fare + sibsp + parch, data = titanic)

  explain_titanic_rf <- explain(model_titanic_rf,
                                data = titanic[,-9],
                                y = titanic$survived == "yes",
                                label = "Random Forest v7")

  bd_explanation <- break_down(explain_titanic_rf, titanic[1, ], keep_distributions = TRUE)
  plot(bd_explanation)

  description <- describe(bd_explanation,
                          label = "the passanger will survive with probability",
                          short_description = FALSE,
                          display_values = TRUE,
library("DALEX")
library("iBreakDown")
titanic <- na.omit(titanic)
model_titanic_glm <- glm(titanic$survived == "yes" ~ age + gender + class + fare + sibsp,
                         data = titanic[, -9], family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
                               data = titanic[, -9],
                               y = titanic$survived == "yes",
                               label = "glm")
passanger <- titanic[1, -9]
shap_glm <- shap(explain_titanic_glm, passanger)
plot(shap_glm)

describe(shap_glm,
         label = "the selected passanger survives with probability",
         display_shap = TRUE,
         display_numbers = TRUE)

local_attributions

Model Agnostic Sequential Variable attributions

Description
This function finds Variable attributions via Sequential Variable Conditioning. The complexity of this function is O(2^p). This function works in a similar way to step-up and step-down greedy approximations in function \texttt{break_down}. The main difference is that in the first step the order of variables is determined. And in the second step the impact is calculated.

Usage
local_attributions(x, ...)

## S3 method for class 'explainer'
local_attributions(x, new_observation, keep_distributions = FALSE, ...)

## Default S3 method:
local_attributions(
  x,
  data,
  predict_function = predict,
  new_observation,
  label = class(x)[1],
  keep_distributions = FALSE,
Arguments

x an explainer created with function explain or a model.
...
other parameters.
new_observation a new observation with columns that correspond to variables used in the model.
keep_distributions if TRUE, then distribution of partial predictions is stored and can be plotted with the generic plot().
data validation dataset, will be extracted from x if it is an explainer.
predict_function predict function, will be extracted from x if it is an explainer.
label name of the model. By default it’s extracted from the 'class' attribute of the model.
order if not NULL, then it will be a fixed order of variables. It can be a numeric vector or vector with names of variables.

Value

an object of the break_down class.

References


See Also

break_down, local_interactions

Examples

library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare, data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm, data = titanic_imputed, y = titanic_imputed$survived, label = "glm")

bd_glm <- local_attributions(explain_titanic_glm, titanic_imputed[1, ])
bd_glm
plot(bd_glm, max_features = 3)
## Not run:
## Not run:
library("randomForest")
set.seed(1313)
# example with interaction
# classification for HR data
model <- randomForest(status ~ . , data = HR)
new_observation <- HR_test[1,]

explainer_rf <- explain(model,
data = HR[1:1000,1:5])

bd_rf <- local_attributions(explainer_rf,
new_observation)

bd_rf
plot(bd_rf)
plot(bd_rf, baseline = 0)

# example for regression - apartment prices
# here we do not have interactions
model <- randomForest(m2.price ~ . , data = apartments)
explainer_rf <- explain(model,
data = apartments_test[1:1000,2:6],
y = apartments_test$m2.price[1:1000])

bd_rf <- local_attributions(explainer_rf,
apartments_test[1,])

bd_rf
plot(bd_rf, digits = 1)

bd_rf <- local_attributions(explainer_rf,
apartments_test[1,],
keep_distributions = TRUE)
plot(bd_rf, plot_distributions = TRUE)

## End(Not run)

---

**local_interactions**  
*Model Agnostic Sequential Variable Attributions with Interactions*

### Description

This function implements decomposition of model predictions with identification of interactions. The complexity of this function is $O(2^p)$ for additive models and $O(2^p p^2)$ for interactions. This function works in a similar way to step-up and step-down greedy approximations in function `break_down()`. The main difference is that in the first step the order of variables and interactions is determined. And in the second step the impact is calculated.
Usage

local_interactions(x, ...)

## S3 method for class 'explainer'
local_interactions(x, new_observation, keep_distributions = FALSE, ...)

## Default S3 method:
local_interactions(
  x,
  data,
  predict_function = predict,
  new_observation,
  label = class(x)[1],
  keep_distributions = FALSE,
  order = NULL,
  interaction_preference = 1,
  ...
)

Arguments

x an explainer created with function explain or a model.
...
other parameters.
new_observation a new observation with columns that correspond to variables used in the model.
keep_distributions if TRUE, then the distribution of partial predictions is stored in addition to the average.
data validation dataset, will be extracted from x if it's an explainer.
predict_function predict function, will be extracted from x if it's an explainer.
label character - the name of the model. By default it's extracted from the 'class' attribute of the model.
order if not NULL, then it will be a fixed order of variables. It can be a numeric vector or vector with names of variables/interactions.
interaction_preference an integer specifying which interactions will be present in an explanation. The larger the integer, the more frequently interactions will be presented.

Value

an object of the break_down class.

References

local_interactions

See Also

break_down, local_attributions

Examples

library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare,
data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
data = titanic_imputed,
y = titanic_imputed$survived,
label = "glm")

bd_glm <- local_interactions(explain_titanic_glm, titanic_imputed[1, ],
interaction_preference = 500)
bd_glm
plot(bd_glm, max_features = 2)

## Not run:
library("randomForest")
# example with interaction
# classification for HR data
model <- randomForest(status ~ . , data = HR)
new_observation <- HR_test[1,]
explainer_rf <- explain(model,
data = HR[1:1000,1:5])

bd_rf <- local_interactions(explainer_rf,
new_observation)

bd_rf
plot(bd_rf)

# example for regression - apartment prices
# here we do not have intreactions
model <- randomForest(m2.price ~ . , data = apartments)
explainer_rf <- explain(model,
data = apartments_test[1:1000,2:6],
y = apartments_test$m2.price[1:1000])

new_observation <- apartments_test[1,]

bd_rf <- local_interactions(explainer_rf,
new_observation, keep_distributions = TRUE)

bd_rf
plot(bd_rf)
plot(bd_rf, plot_distributions = TRUE)
plot.break_down

## End(Not run)

---

### plot.break_down

#### Plot Generic for Break Down Objects

**Description**

Displays a waterfall break down plot for objects of `break_down` class.

**Usage**

```r
## S3 method for class 'break_down'
plot(
x, ...
baseline = NA,
max_features = 10,
min_max = NA,
vcolors = DALEX::colors_breakdown_drwhy(),
digits = 3,
rounding_function = round,
add_contributions = TRUE,
shift_contributions = 0.05,
plot_distributions = FALSE,
vnames = NULL,
title = "Break Down profile",
subtitle = "",
max_vars = NULL
)
```

**Arguments**

- `x` an explanation created with `break_down`
- `...` other parameters.
- `baseline` if numeric then veritical line starts in baseline.
- `max_features` maximal number of features to be included in the plot. default value is 10.
- `min_max` a range of OX axis. By default NA, therefore it will be extracted from the contributions of x. But it can be set to some constants, useful if these plots are to be used for comparisons.
- `vcolors` If NA (default), DrWhy colors are used.
- `digits` number of decimal places (`round`) or significant digits (`signif`) to be used. See the `rounding_function` argument.
plot.break_down

rounding_function
a function to be used for rounding numbers. This should be signif which keeps a specified number of significant digits or round (which is default) to have the same precision for all components.

add_contributions
if TRUE, variable contributions will be added to the plot

shift_contributions
number describing how much labels should be shifted to the right, as a fraction of range. By default equal to \(0.05\).

plot_distributions
if TRUE then distributions of conditional proportions will be plotted. This requires keep_distributions=TRUE in the break_down, local_attributions, or local_interactions.

vnames
a character vector, if specified then will be used as labels on OY axis. By default NULL

title
a character. Plot title. By default "Break Down profile".

subtitle
a character. Plot subtitle. By default "".

max_vars
alias for the max_features parameter.

Value
a ggplot2 object.

References

Examples
library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare,
data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
data = titanic_imputed,
y = titanic_imputed$survived,
label = "glm")

bd_glm <- break_down(explain_titanic_glm, titanic_imputed[1, ])
bd_glm
plot(bd_glm, max_features = 3)
plot(bd_glm, max_features = 3,
    vnames = c("average","+ male","+ young","+ cheap ticket","+ other factors","final"))

## Not run:
## Not run:
library("randomForest")
```r
set.seed(1313)
# example with interaction
# classification for HR data
model <- randomForest(status ~ ., data = HR)
new_observation <- HR_test[1,]

explainer_rf <- explain(model,
                         data = HR[1:1000,1:5])

bd_rf <- local_attributions(explainer_rf,
                             new_observation)

plot(bd_rf)
plot(bd_rf, baseline = 0)
plot(bd_rf, min_max = c(0,1))

bd_rf <- local_attributions(explainer_rf,
                             new_observation,
                             keep_distributions = TRUE)

plot(bd_rf, plot_distributions = TRUE)

bd_rf <- local_interactions(explainer_rf,
                             new_observation,
                             keep_distributions = TRUE)

plot(bd_rf)
plot(bd_rf, plot_distributions = TRUE)

# example for regression - apartment prices
# here we do not have interactions
model <- randomForest(m2.price ~ ., data = apartments)
explainer_rf <- explain(model,
                         data = apartments_test[1:1000,2:6],
                         y = apartments_test$m2.price[1:1000])

bd_rf <- local_attributions(explainer_rf,
                             apartments_test[1,])

plot(bd_rf, digits = 1)
plot(bd_rf, digits = 1, baseline = 0)

bd_rf <- local_attributions(explainer_rf,
                             apartments_test[1,],
                             keep_distributions = TRUE)

plot(bd_rf, plot_distributions = TRUE)

bd_rf <- local_interactions(explainer_rf,
                             new_observation = apartments_test[1,],
                             keep_distributions = TRUE)

bd_rf
```
plot(bd_rf)
plot(bd_rf, plot_distributions = TRUE)

## End(Not run)

plot.break_down_uncertainty

Plot Generic for Break Down Uncertainty Objects

Description

Plot Generic for Break Down Uncertainty Objects

Usage

## S3 method for class 'break_down_uncertainty'
plot(
  x,
  ..., vcolors = DALEX::colors_breakdown_drwhy(),
  show_boxplots = TRUE,
  max_features = 10,
  max_vars = NULL
)

Arguments

x
  an explanation created with \texttt{break_down_uncertainty}

... other parameters.

vcolors
  If NA (default), DrWhy colors are used.

show_boxplots
  logical if TRUE (default) boxplot will be plotted to show uncertainty of attribu-
  tions

max_features
  maximal number of features to be included in the plot. By default it’s 10.

max_vars
  alias for the \texttt{max_features} parameter.

Value

a ggplot2 object.

References

drwhy.ai
Examples

```r
library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare,
data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
data = titanic_imputed,
y = titanic_imputed$survived,
label = "glm")

sh_glm <- shap(explain_titanic_glm, titanic_imputed[1,])
sh_glm
plot(sh_glm)
```

```r
## Not run:
## Not run:
library("randomForest")
set.seed(1313)
model <- randomForest(status ~ . , data = HR)
new_observation <- HR_test[1,]
explainer_rf <- explain(model,
data = HR[1:1000,1:5])

bd_rf <- break_down_uncertainty(explainer_rf,
new_observation,
path = c(3,2,4,1,5),
show_boxplots = FALSE)
bd_rf
plot(bd_rf, max_features = 3)
```

```r
# example for regression - apartment prices
# here we do not have interactions
model <- randomForest(m2.price ~ . , data = apartments)
explainer_rf <- explain(model,
data = apartments_test[1:1000,2:6],
y = apartments_test$m2.price[1:1000])

bd_rf <- break_down_uncertainty(explainer_rf,
apartments_test[1,],
path = c("floor", "no.rooms", "district",
"construction.year", "surface"))

bd_rf
plot(bd_rf)
```

```r
bd_rf <- shap(explainer_rf,
apartments_test[1,])
bd_rf
plot(bd_rf)
```
plotD3

Plot Break Down Objects in D3 with r2d3 package.

Description

Plots waterfall break down for objects of the `break_down` class.

Usage

```r
plotD3(x, ...) 
## S3 method for class 'break_down'
plotD3(
  x,
  ..., 
  baseline = NA,
  max_features = 10,
  digits = 3,
  rounding_function = round,
  bar_width = 12,
  margin = 0.2,
  scale_height = FALSE,
  min_max = NA,
  vcolors = NA,
  chart_title = NA,
  time = 0,
  max_vars = NULL,
  reload = FALSE
)
```

Arguments

- `x` an explanation created with `break_down`
- `...` other parameters.
- `baseline` if numeric then vertical line will start in baseline.
- `max_features` maximal number of features to be included in the plot. By default it's 10.
- `digits` number of decimal places (`round`) or significant digits (`signif`) to be used. See the `rounding_function` argument.
- `rounding_function` a function to be used for rounding numbers. This should be `signif` which keeps a specified number of significant digits or `round` (which is default) to have the same precision for all components.
bar_width  width of bars in px. By default it’s 12px
margin     extend x axis domain range to adjust the plot. Usually value between 0.1 and 0.3, by default it’s 0.2
scale_height if TRUE, the height of the plot scales with window size
min_max    a range of OX axis. By default NA therefore will be extracted from the contributions of x. But can be set to some constants, useful if these plots are used for comparisons.
vcolors    If NA (default), DrWhy colors are used.
chart_title a character. Set custom title
time       in ms. Set the animation length
max_vars   alias for the max_features parameter.
reload     Reload the plot on resize. By default it’s FALSE.

Value

a r2d3 object.

References


Examples

```r
library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare, 
data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
data = titanic_imputed,
y = titanic_imputed$survived,
label = "glm")

bd_glm <- local_attributions(explain_titanic_glm, titanic_imputed[1, ])
bd_glm
plotD3(bd_glm)

## Not run:
## Not run:
library("randomForest")

m_rf <- randomForest(status ~ . , data = HR[2:2000,])
new_observation <- HR_test[1,]
new_observation
p_fun <- function(object, newdata){predict(object, newdata=newdata, type = "prob")}
bd_rf <- local_attributions(m_rf,
```
```r
plotD3.shap

```data = HR_test,
  new_observation = new_observation,
  predict_function = p_fun)

```bd_rf
plotD3(bd_rf)

```## End(Not run)

---------------------------------------------------------------------
plotD3.shap  
Plot Shap (Break Down Uncertainty) Objects in D3 with r2d3 package.

Description

Plots Shapley values.

Usage

```## S3 method for class 'shap'
plotD3(
  x,
  ..., baseline = NA,
  max_features = 10,
  digits = 3,
  rounding_function = round,
  bar_width = 12,
  margin = 0.2,
  scale_height = FALSE,
  min_max = NA,
  vcolors = NA,
  chart_title = NA,
  time = 0,
  max_vars = NULL,
  reload = FALSE
)
```

Arguments

- `x` an explanation created with `shap`
- `...` other parameters.
- `baseline` if numeric then vertical line will start in baseline.
- `max_features` maximal number of features to be included in the plot. By default it's 10.
- `digits` number of decimal places (`round`) or significant digits (`signif`) to be used. See the `rounding_function` argument.
rounding_function

a function to be used for rounding numbers. This should be signif which keeps a specified number of significant digits or round (which is default) to have the same precision for all components.

bar_width

width of bars in px. By default it's 12px

margin

extend x axis domain range to adjust the plot. Usually value between 0.1 and 0.3, by default it's 0.2

scale_height

if TRUE, the height of the plot scales with window size.

min_max

a range of OX axis. By default NA therefore will be extracted from the contributions of x. But can be set to some constants, useful if these plots are used for comparisons.

vcolors

If NA (default), DrWhy colors are used.

chart_title

a character. Set custom title

time

in ms. Set the animation length

max_vars

alias for the max_features parameter.

reload

Reload the plot on resize. By default it’s FALSE.

Value

a r2d3 object.

References


Examples

library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare, data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm, data = titanic_imputed, y = titanic_imputed$survived, label = "glm")

s_glm <- shap(explain_titanic_glm, titanic_imputed[1, ])
s_glm
plotD3(s_glm)

## Not run:
## Not run:
library("randomForest")
HR_small <- HR[2:500,]
m_rf <- randomForest(status ~. , data = HR_small)
new_observation <- HR_test[1,]
new_observation

p_fun <- function(object, newdata){predict(object, newdata=newdata, type = "prob")}

s_rf <- shap(m_rf,
    data = HR_small[-6],
    new_observation = new_observation,
    predict_function = p_fun)

plotD3(s_rf, time = 500)

## End(Not run)

---

print.break_down

Print Generic for Break Down Objects

Description
Print Generic for Break Down Objects

Usage
## S3 method for class 'break_down'
print(x, ..., digits = 3, rounding_function = round)

Arguments
- x: an explanation created with `break_down`
- ...: other parameters.
- digits: number of decimal places (round) or significant digits (signif) to be used. See the rounding_function argument.
- rounding_function: a function to be used for rounding numbers. This should be `signif` which keeps a specified number of significant digits or `round` (which is default) to have the same precision for all components.

Value
- a data frame

References
print.break_down_description

*Print Generic for Break Down Objects*

**Description**
Print Generic for Break Down Objects

**Usage**
```r
## S3 method for class 'break_down_description'
print(x, ...)
```

**Arguments**
- `x` a description of `break_down_description` class.
- `...` other parameters.

**Value**
a character

**References**

print.break_down_uncertainty

*Print Generic for Break Down Uncertainty Objects*

**Description**
Print Generic for Break Down Uncertainty Objects

**Usage**
```r
## S3 method for class 'break_down_uncertainty'
print(x, ...)
```

**Arguments**
- `x` an explanation created with `break_down_uncertainty`
- `...` other parameters.
Value

a data frame.

References


Examples

library("DALEX")
library("iBreakDown")
set.seed(1313)
model_titanic_glm <- glm(survived ~ gender + age + fare,
  data = titanic_imputed, family = "binomial")
explain_titanic_glm <- explain(model_titanic_glm,
  data = titanic_imputed,
  y = titanic_imputed$survived,
  label = "glm")

bd_glm <- break_down_uncertainty(explain_titanic_glm, titanic_imputed[1, ])
bd_glm
plot(bd_glm)

## Not run:

library("randomForest")
set.seed(1313)
model <- randomForest(status ~ . , data = HR)
new_observation <- HR_test[1,]
explainer_rf <- explain(model,
  data = HR[1:1000,1:5],
  y = HR$status[1:1000],
  verbose = FALSE)

bd_rf <- break_down_uncertainty(explainer_rf, new_observation)
bd_rf

# example for regression - apartment prices
# here we do not have interactions
model <- randomForest(m2.price ~ . , data = apartments)
explainer_rf <- explain(model,
  data = apartments_test[1:1000,2:6],
  y = apartments_test$m2.price[1:1000])

bd_rf <- break_down_uncertainty(explainer_rf, apartments_test[1,])
bd_rf

## End(Not run)
Index

break_down, 2, 5, 7–10, 13–15, 19, 23
break_down_uncertainty, 4, 17, 24

describe, 7
explain, 3, 5, 10, 12

local_attributions, 2, 3, 5, 9, 13, 15
local_interactions, 2, 3, 10, 11, 15

plot.break_down, 14
plot.break_down_uncertainty, 17
plotD3, 19
plotD3.shap, 21
print.break_down, 23
print.break_down_description, 24
print.break_down_uncertainty, 24

round, 14, 15, 19, 21–23

shap, 7, 8, 21
shap(break_down_uncertainty), 4
signif, 14, 15, 19, 21–23