

Package ‘ceterisParibus’

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Title Ceteris Paribus Profiles

Version 0.3.0

Description Ceteris Paribus Profiles (What-If Plots) are designed to present model responses around selected points in a feature space.
For example around a single prediction for an interesting observation.
Plots are designed to work in a model-agnostic fashion, they are working for any predictive Machine Learning model and allow for model comparisons.
Ceteris Paribus Plots supplement the Break Down Plots from 'breakDown' package.

Depends R (>= 3.3), ggplot2, gower

Suggests randomForest, ggiraph, e1071, testthat, rpart

Imports DALEX, knitr

License GPL-2

Encoding UTF-8

LazyData true

VignetteBuilder knitr

URL <https://pbiecek.github.io/ceterisParibus/>

BugReports <https://github.com/pbiecek/ceterisParibus/issues>

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`+.plot_ceteris_paribus_explainer`
Add More Layers to a Ceteris Paribus Plot

Description

Add More Layers to a Ceteris Paribus Plot

Usage

```
## S3 method for class 'plot_ceteris_paribus_explainer'
e1 + e2
```

Arguments

e1 An object of class 'plot_ceteris_paribus_explainer'.

e2 A plot component

calculate_profiles *Calculate Ceteris Paribus Profiles*

Description

This function calculates ceteris paribus profiles, i.e. series of predictions from a model calculated for observations with altered single coordinate.

Usage

```
calculate_profiles(data, variable_splits, model, predict_function = predict,
  ...)
```

Arguments

data	set of observations. Profile will be calculated for every observation (every row)
variable_splits	named list of vectors. Elements of the list are vectors with points in which profiles should be calculated. See an example for more details.
model	a model that will be passed to the predict_function
predict_function	function that takes data and model and returns numeric predictions. Note that the ... arguments will be passed to this function.
...	other parameters that will be passed to the predict_function

Details

Note that calculate_profiles function is S3 generic. If you want to work on non standard data sources (like H2O ddf, external databases) you should overload it.

Value

a data frame with profiles for selected variables and selected observations

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)
apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)
vars <- c("construction.year", "surface", "floor", "no.rooms", "district")
variable_splits <- calculate_variable_splits(apartments, vars)
new_apartment <- apartmentsTest[1:10, ]
profiles <- calculate_profiles(new_apartment, variable_splits,
  apartments_rf_model)
```

```

profiles

# only subset of observations
small_apartments <- select_sample(apartmentsTest, n = 10)
small_apartments
small_profiles <- calculate_profiles(small_apartments, variable_splits,
                                   apartments_rf_model)

small_profiles

# neighbors for a selected observation
new_apartment <- apartments[1, 2:6]
small_apartments <- select_neighbours(apartmentsTest, new_apartment, n = 10)
small_apartments
small_profiles <- calculate_profiles(small_apartments, variable_splits,
                                   apartments_rf_model)

new_apartment
small_profiles

## End(Not run)

```

calculate_variable_splits

Calculate Split Points for Selected Variables

Description

This function calculate candidate splits for each selected variable. For numerical variables splits are calculated as percentiles (in general uniform quantiles of the length grid_points). For all other variables splits are calculated as unique values.

Usage

```
calculate_variable_splits(data, variables = colnames(data),
                          grid_points = 101)
```

Arguments

data	validation dataset. Is used to determine distribution of observations.
variables	names of variables for which splits shall be calculated
grid_points	number of points used for response path

Details

Note that calculate_variable_splits function is S3 generic. If you want to work on non standard data sources (like H2O ddf, external databases) you should overload it.

Value

A named list with splits for selected variables

Examples

```

library("DALEX")
## Not run:
library("randomForest")
set.seed(59)
apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
                                   no.rooms + district, data = apartments)
vars <- c("construction.year", "surface", "floor", "no.rooms", "district")
calculate_variable_splits(apartments, vars)

## End(Not run)

```

ceteris_paribus

Ceteris Paribus Explainer

Description

This function calculate ceteris paribus profiles for selected data points.

Usage

```

ceteris_paribus(explainer, observations, y = NULL, variable_splits = NULL,
                variables = NULL, grid_points = 101)

```

Arguments

explainer	a model to be explained, preprocessed by function ‘DALEX::explain()’.
observations	set of observation for which profiles are to be calculated
y	true labels for ‘observations’. If specified then will be added to ceteris paribus plots.
variable_splits	named list of splits for variables, in most cases created with ‘calculate_variable_splits()’. If NULL then it will be calculated based on validation data available in the ‘explainer’.
variables	names of variables for which profiles shall be calculated. Will be passed to ‘calculate_variable_splits()’. If NULL then all variables from the validation data will be used.
grid_points	number of points for profile. Will be passed to ‘calculate_variable_splits()’.

Value

An object of the class ‘ceteris_paribus_explainer’. It’s a data frame with calculated average responses.

Examples

```

library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

apartments_small <- select_sample(apartmentsTest, 10)

cp_rf <- ceteris_paribus(explainer_rf, apartments_small)
cp_rf

cp_rf <- ceteris_paribus(explainer_rf, apartments_small, y = apartments_small$m2.price)
cp_rf

## End(Not run)

```

ceteris_paribus_layer *Add Layer to the Ceteris Paribus Plot*

Description

Function ‘ceteris_paribus_layer()’ adds a layer to a plot created with ‘plot.ceteris_paribus_explainer()’ plots. Various parameters help to decide what should be plotted, profiles, aggregated profiles, points or rugs.

Usage

```

ceteris_paribus_layer(x, ..., size = 1, alpha = 0.3, color = "black",
  size_points = 2, alpha_points = 1, color_points = color,
  size_rugs = 0.5, alpha_rugs = 1, color_rugs = color,
  size_residuals = 1, alpha_residuals = 1, color_residuals = color,
  only_numerical = TRUE, show_profiles = TRUE, show_observations = TRUE,
  show_rugs = FALSE, show_residuals = FALSE, aggregate_profiles = NULL,
  as.gg = FALSE, facet_ncol = NULL, selected_variables = NULL,
  init_plot = FALSE)

```

Arguments

x	a ceteris paribus explainer produced with function ‘ceteris_paribus()’
...	other explainers that shall be plotted together
size	a numeric. Size of lines to be plotted

alpha	a numeric between 0 and 1. Opacity of lines
color	a character. Either name of a color or name of a variable that should be used for coloring
size_points	a numeric. Size of points to be plotted
alpha_points	a numeric between 0 and 1. Opacity of points
color_points	a character. Either name of a color or name of a variable that should be used for coloring
size_rugs	a numeric. Size of rugs to be plotted
alpha_rugs	a numeric between 0 and 1. Opacity of rugs
color_rugs	a character. Either name of a color or name of a variable that should be used for coloring
size_residuals	a numeric. Size of line and points to be plotted for residuals
alpha_residuals	a numeric between 0 and 1. Opacity of points and lines for residuals
color_residuals	a character. Either name of a color or name of a variable that should be used for coloring for residuals
only_numerical	a logical. If TRUE then only numerical variables will be plotted. If FALSE then only categorical variables will be plotted.
show_profiles	a logical. If TRUE then profiles will be plotted. Either individual or aggregate (see 'aggregate_profiles')
show_observations	a logical. If TRUE then individual observations will be marked as points
show_rugs	a logical. If TRUE then individual observations will be marked as rugs
show_residuals	a logical. If TRUE then residuals will be plotted as a line ended with a point
aggregate_profiles	function. If NULL (default) then individual profiles will be plotted. If a function (e.g. mean or median) then profiles will be aggregated and only the aggregate profile will be plotted
as.gg	if TRUE then returning plot will have gg class
facet_ncol	number of columns for the 'facet_wrap()'. selected_variables
selected_variables	if not NULL then only 'selected_variables' will be presented
init_plot	technical parameter, do not use.

Value

a ggplot2 object

Examples

```

library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

apartments_small_1 <- apartmentsTest[1,]
apartments_small_2 <- select_sample(apartmentsTest, n = 20)
apartments_small_3 <- select_neighbours(apartmentsTest, apartments_small_1, n = 20)

cp_rf_y1 <- ceteris_paribus(explainer_rf, apartments_small_1, y = apartments_small_1$m2.price)
cp_rf_y2 <- ceteris_paribus(explainer_rf, apartments_small_2, y = apartments_small_2$m2.price)
cp_rf_y3 <- ceteris_paribus(explainer_rf, apartments_small_3, y = apartments_small_3$m2.price)

tmp <- plot(cp_rf_y3, show_profiles = TRUE, show_observations = TRUE,
  show_residuals = TRUE, color = "black",
  alpha = 0.2, color_residuals = "darkred",
  selected_variables = c("construction.year", "surface"))

tmp <- plot(cp_rf_y3, show_profiles = TRUE, show_observations = TRUE,
  show_residuals = TRUE, color = "black",
  alpha = 0.2, color_residuals = "darkred")

tmp

tmp +
  ceteris_paribus_layer(cp_rf_y2, show_profiles = TRUE, show_observations = TRUE,
    alpha = 0.2, color = "darkblue")

tmp +
  ceteris_paribus_layer(cp_rf_y2, show_profiles = TRUE, show_observations = TRUE,
    alpha = 0.2, color = "darkblue") +
  ceteris_paribus_layer(cp_rf_y2, show_profiles = TRUE, show_observations = FALSE,
    alpha = 1, size = 2, color = "blue", aggregate_profiles = mean) +
  ceteris_paribus_layer(cp_rf_y1, show_profiles = TRUE, show_observations = FALSE,
    alpha = 1, size = 2, color = "red", aggregate_profiles = mean)

## End(Not run)

```

local_fit

Local Fit / Wangkardu Explanations

Description

Local Fit / Wangkardu Explanations

Usage

```
local_fit(explainer, observation, selected_variable, grid_points = 101,
          select_points = 0.1)
```

Arguments

```
explainer      a model to be explained, preprocessed by the 'DALEX::explain' function
observation    a new observation for which predictions need to be explained
selected_variable
               variable to be presented in the local fit plot
grid_points    number of points used for response path
select_points  fraction of points from validation data to be presented in local fit plots
```

Value

An object of the class 'local_fit_explainer'. It's a data frame with calculated average responses.

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

new_apartment <- apartmentsTest[1, ]
new_apartment

cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
cr_rf

## End(Not run)
```

```
plot.ceteris_paribus_explainer
```

Plot Ceteris Paribus Explanations

Description

Function 'plot.ceteris_paribus_explainer' plots Ceteris Paribus Plots for selected observations. Various parameters help to decide what should be plotted, profiles, aggregated profiles, points or rugs.

Usage

```
## S3 method for class 'ceteris_paribus_explainer'
plot(x, ..., size = 1, alpha = 0.3,
     color = "black", size_points = 2, alpha_points = 1,
     color_points = color, size_rugs = 0.5, alpha_rugs = 1,
     color_rugs = color, size_residuals = 1, alpha_residuals = 1,
     color_residuals = color, only_numerical = TRUE, show_profiles = TRUE,
     show_observations = TRUE, show_rugs = FALSE, show_residuals = FALSE,
     aggregate_profiles = NULL, as.gg = FALSE, facet_ncol = NULL,
     selected_variables = NULL)
```

Arguments

<code>x</code>	a ceteris paribus explainer produced with function <code>'ceteris_paribus()'</code>
<code>...</code>	other explainers that shall be plotted together
<code>size</code>	a numeric. Size of lines to be plotted
<code>alpha</code>	a numeric between 0 and 1. Opacity of lines
<code>color</code>	a character. Either name of a color or name of a variable that should be used for coloring
<code>size_points</code>	a numeric. Size of points to be plotted
<code>alpha_points</code>	a numeric between 0 and 1. Opacity of points
<code>color_points</code>	a character. Either name of a color or name of a variable that should be used for coloring
<code>size_rugs</code>	a numeric. Size of rugs to be plotted
<code>alpha_rugs</code>	a numeric between 0 and 1. Opacity of rugs
<code>color_rugs</code>	a character. Either name of a color or name of a variable that should be used for coloring
<code>size_residuals</code>	a numeric. Size of line and points to be plotted for residuals
<code>alpha_residuals</code>	a numeric between 0 and 1. Opacity of points and lines for residuals
<code>color_residuals</code>	a character. Either name of a color or name of a variable that should be used for coloring for residuals
<code>only_numerical</code>	a logical. If TRUE then only numerical variables will be plotted. If FALSE then only categorical variables will be plotted.
<code>show_profiles</code>	a logical. If TRUE then profiles will be plotted. Either individual or aggregate (see <code>'aggregate_profiles'</code>)
<code>show_observations</code>	a logical. If TRUE then individual observations will be marked as points
<code>show_rugs</code>	a logical. If TRUE then individual observations will be marked as rugs
<code>show_residuals</code>	a logical. If TRUE then residuals will be plotted as a line ended with a point

aggregate_profiles function. If NULL (default) then individual profiles will be plotted. If a function (e.g. mean or median) then profiles will be aggregated and only the aggregate profile will be plotted

as.gg if TRUE then returning plot will have gg class

facet_ncol number of columns for the 'facet_wrap()'

selected_variables if not NULL then only 'selected_variables' will be presented

Value

a ggplot2 object

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

apartments_small <- apartmentsTest[1:20,]
apartments_small_1 <- apartmentsTest[1,]
apartments_small_2 <- select_sample(apartmentsTest, n = 20)
apartments_small_3 <- select_neighbours(apartmentsTest, apartments_small_1, n = 20)

cp_rf <- ceteris_paribus(explainer_rf, apartments_small)
cp_rf_1 <- ceteris_paribus(explainer_rf, apartments_small_1)
cp_rf_2 <- ceteris_paribus(explainer_rf, apartments_small_2)
cp_rf_3 <- ceteris_paribus(explainer_rf, apartments_small_3)
cp_rf

cp_rf_y <- ceteris_paribus(explainer_rf, apartments_small, y = apartments_small$m2.price)
cp_rf_y1 <- ceteris_paribus(explainer_rf, apartments_small_1, y = apartments_small_1$m2.price)
cp_rf_y2 <- ceteris_paribus(explainer_rf, apartments_small_2, y = apartments_small_2$m2.price)
cp_rf_y3 <- ceteris_paribus(explainer_rf, apartments_small_3, y = apartments_small_3$m2.price)

plot(cp_rf_y, show_profiles = TRUE, show_observations = TRUE,
  show_residuals = TRUE, color = "black",
  alpha = 0.3, alpha_points = 1, alpha_residuals = 0.5,
  size_points = 2, size_rugs = 0.5)

plot(cp_rf_y, show_profiles = TRUE, show_observations = TRUE,
  show_residuals = TRUE, color = "black",
  selected_variables = c("construction.year", "surface"),
  alpha = 0.3, alpha_points = 1, alpha_residuals = 0.5,
```

```

        size_points = 2, size_rugs = 0.5)

plot(cp_rf_y1, show_profiles = TRUE, show_observations = TRUE, show_rugs = TRUE,
      show_residuals = TRUE, alpha = 0.5, size_points = 3,
      alpha_points = 1, size_rugs = 0.5)

plot(cp_rf_y2, show_profiles = TRUE, show_observations = TRUE, show_rugs = TRUE,
      alpha = 0.2, alpha_points = 1, size_rugs = 0.5)

plot(cp_rf_y3, show_profiles = TRUE, show_rugs = TRUE,
      show_residuals = TRUE, alpha = 0.2, color_residuals = "orange", size_rugs = 0.5)

plot(cp_rf_y, show_profiles = TRUE, show_observations = TRUE, show_rugs = TRUE, size_rugs = 0.5,
      show_residuals = TRUE, alpha = 0.5, color = "surface", as.gg = TRUE) +
  scale_color_gradient(low = "darkblue", high = "darkred")

plot(cp_rf_y1, show_profiles = TRUE, show_observations = TRUE, show_rugs = TRUE,
      show_residuals = TRUE, alpha = 0.5, color = "surface", size_points = 3)

plot(cp_rf_y2, show_profiles = TRUE, show_observations = TRUE, show_rugs = TRUE,
      size = 0.5, alpha = 0.5, color = "surface")

plot(cp_rf_y, show_profiles = TRUE, show_rugs = TRUE, size_rugs = 0.5,
      show_residuals = FALSE, aggregate_profiles = mean, color = "darkblue")

## End(Not run)

```

```
plot.local_fit_explainer
```

Local Fit Plots / Wangkardu Explanations

Description

Function 'plot.local_fit_explainer' plots Local Fit Plots for a single prediction / observation.

Usage

```

## S3 method for class 'local_fit_explainer'
plot(x, ..., plot_residuals = TRUE,
      palette = "default")

```

Arguments

x	a local fir explainer produced with the 'local_fit' function
...	other explainers that shall be plotted together
plot_residuals	if TRUE (default) then residuals are plotted as red/blue bars
palette	color palette. Currently the choice is limited to 'wangkardu' and 'default'

Value

a ggplot2 object

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

new_apartment <- apartmentsTest[1, ]
new_apartment

cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
plot(cr_rf, plot_residuals = FALSE)
plot(cr_rf)

cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
plot(cr_rf, plot_residuals = FALSE, palette = "wangkardu")
plot(cr_rf, palette = "wangkardu")

new_apartment <- apartmentsTest[10, ]
cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
plot(cr_rf, plot_residuals = FALSE)
plot(cr_rf)

new_apartment <- apartmentsTest[302, ]
cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
plot(cr_rf, plot_residuals = FALSE)
plot(cr_rf)

new_apartment <- apartmentsTest[720, ]
cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
plot(cr_rf, plot_residuals = FALSE)
plot(cr_rf)

## End(Not run)
```

 plot.what_if_explainer

Plot What If Explanations

Description

Function 'plot.what_if_explainer' plots What-If Plots for a single prediction / observation.

Usage

```
## S3 method for class 'what_if_explainer'
plot(x, ..., quantiles = TRUE, split = "models",
     split_ncol = NULL, color = "variables")
```

Arguments

x	a ceteris paribus explainer produced with the 'what_if' function
...	other explainers that shall be plotted together
quantiles	if TRUE (default) then quantiles will be presented on OX axis. If FALSE then original values will be presented on OX axis
split	a character, either 'models' or 'variables'. Sets the variable for faceting
split_ncol	number of columns for the 'facet_wrap'
color	a character, either 'models' or 'variables'. Sets the variable for coloring

Value

a ggplot2 object

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

new_apartment <- apartmentsTest[1, ]
new_apartment

wi_rf <- what_if(explainer_rf, observation = new_apartment)
wi_rf
```

```

plot(wi_rf, split = "variables", color = "variables")
plot(wi_rf)

## End(Not run)

```

plot_interactive	<i>Plots Interactive What-If Explanations</i>
------------------	---

Description

Function 'plot_interactive.what_if_explainer' plots Ceteris Paribus Plots for a single prediction.

Usage

```

## S3 method for class 'what_if_explainer'
plot_interactive(x, ..., split = "models",
  color = "variables")

plot_interactive(x, ...)

## Default S3 method:
plot_interactive(x, ..., split = "models",
  color = "variables")

```

Arguments

x	a ceteris_paribus explainer produced with the 'ceteris_paribus' function
...	other explainers that shall be plotted together
split	a character, either 'models' or 'variables'. Sets the variable for faceting
color	a character, either 'models' or 'variables'. Sets the variable for coloring

Value

a ggiraph object

Examples

```

library("DALEX")
## Not run:
library("ggiraph")
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

```

```

new_apartment <- apartmentsTest[1, ]
new_apartment

wi_rf <- what_if(explainer_rf, observation = new_apartment)
wi_rf

plot_interactive(wi_rf, split = "variables", color = "variables")

## End(Not run)

```

```

print.ceteris_paribus_explainer
      Print Ceteris Paribus Explainer Summary

```

Description

Print Ceteris Paribus Explainer Summary

Usage

```

## S3 method for class 'ceteris_paribus_explainer'
print(x, ...)

```

Arguments

```

x           a ceteris_paribus explainer produced with the 'ceteris_paribus()' function
...        other arguments that will be passed to 'head()'

```

Examples

```

library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

apartments_small <- select_sample(apartmentsTest, 10)

cp_rf <- ceteris_paribus(explainer_rf, apartments_small)
cp_rf

## End(Not run)

```

```
print.ceteris_paribus_profile
      Print Ceteris Paribus Profiles
```

Description

Print Ceteris Paribus Profiles

Usage

```
## S3 method for class 'ceteris_paribus_profile'
print(x, ...)
```

Arguments

x a ceteris paribus profile produced with the 'calculate_profiles' function
... other arguments that will be passed to head()

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)
apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
                                   no.rooms + district, data = apartments)
vars <- c("construction.year", "surface", "floor", "no.rooms", "district")
variable_splits <- calculate_variable_splits(apartments, vars)
new_apartment <- apartmentsTest[1:10, ]
profiles <- calculate_profiles(new_apartment, variable_splits,
                              apartments_rf_model)

profiles

# only subset of observations
small_apartments <- select_sample(apartmentsTest, n = 10)
small_apartments
small_profiles <- calculate_profiles(small_apartments, variable_splits,
                                   apartments_rf_model)

small_profiles

# neighbors for a selected observation
new_apartment <- apartments[1, 2:6]
small_apartments <- select_neighbours(apartmentsTest, new_apartment, n = 10)
small_apartments
small_profiles <- calculate_profiles(small_apartments, variable_splits,
                                   apartments_rf_model)

new_apartment
small_profiles

## End(Not run)
```

```
print.local_fit_explainer
      Prints Local Fit / Wangkardu Summary
```

Description

Prints Local Fit / Wangkardu Summary

Usage

```
## S3 method for class 'local_fit_explainer'
print(x, ...)
```

Arguments

x	a local fit explainer produced with the 'local_fit' function
...	other arguments that will be passed to 'head' function

Examples

```
library("DALEX")
## Not run:
library("randomForest")
apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)
explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)
new_apartment <- apartmentsTest[1, ]
new_apartment
cr_rf <- local_fit(explainer_rf, observation = new_apartment,
  select_points = 0.002, selected_variable = "surface")
cr_rf

## End(Not run)
```

```
print.plot_ceteris_paribus_explainer
      Print Ceteris Paribus Explainer Summary
```

Description

See more examples in the ceteris_paribus_layer function

Usage

```
## S3 method for class 'plot_ceteris_paribus_explainer'
print(x, ...)
```

Arguments

x a plot_ceteris_paribus_explainer object to plot
... other arguments that will be passed to 'print.ggplot()'

`print.what_if_explainer`*Print What If Explainer Summary*

Description

Print What If Explainer Summary

Usage

```
## S3 method for class 'what_if_explainer'  
print(x, ...)
```

Arguments

x a what_if explainer produced with the 'what_if' function
... other arguments that will be passed to head()

Examples

```
library("DALEX")  
## Not run:  
library("randomForest")  
apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +  
  no.rooms + district, data = apartments)  
explainer_rf <- explain(apartments_rf_model,  
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)  
new_apartment <- apartmentsTest[1, ]  
new_apartment  
  
## End(Not run)
```

select_neighbours	<i>Select Subset of Rows Closest to a Specified Observation</i>
-------------------	---

Description

This function selects subset of rows from data set. This is useful if data is large and we need just a sample to calculate profiles.

Usage

```
select_neighbours(data, observation, variables = NULL,  
  distance = gower::gower_dist, n = 20, frac = NULL)
```

Arguments

data	set of observations
observation	single observation
variables	variables that shall be used for calculation of distance. By default these are all variables present in 'data' and 'observation'
distance	distance function, by default the 'gower_dist' function.
n	number of neighbours to select
frac	if 'n' is not specified (NULL), then will be calculated as 'frac' * number of rows in 'data'. Either 'n' or 'frac' need to be specified.

Details

Note that select_neighbours function is S3 generic. If you want to work on non standard data sources (like H2O ddf, external databases) you should overload it.

Value

a data frame with selected rows

Examples

```
library("DALEX")  
  
new_apartment <- apartments[1, 2:6]  
small_apartments <- select_neighbours(apartmentsTest, new_apartment, n = 10)  
new_apartment  
small_apartments
```

select_sample	<i>Select Subset of Rows</i>
---------------	------------------------------

Description

This function selects subset of rows from data set. This is useful if data is large and we need just a sample to calculate profiles.

Usage

```
select_sample(data, n = 100, seed = 1313)
```

Arguments

data	set of observations. Profile will be calculated for every observation (every row)
n	named list of vectors. Elements of the list are vectors with points in which profiles should be calculated. See an example for more details.
seed	seed for random number generator.

Details

Note that select_subsample function is S3 generic. If you want to work on non standard data sources (like H2O ddf, external databases) you should overload it.

Value

a data frame with selected rows

Examples

```
library("DALEX")
small_apartments <- select_sample(apartmentsTest)
head(small_apartments)
```

what_if	<i>What-If Plot</i>
---------	---------------------

Description

What-If Plot

Usage

```
what_if(explainer, observation, grid_points = 101,
        selected_variables = NULL)
```

Arguments

`explainer` a model to be explained, preprocessed by the 'DALEX::explain' function
`observation` a new observation for which predictions need to be explained
`grid_points` number of points used for response path
`selected_variables`
if specified, then only these variables will be explained

Value

An object of the class 'what_if_explainer'. It's a data frame with calculated average responses.

Examples

```
library("DALEX")
## Not run:
library("randomForest")
set.seed(59)

apartments_rf_model <- randomForest(m2.price ~ construction.year + surface + floor +
  no.rooms + district, data = apartments)

explainer_rf <- explain(apartments_rf_model,
  data = apartmentsTest[,2:6], y = apartmentsTest$m2.price)

new_apartment <- apartmentsTest[1, ]
new_apartment

wi_rf <- what_if(explainer_rf, observation = new_apartment)
wi_rf
wi_rf <- what_if(explainer_rf, observation = new_apartment,
  selected_variables = c("surface", "floor", "no.rooms"))
wi_rf

## End(Not run)
```

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