

# Package ‘ceterisParibus’

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**Title** Ceteris Paribus Profiles

**Version** 0.4.2

**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

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

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

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**NeedsCompilation** no

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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_oscillations`*Calculate Oscillations for Ceteris Paribus Explainer*

---

**Description**

Calculate Oscillations for Ceteris Paribus Explainer

**Usage**

```
calculate_oscillations(x, sort = TRUE, ...)
```

**Arguments**

<code>x</code>	a <code>ceteris_paribus</code> explainer produced with the <code>'ceteris_paribus()'</code> function
<code>sort</code>	a logical value. If <code>TRUE</code> then rows are sorted along the oscillations
<code>...</code>	other arguments

**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, y = apartmentsTest$m2.price)

apartment <- apartmentsTest[1,]

cp_rf <- ceteris_paribus(explainer_rf, apartment)
calculate_oscillations(cp_rf)

## End(Not run)
```

---

`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**

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

**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\_profiles\_lce

*Calculate Local Conditional Expectation profiles*

---

## Description

This function Local Conditional Expectation profiles

## Usage

```

calculate_profiles_lce(
  data,
  variable_splits,
  model,
  dataset,
  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
dataset	a data.frame, usually training data of a model, used for calculation of LCE profiles
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_lce` 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)
explainer_rf <- explain(apartments_rf_model,
                       data = apartments[,2:6], y = apartments$m2.price)
vars <- c("construction.year", "surface", "floor", "no.rooms", "district")
variable_splits <- calculate_variable_splits(apartments, vars)
new_apartment <- apartments[1, ]

profiles <- calculate_profiles_lce(new_apartment, variable_splits,
                                  apartments_rf_model, explainer_rf$data)

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**

<code>data</code>	validation dataset. Is used to determine distribution of observations.
<code>variables</code>	names of variables for which splits shall be calculated
<code>grid_points</code>	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

<code>explainer</code>	a model to be explained, preprocessed by function <code>'DALEX::explain()'</code> .
<code>observations</code>	set of observation for which profiles are to be calculated
<code>y</code>	true labels for <code>'observations'</code> . 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**

<code>x</code>	a ceteris paribus explainer produced with function ‘ceteris_paribus()’
<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

`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` 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\_conditional\_expectations

*Local Conditional Expectation Explainer*

---

## Description

This explainer works for individual observations. For each observation it calculates Local Conditional Expectation (LCE) profiles for selected variables.

## Usage

```

local_conditional_expectations(
  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

<code>y</code>	true labels for ‘observations’. If specified then will be added to local conditional expectations plots.
<code>variable_splits</code>	named list of splits for variables, in most cases created with ‘ <code>calculate_variable_splits()</code> ’. If NULL then it will be calculated based on validation data available in the ‘explainer’.
<code>variables</code>	names of variables for which profiles shall be calculated. Will be passed to ‘ <code>calculate_variable_splits()</code> ’. If NULL then all variables from the validation data will be used.
<code>grid_points</code>	number of points for profile. Will be passed to ‘ <code>calculate_variable_splits()</code> ’.

**Value**

An object of the class ‘`ceteris_paribus_explainer`’. A data frame with calculated LCE profiles.

**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 = apartments[,2:6], y = apartments$m2.price)

new_apartment <- apartments[1, ]

cp_rf <- ceteris_paribus(explainer_rf, new_apartment)
lce_rf <- local_conditional_expectations(explainer_rf, new_apartment)
lce_rf

lce_rf <- local_conditional_expectations(explainer_rf, new_apartment, y = new_apartment$m2.price)
lce_rf

# Plot LCE
sel_vars <- c("surface", "no.rooms")
plot(lce_rf, selected_variables = sel_vars)

# Compare ceteris paribus profiles with LCE profiles
plot(cp_rf, selected_variables = sel_vars) +
  ceteris_paribus_layer(lce_rf, selected_variables = sel_vars, color = "red")

## 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

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	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.ceteris_paribus_oscillations
      Plot Ceteris Paribus Oscillations

```

---

## Description

Function 'plot.ceteris\_paribus\_oscillations' plots variable importance plots.

## Usage

```

## S3 method for class 'ceteris_paribus_oscillations'
plot(x, ...)

```

## Arguments

x a ceteris paribus oscillation explainer produced with function 'calculate\_oscillations()'  
 ... other explainers that shall be plotted together

## 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, y = apartmentsTest$m2.price)

apartment <- apartmentsTest[1:2,]

```

```

cp_rf <- ceteris_paribus(explainer_rf, apartment)
plot(cp_rf, color = "_ids_")

vips <- calculate_oscillations(cp_rf)
vips
plot(vips)

## 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 fit 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\_2d\_explainer

*Plot What If 2D Explanations*

---

### Description

Function 'plot.what\_if\_2d\_explainer' plots What-If Plots for a single prediction / observation.

### Usage

```

## S3 method for class 'what_if_2d_explainer'
plot(
  x,
  ...,
  split_ncol = NULL,
  add_raster = TRUE,
  add_contour = TRUE,
  add_observation = TRUE,
  bins = 3
)

```

**Arguments**

<code>x</code>	a ceteris paribus explainer produced with the 'what_if_2d' function
<code>...</code>	currently will be ignored
<code>split_ncol</code>	number of columns for the 'facet_wrap'
<code>add_raster</code>	if TRUE then 'geom_raster' will be added to present levels with diverging colors
<code>add_contour</code>	if TRUE then 'geom_contour' will be added to present contours
<code>add_observation</code>	if TRUE then 'geom_point' will be added to present observation that is explained
<code>bins</code>	number of contours to be added

**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_2d <- what_if_2d(explainer_rf, observation = new_apartment)
wi_rf_2d

plot(wi_rf_2d)
plot(wi_rf_2d, add_contour = FALSE)
plot(wi_rf_2d, add_observation = FALSE)
plot(wi_rf_2d, add_raster = FALSE)

# HR data
model <- randomForest(status ~ gender + age + hours + evaluation + salary, data = HR)
pred1 <- function(m, x) predict(m, x, type = "prob")[,1]
explainer_rf_fired <- explain(model, data = HR[,1:5],
  y = HR$status == "fired",
  predict_function = pred1, label = "fired")

new_emp <- HR[1, ]
new_emp
```

```

wi_rf_2d <- what_if_2d(explainer_rf_fired, observation = new_emp)
wi_rf_2d

plot(wi_rf_2d)

## 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

*Plots Interactive What-If Explanations*


---

**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**

<code>x</code>	a <code>ceteris_paribus</code> explainer produced with the <code>'ceteris_paribus'</code> function
<code>...</code>	other explainers that shall be plotted together
<code>split</code>	a character, either <code>'models'</code> or <code>'variables'</code> . Sets the variable for faceting
<code>color</code>	a character, either <code>'models'</code> or <code>'variables'</code> . 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_2d_explainer
  Print What If 2D Explainer Summary

```

---

**Description**

Print What If 2D Explainer Summary

**Usage**

```

## S3 method for class 'what_if_2d_explainer'
print(x, ...)

```

**Arguments**

`x` a `what_if_2d` explainer produced with the `'what_if_2d'` 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)
```

---

```
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      *Select Subset of Rows Closest to a Specified Observation*

---

### 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 usefull 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)

```

---

 what\_if\_2d

*What-If 2D Plot*


---

**Description**

This function calculates what if scores for grid of values spanned by two variables.

**Usage**

```

what_if_2d(
  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. Will be used for both variables  
selected\_variables  
                 if specified, then only these variables will be explained

**Value**

An object of the class 'what\_if\_2d\_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_2d <- what_if_2d(explainer_rf, observation = new_apartment,
  selected_variables = c("surface", "floor", "no.rooms"))
wi_rf_2d
plot(wi_rf_2d)

## End(Not run)
```

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