

Package ‘easyalluvial’

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Title Generate Alluvial Plots with a Single Line of Code

Version 0.4.0

URL <https://github.com/erblast/easyalluvial/>,
<https://erblast.github.io/easyalluvial/>

BugReports <https://github.com/erblast/easyalluvial/issues>

Description Alluvial plots are similar to sankey diagrams and visualise categorical data over multiple dimensions as flows. (Rosvall M, Bergstrom CT (2010) Mapping Change in Large Networks. PLoS ONE 5(1): e8694. <[doi:10.1371/journal.pone.0008694](https://doi.org/10.1371/journal.pone.0008694)> Their graphical grammar however is a bit more complex than that of a regular x/y plots. The 'ggalluvial' package made a great job of translating that grammar into 'ggplot2' syntax and gives you many options to tweak the appearance of an alluvial plot, however there still remains a multi-layered complexity that makes it difficult to use 'ggalluvial' for explorative data analysis. 'easyalluvial' provides a simple interface to this package that allows you to produce a decent alluvial plot from any dataframe in either long or wide format from a single line of code while also handling continuous data. It is meant to allow a quick visualisation of entire dataframes with a focus on different colouring options that can make alluvial plots a great tool for data exploration.

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Encoding UTF-8

LazyData true

Depends R(>= 3.5)

Suggests testthat, covr, ISLR, nycflights13, vdiffr (>= 0.3.1),
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caret, parsnip, vip, rpart, glmnet, xgboost

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Imports purrr , tidyr (>= 1.0.0) , dplyr , forcats , ggalluvial (>= 0.9.1) , ggplot2 (>= 3.2.0) , ggridges , RColorBrewer , recipes (>= 0.1.5) , rlang , stringr , magrittr , tibble , gridExtra , randomForest , progressr , progress

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add_imp_plot	<i>add bar plot of important features to model response alluvial plot</i>
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Description

adds bar plot of important features to model response alluvial plot

Usage

```
add_imp_plot(grid, p = NULL, data_input, plot = T, ...)
```

Arguments

grid	gtable or ggplot
p	alluvial plot, optional if alluvial plot has already been passed as grid. Default: NULL
data_input	dataframe used to generate alluvial plot
plot	logical if plot should be drawn or not
...	additional parameters passed to plot_imp

Value

gtable

See Also

[arrangeGrob](#) [plot_imp](#)

Examples

```
## Not run:
df = mtcars2[, ! names(mtcars2) %in% 'ids' ]

train = caret::train( disp ~ .
                      , df
                      , method = 'rf'
                      , trControl = caret::trainControl( method = 'none' )
                      , importance = TRUE )

pred_train = caret::predict.train(train, df)

p = alluvial_model_response_caret(train, degree = 4, pred_train = pred_train)

p_grid = add_marginal_histograms(p, data_input = df)

p_grid = add_imp_plot(p_grid, p, data_input = df)

## End(Not run)
```

```
add_marginal_histograms
```

add marginal histograms to alluvial plot

Description

will add density histograms and frequency plots of original data to alluvial plot

Usage

```
add_marginal_histograms(
  p,
  data_input,
  top = TRUE,
  keep_labels = FALSE,
  plot = TRUE,
  ...
)
```

Arguments

<code>p</code>	alluvial plot
<code>data_input</code>	dataframe, input data that was used to create dataframe
<code>top</code>	logical, position of histograms, if FALSE adds them at the bottom, Default: TRUE
<code>keep_labels</code>	logical, keep title and caption, Default: FALSE
<code>plot</code>	logical if plot should be drawn or not
<code>...</code>	additional arguments for model response alluvial plot concerning the response variable
pred_train	display training prediction, not necessary if pred_train has already been passed to alluvial_model_response()
scale	int, y-axis distance between the ridge plots, Default: 400
resp_var	character vector, specify response variable in data_input, if not set response variable will try to be inferred, Default: NULL

Value

gtable

See Also

[arrangeGrob](#)

Examples

```
## Not run:
p = alluvial_wide(mtcars2, max_variables = 3)
p_grid = add_marginal_histograms(p, mtcars2)

## End(Not run)
```

alluvial_long

alluvial plot of data in long format

Description

Plots two variables of a dataframe on an alluvial plot. A third variable can be added either to the left or the right of the alluvial plot to provide coloring of the flows. All numerical variables are scaled, centered and YeoJohnson transformed before binning.

Usage

```
alluvial_long(
  data,
  key,
  value,
  id,
  fill = NULL,
  fill_right = T,
  bins = 5,
  bin_labels = c("LL", "ML", "M", "MH", "HH"),
  NA_label = "NA",
  order_levels_value = NULL,
  order_levels_key = NULL,
  order_levels_fill = NULL,
  complete = TRUE,
  fill_by = "first_variable",
  col_vector_flow = palette_qualitative() %>% palette_filter(greys = F),
  col_vector_value = RColorBrewer::brewer.pal(9, "Greys")[c(3, 6, 4, 7, 5)],
  verbose = F,
  stratum_labels = T,
  stratum_label_type = "label",
  stratum_label_size = 4.5,
  stratum_width = 1/4,
  auto_rotate_xlabs = T,
  ...
)
```

Arguments

<code>data</code>	a dataframe
<code>key</code>	unquoted column name or string of x axis variable
<code>value</code>	unquoted column name or string of y axis variable
<code>id</code>	unquoted column name or string of id column
<code>fill</code>	unquoted column name or string of fill variable which will be used to color flows, Default: NULL
<code>fill_right</code>	logical, TRUE fill variable is added to the right FALSE to the left, Default: T
<code>bins</code>	number of bins for automatic binning of numerical variables, Default: 5
<code>bin_labels</code>	labels for bins, Default: c("LL", "ML", "M", "MH", "HH")
<code>NA_label</code>	character vector define label for missing data
<code>order_levels_value</code>	character vector denoting order of y levels from low to high, does not have to be complete can also just be used to bring levels to the front, Default: NULL
<code>order_levels_key</code>	character vector denoting order of x levels from low to high, does not have to be complete can also just be used to bring levels to the front, Default: NULL
<code>order_levels_fill</code>	character vector denoting order of color fill variable levels from low to high, does not have to be complete can also just be used to bring levels to the front, Default: NULL
<code>complete</code>	logical, insert implicitly missing observations, Default: TRUE
<code>fill_by</code>	one_of(c('first_variable', 'last_variable', 'all_flows', 'values')), Default: 'first_variable'
<code>col_vector_flow</code>	HEX color values for flows, Default: palette_filter(greys = F)
<code>col_vector_value</code>	HEX color values for y levels/values, Default: RColorBrewer::brewer.pal(9, 'Greys')[c(3,6,4,7,5)]
<code>verbose</code>	logical, print plot summary, Default: F
<code>stratum_labels</code>	logical, Default: TRUE
<code>stratum_label_type</code>	character, Default: "label"
<code>stratum_label_size</code>	numeric, Default: 4.5
<code>stratum_width</code>	double, Default: 1/4
<code>auto_rotate_xlabs</code>	logical, Default: TRUE
<code>...</code>	additional parameter passed to manip_bin_numerics

Value

ggplot2 object

See Also

[alluvial_wide](#), [geom_flow](#), [geom_stratum](#), [manip_bin_numerics](#)

Examples

```
## Not run:
data = quarterly_flights

alluvial_long( data, key = qu, value = mean_arr_delay, id = tailnum, fill_by = 'last_variable' )

# more flow coloring variants -----

alluvial_long( data, key = qu, value = mean_arr_delay, id = tailnum, fill_by = 'first_variable' )
alluvial_long( data, key = qu, value = mean_arr_delay, id = tailnum, fill_by = 'all_flows' )
alluvial_long( data, key = qu, value = mean_arr_delay, id = tailnum, fill_by = 'value' )

# color by additional variable carrier -----

alluvial_long( data, key = qu, value = mean_arr_delay, fill = carrier, id = tailnum )

# use same color coding for flows and y levels -----

palette = c('green3', 'tomato')

alluvial_long( data, qu, mean_arr_delay, tailnum, fill_by = 'value'
              , col_vector_flow = palette
              , col_vector_value = palette )

# reorder levels -----

alluvial_long( data, qu, mean_arr_delay, tailnum, fill_by = 'first_variable'
              , order_levels_value = c('on_time', 'late') )

alluvial_long( data, qu, mean_arr_delay, tailnum, fill_by = 'first_variable'
              , order_levels_key = c('Q4', 'Q3', 'Q2', 'Q1') )

require(dplyr)
require(magrittr)

order_by_carrier_size = data %>%
  group_by(carrier) %>%
  count() %>%
  arrange( desc(n) ) %>%
  .[['carrier']]

alluvial_long( data, qu, mean_arr_delay, tailnum, carrier
              , order_levels_fill = order_by_carrier_size )

## End(Not run)
```

```
alluvial_model_response
    create model response plot
```

Description

alluvial plots are capable of displaying higher dimensional data on a plane, thus lend themselves to plot the response of a statistical model to changes in the input data across multiple dimensions. The practical limit here is 4 dimensions. We need the data space (a sensible range of data calculated based on the importance of the explanatory variables of the model as created by [get_data_space](#) and the predictions returned by the model in response to the data space.

Usage

```
alluvial_model_response(
  pred,
  dspace,
  imp,
  degree = 4,
  bin_labels = c("LL", "ML", "M", "MH", "HH"),
  col_vector_flow = c("#FF0065", "#009850", "#A56F2B", "#005EAA", "#710500", "#7B5380",
    "#9DD1D1"),
  method = "median",
  force = FALSE,
  params_bin_numeric_pred = list(bins = 5),
  pred_train = NULL,
  stratum_label_size = 3.5,
  ...
)
```

Arguments

pred	vector, predictions, if method = 'pdp' use get_pdp_predictions to calculate predictions
dspace	data frame, returned by get_data_space
imp	dataframe, with not more then two columns one of them numeric containing importance measures and one character or factor column containing corresponding variable names as found in training data.
degree	integer, number of top important variables to select. For plotting more than 4 will result in too many flows and the alluvial plot will not be very readable, Default: 4
bin_labels	labels for prediction bins from low to high, Default: c("LL", "ML", "M", "MH", "HH")
col_vector_flow	character vector, defines flow colours, Default: c('#FF0065', '#009850', '#A56F2B', '#005EAA', '#710500')

method character vector, one of c('median', 'pdp')

median sets variables that are not displayed to median mode, use with regular predictions

pdp partial dependency plot method, for each observation in the training data the displayed variable as are set to the indicated values. The predict function is called for each modified observation and the result is averaged, calculate predictions using [get_pdp_predictions](#)

. Default: 'median'

force logical, force plotting of over 1500 flows, Default: FALSE

params_bin_numeric_pred list, additional parameters passed to [manip_bin_numerics](#) which is applied to the pred parameter. Default: list(bins = 5, center = T, transform = T, scale = T)

pred_train numeric vector, base the automated binning of the pred vector on the distribution of the training predictions. This is useful if marginal histograms are added to the plot later. Default = NULL

stratum_label_size numeric, Default: 3.5

... additional parameters passed to [alluvial_wide](#)

Details

this model visualisation approach follows the "visualising the model in the dataspace" principle as described in Wickham H, Cook D, Hofmann H (2015) Visualizing statistical models: Removing the blindfold. Statistical Analysis and Data Mining 8(4) <doi:10.1002/sam.11271>

Value

ggplot2 object

See Also

[alluvial_wide](#), [get_data_space](#), [alluvial_model_response_caret](#)

Examples

```
df = mtcars2[, ! names(mtcars2) %in% 'ids' ]
m = randomForest::randomForest( disp ~ ., df)
imp = m$importance
dspace = get_data_space(df, imp, degree = 3)
pred = predict(m, newdata = dspace)
alluvial_model_response(pred, dspace, imp, degree = 3)

# partial dependency plotting method
## Not run:
pred = get_pdp_predictions(df, imp
                           , .f_predict = randomForest::predict.randomForest
                           , m
                           , degree = 3)
```

```

, bins = 5)

alluvial_model_response(pred, dspace, imp, degree = 3, method = 'pdp')

## End(Not run)

```

```

alluvial_model_response_caret
  create model response plot for caret models

```

Description

Wraps [alluvial_model_response](#) and [get_data_space](#) into one call for caret models.

Usage

```

alluvial_model_response_caret(
  train,
  data_input,
  degree = 4,
  bins = 5,
  bin_labels = c("LL", "ML", "M", "MH", "HH"),
  col_vector_flow = c("#FF0065", "#009850", "#A56F2B", "#005EAA", "#710500", "#7B5380",
    "#9DD1D1"),
  method = "median",
  parallel = FALSE,
  params_bin_numeric_pred = list(bins = 5),
  pred_train = NULL,
  stratum_label_size = 3.5,
  force = F,
  resp_var = NULL,
  ...
)

```

Arguments

train	caret train object
data_input	dataframe, input data
degree	integer, number of top important variables to select. For plotting more than 4 will result in too many flows and the alluvial plot will not be very readable, Default: 4
bins	integer, number of bins for numeric variables, increasing this number might result in too many flows, Default: 5
bin_labels	labels for the bins from low to high, Default: c("LL", "ML", "M", "MH", "HH")

col_vector_flow	character vector, defines flow colours, Default: c('#FF0065', '#009850', '#A56F2B', '#005EAA', '#710500')
method	character vector, one of c('median', 'pdp') median sets variables that are not displayed to median mode, use with regular predictions pdp partial dependency plot method, for each observation in the training data the displayed variables are set to the indicated values. The predict function is called for each modified observation and the result is averaged . Default: 'median'
parallel	logical, turn on parallel processing for pdp method. Default: FALSE
params_bin_numeric_pred	list, additional parameters passed to manip_bin_numerics which is applied to the pred parameter. Default: list(bins = 5, center = T, transform = T, scale = T)
pred_train	numeric vector, base the automated binning of the pred vector on the distribution of the training predictions. This is useful if marginal histograms are added to the plot later. Default = NULL
stratum_label_size	numeric, Default: 3.5
force	logical, force plotting of over 1500 flows, Default: FALSE
resp_var	character, sometimes target variable cannot be inferred and needs to be passed. Default NULL
...	additional parameters passed to alluvial_wide

Details

this model visualisation approach follows the "visualising the model in the dataspace" principle as described in Wickham H, Cook D, Hofmann H (2015) Visualizing statistical models: Removing the blindfold. Statistical Analysis and Data Mining 8(4) <doi:10.1002/sam.11271>

Value

ggplot2 object

Parallel Processing

We are using 'furrr' and the 'future' package to paralelize some of the computational steps for calculating the predictions. It is up to the user to register a compatible backend (see [plan](#)).

See Also

[alluvial_wide](#), [get_data_space](#), [varImp](#), [extractPrediction](#), [get_data_space](#), [get_pdp_predictions](#)

Examples

```

if(check_pkg_installed("caret", raise_error = FALSE)) {
  df = mtcars2[, ! names(mtcars2) %in% 'ids' ]

  train = caret::train( disp ~ .,
                        df,
                        method = 'rf',
                        trControl = caret::trainControl( method = 'none' ),
                        importance = TRUE )

  alluvial_model_response_caret(train, df, degree = 3)
}
# partial dependency plotting method
## Not run:
future::plan("multisession")
alluvial_model_response_caret(train, df, degree = 3, method = 'pdp', parallel = TRUE)

## End(Not run)

```

alluvial_model_response_parsnip

create model response plot for parsnip models

Description

Wraps [alluvial_model_response](#) and [get_data_space](#) into one call for parsnip models.

Usage

```

alluvial_model_response_parsnip(
  m,
  data_input,
  degree = 4,
  bins = 5,
  bin_labels = c("LL", "ML", "M", "MH", "HH"),
  col_vector_flow = c("#FF0065", "#009850", "#A56F2B", "#005EAA", "#710500", "#7B5380",
    "#9DD1D1"),
  method = "median",
  parallel = FALSE,
  params_bin_numeric_pred = list(bins = 5),
  pred_train = NULL,
  stratum_label_size = 3.5,
  force = F,
  resp_var = NULL,
  .f_imp = vip::vi_model,
  ...
)

```

Arguments

<code>m</code>	parsnip model or trained workflow
<code>data_input</code>	dataframe, input data
<code>degree</code>	integer, number of top important variables to select. For plotting more than 4 will result in too many flows and the alluvial plot will not be very readable, Default: 4
<code>bins</code>	integer, number of bins for numeric variables, increasing this number might result in too many flows, Default: 5
<code>bin_labels</code>	labels for the bins from low to high, Default: <code>c("LL", "ML", "M", "MH", "HH")</code>
<code>col_vector_flow</code>	character vector, defines flow colours, Default: <code>c('#FF0065', '#009850', '#A56F2B', '#005EAA', '#710500')</code>
<code>method</code>	character vector, one of <code>c('median', 'pdp')</code> median sets variables that are not displayed to median mode, use with regular predictions pdp partial dependency plot method, for each observation in the training data the displayed variables are set to the indicated values. The predict function is called for each modified observation and the result is averaged . Default: 'median'
<code>parallel</code>	logical, turn on parallel processing for pdp method. Default: FALSE
<code>params_bin_numeric_pred</code>	list, additional parameters passed to <code>manip_bin_numerics</code> which is applied to the pred parameter. Default: <code>list(bins = 5, center = T, transform = T, scale = T)</code>
<code>pred_train</code>	numeric vector, base the automated binning of the pred vector on the distribution of the training predictions. This is useful if marginal histograms are added to the plot later. Default = NULL
<code>stratum_label_size</code>	numeric, Default: 3.5
<code>force</code>	logical, force plotting of over 1500 flows, Default: FALSE
<code>resp_var</code>	character, sometimes target variable cannot be inferred and needs to be passed. Default NULL
<code>.f_imp</code>	vip function that calculates feature importance, Default: <code>vip::vi_model</code>
<code>...</code>	additional parameters passed to <code>alluvial_wide</code>

Details

this model visualisation approach follows the "visualising the model in the dataspace" principle as described in Wickham H, Cook D, Hofmann H (2015) Visualizing statistical models: Removing the blindfold. Statistical Analysis and Data Mining 8(4) <doi:10.1002/sam.11271>

Value

ggplot2 object

Parallel Processing

We are using ‘furr’ and the ‘future’ package to paralelize some of the computational steps for calculating the predictions. It is up to the user to register a compatible backend (see [plan](#)).

See Also

[alluvial_wide](#), [get_data_space](#), [varImp](#), [extractPrediction](#), [get_data_space](#), [get_pdp_predictions](#)

Examples

```
if(check_pkg_installed("parsnip", raise_error = FALSE) &
  check_pkg_installed("vip", raise_error = FALSE)) {
  df = mtcars2[, ! names(mtcars2) %in% 'ids' ]

  m = parsnip::rand_forest(mode = "regression") %>%
    parsnip::set_engine("randomForest") %>%
    parsnip::fit(dis ~ ., data = df)

  alluvial_model_response_parsnip(m, df, degree = 3)
}
## Not run:
# workflow -----
m <- parsnip::rand_forest(mode = "regression") %>%
  parsnip::set_engine("randomForest")

rec_prep = recipes::recipe(dis ~ ., df)

wf <- workflows::workflow() %>%
  workflows::add_model(m) %>%
  workflows::add_recipe(rec_prep) %>%
  parsnip::fit(df)

alluvial_model_response_parsnip(wf, df, degree = 3)

# partial dependence plotting method -----
future::plan("multisession")
alluvial_model_response_parsnip(m, df, degree = 3, method = 'pdp', parallel = TRUE)

## End(Not run)
```

alluvial_wide

alluvial plot of data in wide format

Description

plots a dataframe as an alluvial plot. All numerical variables are scaled, centered and YeoJohnson transformed before binning. Plots all variables in the sequence as they appear in the dataframe until maximum number of values is reached.

Usage

```
alluvial_wide(
  data,
  id = NULL,
  max_variables = 20,
  bins = 5,
  bin_labels = c("LL", "ML", "M", "MH", "HH"),
  NA_label = "NA",
  order_levels = NULL,
  fill_by = "first_variable",
  col_vector_flow = palette_qualitative() %>% palette_filter(greys = F),
  col_vector_value = RColorBrewer::brewer.pal(9, "Greys")[c(4, 7, 5, 8, 6)],
  colorful_fill_variable_stratum = T,
  verbose = F,
  stratum_labels = T,
  stratum_label_type = "label",
  stratum_label_size = 4.5,
  stratum_width = 1/4,
  auto_rotate_xlabs = T,
  ...
)
```

Arguments

<code>data</code>	a dataframe
<code>id</code>	unquoted column name of id column or character vector with id column name
<code>max_variables</code>	maximum number of variables, Default: 20
<code>bins</code>	number of bins for numerical variables, Default: 5
<code>bin_labels</code>	labels for the bins from low to high, Default: c("LL", "ML", "M", "MH", "HH")
<code>NA_label</code>	character vector, define label for missing data, Default: 'NA'
<code>order_levels</code>	character vector denoting levels to be reordered from low to high
<code>fill_by</code>	one_of(c('first_variable', 'last_variable', 'all_flows', 'values')), Default: 'first_variable'
<code>col_vector_flow</code>	HEX colors for flows, Default: palette_filter(greys = F)
<code>col_vector_value</code>	Hex colors for y levels/values, Default: RColorBrewer::brewer.pal(9, "Greys")[c(3, 6, 4, 7, 5)]
<code>colorful_fill_variable_stratum</code>	logical, use flow colors to colorize fill variable stratum, Default: TRUE
<code>verbose</code>	logical, print plot summary, Default: F
<code>stratum_labels</code>	logical, Default: TRUE
<code>stratum_label_type</code>	character, Default: "label"
<code>stratum_label_size</code>	numeric, Default: 4.5

```

stratum_width  double, Default: 1/4
auto_rotate_xlabs
                logical, Default: TRUE
...            additional arguments passed to manip\_bin\_numerics

```

Details

Under the hood this function converts the wide format into long format. `ggalluvial` also offers a way to make alluvial plots directly from wide format tables but it does not allow individual colouring of the stratum segments. The tradeoff is that we can only order levels as a whole and not individually by variable, Thus if some variables have levels with the same name the order will be the same. If we want to change level order independently we have to assign unique level names first.

Value

ggplot2 object

See Also

[alluvial_wide](#), [geom_flow](#), [geom_stratum](#), [manip_bin_numerics](#)

Examples

```

## Not run:
alluvial_wide( data = mtcars2, id = ids
               , max_variables = 3
               , fill_by = 'first_variable' )#'
# more coloring variants-----
alluvial_wide( data = mtcars2, id = ids
               , max_variables = 5
               , fill_by = 'last_variable' )

alluvial_wide( data = mtcars2, id = ids
               , max_variables = 5
               , fill_by = 'all_flows' )

alluvial_wide( data = mtcars2, id = ids
               , max_variables = 5
               , fill_by = 'first_variable' )

# manually order variable values and colour by stratum value

alluvial_wide( data = mtcars2, id = ids
               , max_variables = 5
               , fill_by = 'values'
               , order_levels = c('4', '8', '6') )

## End(Not run)

```

check_pkg_installed	<i>check if package is installed</i>
---------------------	--------------------------------------

Description

check if package is installed

Usage

```
check_pkg_installed(pkg, raise_error = TRUE)
```

Arguments

pkg	character, package name
raise_error	logical

Value

logical

Examples

```
check_pkg_installed("easyalluvial")
```

get_data_space	<i>calculate data space</i>
----------------	-----------------------------

Description

calculates a dataspace based on the modeling dataframe and the importance of the explanatory variables. It only considers the most important variables as defined by the degree parameter. It selects a number (defined by bins) of sensible single values spread over the range of the numeric variables and creates all possible value combinations among the most important variables. The values of the remaining variables are set to mode(factors) or median(numerics).

Usage

```
get_data_space(df, imp, degree = 4, bins = 5, max_levels = 10)
```

Arguments

df	dataframe, training data
imp	dataframe, with not more then two columns one of them numeric containing importance measures and one character or factor column containing corresponding variable names as found in training data.
degree	integer, number of top important variables to select. For plotting more than 4 will result in two many flows and the alluvial plot will not be very readable, Default: 4
bins	integer, number of bins for numeric variables, and maximum number of levels for factor variables, increasing this number might result in too many flows, Default: 5
max_levels	integer, maximum number of levels per factor variable, Default: 10

Details

It selects a the top most important variables based on the degree parameter and bins the numeric variables using [manip_bin_numerics](#), while leaving categoric variables unchanged. The number of bins for each numeric variable is set to bins -2. Next the median is picked for each of the bins and the min and the max value is added for each numeric variable So that we get (median(bin) X bins -2, max, min) for each numeric variable. Then all possible combinations between those values and the categoric factor levels are created. The total number of all possible combinations defines the range of the data space. The values of the remaining variables are set to mode(factors) or median(numerics).

this model visualisation approach follows the "visualising the model in the dataspace" principle as described in Wickham H, Cook D, Hofmann H (2015) Visualizing statistical models: Removing the blindfold. Statistical Analysis and Data Mining 8(4) <doi:10.1002/sam.11271>

Value

data frame

See Also

[alluvial_wide](#), [manip_bin_numerics](#)

Examples

```
df = mtcars2[, ! names(mtcars2) %in% 'ids' ]
m = randomForest::randomForest( disp ~ ., df)
imp = m$importance
dspace = get_data_space(df, imp)
```

get_pdp_predictions	<i>get predictions compatible with the partial dependence plotting method</i>
---------------------	---

Description

Alluvial plots are capable of displaying higher dimensional data on a plane, thus lend themselves to plot the response of a statistical model to changes in the input data across multiple dimensions. The practical limit here is 4 dimensions while conventional partial dependence plots are limited to 2 dimensions.

Briefly the 4 variables with the highest feature importance for a given model are selected and 5 values spread over the variable range are selected for each. Then a grid of all possible combinations is created. All none-plotted variables are set to the values found in the first row of the training data set. Using this artificial data space model predictions are being generated. This process is then repeated for each row in the training data set and the overall model response is averaged in the end. Each of the possible combinations is plotted as a flow which is coloured by the bin corresponding to the average model response generated by that particular combination.

Usage

```
get_pdp_predictions(
  df,
  imp,
  m,
  degree = 4,
  bins = 5,
  .f_predict = predict,
  parallel = FALSE
)
```

Arguments

df	dataframe, training data
imp	dataframe, with not more then two columns one of them numeric containing importance measures and one character or factor column containing corresponding variable names as found in training data.
m	model object
degree	integer, number of top important variables to select. For plotting more than 4 will result in too many flows and the alluvial plot will not be very readable, Default: 4
bins	integer, number of bins for numeric variables, increasing this number might result in too many flows, Default: 5
.f_predict	corresponding model predict() function. Needs to accept 'm' as the first parameter and use the 'newdata' parameter. Supply a wrapper for predict functions with x-y syntax. For parallel processing the predict method of object

classes will not always get imported correctly to the worker environment. We can pass the correct predict method via this parameter for example `randomForest::predict.randomForest`. Note that a lot of modeling packages do not export the predict method explicitly and it can only be found using `:::`.

`parallel` logical, turn on parallel processing. Default: FALSE

Details

For more on partial dependency plots see [<https://christophm.github.io/interpretable-ml-book/pdp.html>].

Value

vector, predictions

Parallel Processing

We are using ‘furry’ and the ‘future’ package to paralelize some of the computational steps for calculating the predictions. It is up to the user to register a compatible backend (see [plan](#)).

Examples

```
df = mtcars2[, ! names(mtcars2) %in% 'ids' ]
m = randomForest::randomForest( disp ~ ., df)
imp = m$importance

pred = get_pdp_predictions(df, imp
                           , m
                           , degree = 3
                           , bins = 5)

# parallel processing -----
## Not run:
future::plan("multisession")

# note that we have to pass the predict method via .f_predict otherwise
# it will not be available in the worker's environment.

pred = get_pdp_predictions(df, imp
                           , m
                           , degree = 3
                           , bins = 5,
                           , parallel = TRUE
                           , .f_predict = randomForest::predict.randomForest)

## End(Not run)
```

`get_pdp_predictions_seq`*get predictions compatible with the partial dependence plotting method, sequential variant that only works for numeric predictions.*

Description

has been replaced by `pdp_predictions` which can be paralelized and also handles factor predictions. It is still used to test results.

Usage

```
get_pdp_predictions_seq(df, imp, m, degree = 4, bins = 5, .f_predict = predict)
```

Arguments

<code>df</code>	dataframe, training data
<code>imp</code>	dataframe, with not more then two columns one of them numeric containing importance measures and one character or factor column containing corresponding variable names as found in training data.
<code>m</code>	model object
<code>degree</code>	integer, number of top important variables to select. For plotting more than 4 will result in two many flows and the alluvial plot will not be very readable, Default: 4
<code>bins</code>	integer, number of bins for numeric variables, increasing this number might result in too many flows, Default: 5
<code>.f_predict</code>	corresponding model <code>predict()</code> function. Needs to accept 'm' as the first parameter and use the 'newdata' parameter. Supply a wrapper for predict functions with x-y syntax. For parallel processing the predict method of object classes will not always get imported correctly to the worker environment. We can pass the correct predict method via this parameter for example <code>randomForest::predict.randomForest</code> . Note that a lot of modeling packages do not export the predict method explicitly and it can only be found using <code>:::</code> .

See Also

[get_pdp_predictions](#)

manip_bin_numerics *bin numerical columns*

Description

centers, scales and Yeo Johnson transforms numeric variables in a dataframe before binning into n bins of equal range. Outliers based on boxplot stats are capped (set to min or max of boxplot stats).

Usage

```
manip_bin_numerics(
  x,
  bins = 5,
  bin_labels = c("LL", "ML", "M", "MH", "HH"),
  center = T,
  scale = T,
  transform = T,
  round_numeric = T,
  digits = 2,
  NA_label = "NA"
)
```

Arguments

x	dataframe with numeric variables, or numeric vector
bins	number of bins for numerical variables, passed to cut as breaks parameter, Default: 5
bin_labels	labels for the bins from low to high, Default: c("LL", "ML", "M", "MH", "HH"). Can also be one of c('mean', 'median', 'min_max', 'cuts'), the corresponding summary function will supply the labels.
center	logical, Default: T
scale	logical, Default: T
transform	logical, apply Yeo Johnson Transformation, Default: T
round_numeric	logical, rounds numeric results if bin_labels is supplied with a supported summary function name.
digits	integer, number of digits to round to
NA_label	character vector, define label for missing data, Default: 'NA'

Value

dataframe

Examples

```
summary( mtcars2 )
summary( manip_bin_numerics(mtcars2) )
summary( manip_bin_numerics(mtcars2, bin_labels = 'mean'))
summary( manip_bin_numerics(mtcars2, bin_labels = 'cuts'
, scale = FALSE, center = FALSE, transform = FALSE))
```

manip_factor_2_numeric

converts factor to numeric preserving numeric levels and order in character levels.

Description

before converting we check whether the levels contain a number, if they do the number will be preserved.

Usage

```
manip_factor_2_numeric(vec)
```

Arguments

vec vector

Value

vector

See Also

[str_detect](#)

Examples

```
fac_num = factor( c(1,3,8) )
fac_chr = factor( c('foo','bar') )
fac_chr_ordered = factor( c('a','b','c'), ordered = TRUE )

manip_factor_2_numeric( fac_num )
manip_factor_2_numeric( fac_chr )
manip_factor_2_numeric( fac_chr_ordered )
# does not work for decimal numbers
manip_factor_2_numeric(factor(c("A12", "B55", "10e4"))))
manip_factor_2_numeric(factor(c("1.56", "4.56", "8.4")))
```

`manip_get_ggplot_data` *Get ggplot data*

Description

retrieve ggplot 2 plot data from S3 and S7 ($\geq$ ggplot2 v4.0)

Usage

`manip_get_ggplot_data(p)`

Arguments

`p` ggplot object

`mtcars2` *mtcars dataset with cyl, vs, am, gear, carb as factor variables and car model names as id*

Description

mtcars dataset with cyl, vs, am, gear, carb as factor variables and car model names as id

Usage

`mtcars2`

Format

A data frame with 32 rows and 12 variables

mpg Miles/(US) gallon

cyl Number of cylinders

disp Displacement (cu.in.)

hp Gross horsepower

drat Rear axle ratio

wt Weight (1000 lbs)

qsec 1/4 mile time

vs Engine

am Transmission

gear Number of forward gears

carb Number of carburetors

ids car model name

Source

datasets

palette_filter	<i>color filters for any vector of hex color values</i>
----------------	---

Description

filters are based on rgb values

Usage

```
palette_filter(
  palette = palette_qualitative(),
  similar = F,
  greys = T,
  reds = T,
  greens = T,
  blues = T,
  dark = T,
  medium = T,
  bright = T,
  thresh_similar = 25
)
```

Arguments

palette	any vector with hex color values, Default: palette_qualitative()
similar	logical, allow similar colours, similar colours are detected using a threshold (thresh_similar), two colours are similar when each value for RGB is within threshold range of the corresponding RGB value of the second colour, Default: F
greys	logical, allow grey colours, blue == green == blue , Default: T
reds	logical, allow red colours, blue < 50 & green < 50 & red > 200 , Default: T
greens	logical, allow green colours, green > red & green > blue, Default: T
blues	logical, allow blue colours, blue > green & green > red, Default: T
dark	logical, allow colours of dark intensity, sum(red, green, blue) < 420 , Default: T
medium	logical, allow colours of medium intensity, between(sum(red, green, blue), 420, 600) , Default: T
bright	logical, allow colours of bright intensity, sum(red, green, blue) > 600, Default: T
thresh_similar	int, threshold for defining similar colours, see similar, Default: 25

Value

vector with hex colors

Examples

```
require(magrittr)

palette_qualitative() %>%
  palette_filter(thresh_similar = 0) %>%
  palette_plot_intensity()

## Not run:
# more examples-----

palette_qualitative() %>%
  palette_filter(thresh_similar = 25) %>%
  palette_plot_intensity()

palette_qualitative() %>%
  palette_filter(thresh_similar = 0, blues = FALSE) %>%
  palette_plot_intensity()

## End(Not run)
```

palette_increase_length

increases length of palette by repeating colours

Description

works for any vector

Usage

```
palette_increase_length(palette = palette_qualitative(), n = 100)
```

Arguments

palette	any vector, Default: palette_qualitative()
n	int, length, Default: 100

Value

vector with increased length

Examples

```
require(magrittr)

length(palette_qualitative())

palette_qualitative() %>%
  palette_increase_length(100) %>%
  length()
```

palette_plot_intensity
plot colour intensity of palette

Description

sum of red green and blue values

Usage

```
palette_plot_intensity(palette)
```

Arguments

palette any vector containing color hex values

Value

ggplot2 plot

See Also

[palette_plot_rgp](#)

Examples

```
## Not run:
if(interactive()){
  palette_qualitative() %>%
    palette_filter( thresh = 25) %>%
    palette_plot_intensity()
}

## End(Not run)
```

palette_plot_rgp	<i>plot rgb values of palette</i>
------------------	-----------------------------------

Description

grouped bar chart

Usage

```
palette_plot_rgp(palette)
```

Arguments

palette any vector containing color hex values

Value

ggplot2 plot

See Also

[palette_plot_intensity](#)

Examples

```
## Not run:
if(interactive()){
  palette_qualitative() %>%
    palette_filter( thresh = 50) %>%
    palette_plot_rgp()
}

## End(Not run)
```

palette_qualitative	<i>compose palette from qualitative RColorBrewer palettes</i>
---------------------	---

Description

uses `c('#FF0065', '#009850', '#A56F2B', '#005EAA', '#710500', '#7B5380', '#9DD1D1')` and then adds all unique values found in all qualitative RColorBrewer palettes

Usage

```
palette_qualitative()
```

Value

vector with hex values

See Also

[RColorBrewer](#)

Examples

```
palette_qualitative()
```

plot_all_hists	<i>plot marginal histograms of alluvial plot</i>
----------------	--

Description

will create gtable with density histograms and frequency plots of all variables of a given alluvial plot.

Usage

```
plot_all_hists(p, data_input, top = TRUE, keep_labels = FALSE, ...)
```

Arguments

p	alluvial plot
data_input	dataframe, input data that was used to create dataframe
top	logical, position of histograms, if FALSE adds them at the bottom, Default: TRUE
keep_labels	logical, keep title and caption, Default: FALSE
...	additional arguments for specific alluvial plot types: pred_train can be used to pass training predictions for model response alluvials

Value

gtable

See Also

[arrangeGrob](#)

[add_marginal_histograms](#)

Examples

```
## Not run:
p = alluvial_wide(mtcars2, max_variables = 3)
plot_all_hists(p, mtcars2)

## End(Not run)
```

plot_condensation	<i>Plot dataframe condensation potential</i>
-------------------	--

Description

plotting the condensation potential is meant as a decision aid for which variables to include in an alluvial plot. All variables are transformed to categoric variables and then two variables are selected by which the dataframe will be grouped and summarized by. The pair that results in the greatest condensation of the original dataframe is selected. Then the next variable which offers the greatest condensation potential is chosen until all variables have been added. The condensation in percent is then plotted for each step along with the number of groups (flows) in the dataframe. By experience it is not advisable to have more than 1500 flows because then the alluvial plot will take a long time to render. If there is a particular variable of interest in the dataframe this variable can be chosen as a starting variable.

Usage

```
plot_condensation(df, first = NULL)
```

Arguments

df	dataframe
first	unquoted expression or string denoting the first variable to be picked for condensation, Default: NULL

Value

ggplot2 plot

See Also

[quosure reexports RColorBrewer](#)

Examples

```
plot_condensation(mtcars2)

plot_condensation(mtcars2, first = 'disp')
```

plot_hist	<i>plot histogram of alluvial plot variable</i>
-----------	---

Description

helper function used by add_marginal_histograms

Usage

```
plot_hist(var, p, data_input, ...)
```

Arguments

var	character vector, variable name
p	alluvial plot
data_input	dataframe used to create alluvial plot
...	additional arguments for specific alluvial plot types: pred_train can be used to pass training predictions for model response alluvials

Value

ggplot object

plot_imp	<i>plot feature importance</i>
----------	--------------------------------

Description

plot important features of model response alluvial as bars

Usage

```
plot_imp(p, data_input, truncate_at = 50, color = "darkgrey")
```

Arguments

p	alluvial plot
data_input	dataframe used to generate alluvial plot
truncate_at	integer, limit number of features to that value, Default: 50
color	character vector, Default: 'darkgrey'

Value

ggplot object

Examples

```
## Not run:
df = mtcars2[, ! names(mtcars2) %in% 'ids' ]

train = caret::train( disp ~ .
                      , df
                      , method = 'rf'
                      , trControl = caret::trainControl( method = 'none' )
                      , importance = TRUE )

pred_train = caret::predict.train(train, df)

p = alluvial_model_response_caret(train, degree = 3, pred_train = pred_train)

plot_imp(p, mtcars2)

## End(Not run)
```

quarterly_flights

Quarterly mean arrival delay times for a set of 402 flights

Description

Created from nycflights13::flights

Usage

```
quarterly_flights
```

Format

A data frame with 1608 rows and 6 variables

tailnum a unique identifier created from tailnum, origin, destination and carrier

carrier carrier code

origin origin code

dest destination code

qu quarter

mean_arr_delay average delay on arrival as either on_time or late

Source

nycflights13::flights

quarterly_sunspots	<i>Quarterly mean relative sunspots number from 1749-1983</i>
--------------------	---

Description

Quarterly mean relative sunspots number from 1749-1983

Usage

```
quarterly_sunspots
```

Format

A data frame with 940 rows and 4 variables

year

qu quarter

spots total number of sunspots

mean_spots_per_year

Source

Andrews, D. F. and Herzberg, A. M. (1985) Data: A Collection of Problems from Many Fields for the Student and Research Worker. New York: Springer-Verlag.

tidy_imp	<i>tidy up dataframe containing model feature importance</i>
----------	--

Description

returns dataframe with exactly two columns, vars and imp and aggregates dummy encoded variables. Helper function called by all functions that take an imp parameter. Can be called manually if formula for aggregating dummy encoded variables must be modified.

Usage

```
tidy_imp(imp, df, .f = max, resp_var = NULL)
```

Arguments

imp	dataframe or matrix with feature importance information
df	dataframe, modeling training data
.f	window function, Default: max
resp_var	character, prediction variable, can usually be inferred from imp and df. It does not work for all models and needs to be specified in those cases.

Value

dataframe

vars character column with feature names

imp numerical column, importance values

Examples

```
# randomforest
df = mtcars2[, ! names(mtcars2) %in% 'ids' ]
m = randomForest::randomForest( disp ~ ., df)
imp = m$importance
tidy_imp(imp, df)
```

titanic	<i>titanic data set</i>
---------	-------------------------

Description

titanic data set'

Usage

titanic

Format

A data frame with 891 rows and 10 variables

Survived Survived

Pclass Pclass

Sex Sex

Age Age

SibSp SibSp

Parch Parch

Fare Fare

Cabin Cabin

Embarked Embarked

title title

Source

datasets

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