Package 'easyalluvial'

April 1, 2019

Title Generate Alluvial Plots with a Single Line of Code

Version 0.2.0

URL https://github.com/erblast/easyalluvial

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> Their graphical grammar however is a bit more complex then 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(>= 2.10)

Suggests testthat, covr, ISLR, nycflights13, vdiffr

RoxygenNote 6.1.1

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R topics documented:

`add_imp_plot` *add bar plot of important features to model response alluvial plot*

Description

adds bar plot of important features to model respoinse 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

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
plot	logical if plot should be drawn or not
...	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
	pred_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

```
p = alluvial_wide(mtcars2, max_variables = 4)
p_grid = add_marginal_histograms(p, mtcars2)
```

---

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_size = 4.5,
  stratum_width = 1/4, auto_rotate_xlabs = T, ...)
```

Arguments

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

stratum_width double, Default: 1/4

auto_rotate_xlabs logical, Default: TRUE

... additional parameter passed to manip_bin_numerics

Value

ggplot2 object

See Also

alluvial_wide ,geom_flow, geom_stratum ,manip_bin_numerics

Examples

```
 data = quarterly_flights

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

## Not run:
 # 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, bins = 5,
  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(center = T, transform = T, scale = T),
  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 two 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 variableas 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, 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",
  params_bin_numeric_pred = list(center = T, transform = T, scale = T),
  pred_train = NULL, stratum_label_size = 3.5, force = F, ...)
```

Arguments

`train`	caret train object
`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, 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 variableas are set to the indicated values. The predict function is called for each modified observation and the result is averaged . Default: 'median'

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

...
: 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, varImp, extractPrediction, get_data_space, get_pdp_predictions

Examples

```
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, degree = 3)

# partial dependency plotting method
## Not run:
alluvial_model_response_caret(train, degree = 3, method = 'pdp')

## 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_size = 4.5, stratum_width = 1/4,
  auto_rotate_xlabs = T, ...)
```

Arguments

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

```
alluvial_wide( data = mtcars2, id = ids
                , max_variables = 5
                , fill_by = 'first_variable' )
## Not run:

# 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)
```

---

get_data_space *calculate data space*

Description

calculates a dataspace based on the modelling 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)
```

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, increasing this number might result in too many flows, Default: 5

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` *get predictions compatibel with the partial dependence plotting method*

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)
```

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 two 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 synthax.

Details

see https://christophm.github.io/interpretable-ml-book/pdp.html

Value

vector, predictions

See Also

progress_bar

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)
```

---

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, 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 corresppnding 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 )
```

`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 *color filters for any vector of hex color values*

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 *plot rgb values of palette*

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 *compose palette from qualitative RColorBrewer palettes*

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 *plot marginal histograms of alluvial plot*

Description

will create gtable with densitiy 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

```
p = alluvial_wide(mtcars2, max_variables = 4)
plot_all_hists(p, mtcars2)
```

`plot_condensation` *Plot dataframe condensation potential*

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 *plot histogram of alluvial plot variable*

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	datafram 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 *plot feature importance*

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 *Quarterly mean relative sunspots number from 1749-1983*

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 *tidy up dataframe containing model feature importance*

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)
```

Arguments

imp	dataframe or matrix with feature importance information
df	dataframe, modelling training data
.f	window function, Default: max

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, mtcars2)
```

---

use_e1071 *calls e1071::skewness*

Description

if e1071 is not listed a a dependency I get an error. I assume caret uses it to calculate feature importance. However, e1071 is not listed as a caret dependency. I have to add a function that directly calls it, so I do not get a NOTE from RMD Check on Linux.

Usage

```
use_e1071(x)
```

Arguments

x	PARAM_DESCRIPTION

Value

OUTPUT_DESCRIPTION

See Also

skewness

Index