Package ‘ggasym’

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diagonal. It operates within the Grammar of Graphics paradigm implemented
in 'ggplot2'.
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add_missing_combinations

Add missing combinations of \( x \) and \( y \) to a data frame

Description

Add rows to \( df \) to complete all combinations of columns \( .x \) and \( .y \). Importantly, this function observes and maintains any groups created by \texttt{dplyr::group_by()}.

Usage

\texttt{add_missing_combinations(df, .x, .y)}

Arguments

- \texttt{df} a data frame (or tibble) object
- \texttt{.x, .y} column names to make combinations of

Value

a data frame (or tibble) with additional columns
Examples

df <- data.frame(
  a = c("A", "B"),
  b = c("C", "D"),
  untouched = c(1, 2)
)
df

add_missing_combinations(df, a, b)

---

asymmetrise  
Add all missing comparisons between two columns

Description

This function prepares input data for `geom_asymmat()` by adding in any missing comparisons to be plotted. Note that this function observes groups created with the `dplyr::group_by()` function. For the 'ggasym' package, this is useful for when you want to facet the plot: before "asymmetrizing" the data table, use `dplyr::group_by()`, passing the column name you wish to later facet by. This functionality is demonstrated in the second example, below.

Usage

asymmetrise(df, .x, .y)
asymmetrize(df, .x, .y)

Arguments

df  
a tidy data.frame or tibble
.x, .y  
the data to add all comparisons between (ie. will be the x and y-axes for `geom_asymmat()`)

Value

a data table with new rows for the added comparisons

Warning

This function does it’s best when x or y are factors. If they have the same levels, then they are maintained. If the levels partially overlap, they are merged. Otherwise, the values are turned into characters and all levels dropped. If you are using factors, save yourself the headache and make both columns factors with the desired levels.
Examples

```r
df <- data.frame(
  a = c("A", "B", "C"),
  b = c("C", "A", "B"),
  untouched = c(1, 2, 3),
  grouping_value = c("group1", "group1", "group2"),
  stringsAsFactors = FALSE
)
df

asymmetrise(df, a, b)

grouped_df <- dplyr::group_by(df, grouping_value)
asymmetrise(grouped_df, a, b)
```

---

**asymmetrise_stats**

*Prepare an asymmetric data table from a statistical test*

**Description**

This function prepares the results of a statistical test for plotting using 'geom_asymmat' from the ggasym package. For more information, see vignette(ggasym-stats)

**Usage**

```r
asymmetrise_stats(df, contrast_sep = "-")

asymmetrize_stats(df, contrast_sep = "-")
```

**Arguments**

- `df` either the results of a statistical test or the tidy tibble from using the broom::tidy() function
- `contrast_sep` the separation used between the names being compared; it is usually a hyphen (set as default here); since it is passed as the pattern parameter to stringr::str_split_fixed(), this can be any regular expression that will reliably split the "contrast" (or "comparison" in 'broom' version <0.70) column returned by broom::tidy().

**Value**

a tibble object that can be used as direct input for 'ggplot2' for use with the geom_asymmat geom
bind_missing_combs

Add the missing combinations of x and y

Description

Adds rows to the input data table to include any combinations of x and y that are not already present. All other columns (if any) are set to NA.

Usage

bind_missing_combs(df, .x, .y)

Arguments

df input data table
.x, .y names of the columns for which to add missing comparisons

Value

a data table with the new rows

Examples

df <- data.frame(
a = c("A", "B"),
b = c("C", "A"),
untouched = c(1, 2),
stringsAsFactors = FALSE
)
df

bind_missing_combs(df, a, b)

continuous_scale_asym Continuous scale constructor for 'ggasym'

Description

This is a this wrapper around continuous_scale() from the 'ggplot2' package. It is generally best to call this function implicitly using one of the wrappers that have the general naming scheme of scale_*_tl/br_*() (such as scale_fill_tl_gradient()).

Usage

continuous_scale_asym(aesthetics, scale_name, palette, na.value, guide, ...)
Arguments

- **aesthetics**: The names of the aesthetics that this scale works with.
- **scale_name**: The name of the scale.
- **palette**: A palette function that when called with a numeric vector with values between 0 and 1 returns the corresponding values in the range the scale maps to.
- **na.value**: Missing values will be replaced with this value.
- **guide**: A function used to create a guide or its name. See guides() for more info.
- **...**: Other input is passed on to ggplot2::continuous_scale(); see ?ggplot2::continuous_scale for complete documentation.

Examples

```r
library(tibble)
library(ggplot2)
tib <- tibble(
  g1 = c("A", "A", "B"),
  g2 = c("B", "C", "C"),
  val_1 = c(1, 2, 3),
  val_2 = c(-1, 0, 1)
)

... 

ggplot(tib) +
  geom_asymmat(aes(x = g1, y = g2, fill_tl = val_1, fill_br = val_2)) +
  scale_fill_tl_gradient(low = "lightpink", high = "tomato") +
  scale_fill_br_gradient(low = "lightblue1", high = "dodgerblue") +
  labs(fill_tl = "top-left fill", fill_br = "bottom-right fill")
```

---

**factor_is_greater**

*Determines if the level of a is greater than that of b*

Description

Determines if the level of a is greater than that of b.

Usage

`factor_is_greater(a, b)`

Arguments

- **a, b**: Two same-length, same-leveled vectors of type factor.

Value

A single boolean vector of type logical.
## Examples

```r
first <- c("J", "O", "S", "H")
last <- c("C", "O", "O", "K")
first <- factor(first, LETTERS)
last <- factor(last, LETTERS)
factor_is_greater(first, last)
```

## Description

A `ggproto` object for the `ggasym` package and used by `geom_asymmat()`

## Usage

`GeomAsymmat`

## Format

An object of class `GeomAsymmat` (inherits from `GeomRect`, `Geom`, `ggproto`, `gg`) of length 7.

## Warning

`GeomAsymmat` is subject to change in future versions. Use at your own risk. If dependent on `GeomAsymmat`, it is advisable to include tests with a cached version to test for equivalence.

## Description

Generate an asymmetric matrix with different fill values for top-left and bottom-right triangles and along the diagonal as a `ggplot()` object

## Usage

```r
geom_asymmat(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  ...,
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE
)
```
Arguments

mapping Set of aesthetic mappings created by aes() or aes_(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.

data The data to be displayed in this layer. There are three options: If NULL (the default) the data is inherited from the plot data as specified in the call to ggplot(). A data frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created. A function will be called with a single argument, the plot data. The return value must be a data frame, and will be used as the layer data.

stat The statistical transformation to use on the data for this layer, as a string.

position Position adjustment, either as a string, or the result of a call to a position adjustment function.

... Other arguments passed on to layer(). These are often aesthetics, used to set an aesthetic to a fixed value, like colour = "red" or size = 3. They may also be parameters to the paired geom/stat.

na.rm If FALSE, the default, missing values are removed with a warning. If TRUE, missing values are silently removed.

show.legend logical. Should this layer be included in the legends? NA (the default) includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes. It can also be a named logical vector to finely select the aesthetics to display.

inherit.aes If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and should not inherit behaviour from the default plot specification, e.g. borders().

Examples

library(tibble)
library(ggplot2)
suppressMessages(library(dplyr))
tib <- tibble(
  g1 = c("A", "A", "B"),
  g2 = c("B", "C", "C"),
  val_1 = c(1, 2, 3),
  val_2 = c(-1, 0, 1)
)
tib

tib <- asymmetrise(tib, g1, g2)
tib$val_3 <- NA
tib$val_3[tib$g1 == tib$g2] <- c(1, 2, 3)
ggplot(tib, aes(x = g1, y = g2)) +
  geom_asymmat(aes(fill_tl = val_1, fill_br = val_2, fill_diag = val_3)) +
  scale_fill_br_gradient(low = "lightblue1", high = "dodgerblue") +
  scale_fill_tl_gradient(low = "lightpink", high = "tomato") +
  scale_fill_diag_gradient(low = "aquamarine", high = "forestgreen") +
  labs(fill_tl = "top-left fill", fill_br = "bottom-right fill")
get_other_combs

Description

Get all combinations of the values between two vectors.

Usage

get_other_combs(x, y)

Arguments

x, y
two vectors

Value

data.frame of other possible combinations stored in Var1 and Var2 for x and y, respectively

Examples

get_other_combs(LETTERS[1:2], LETTERS[1:2])

ggasym

ggasym: Asymmetric Matrix Plotting in ggplot

Description

This package plots a symmetric matrix with two different fill aesthetics for the top-left and bottom-right triangles and a third along the diagonal. It operates within the Grammar of Graphics paradigm implemented in `ggplot2`.

is_grouped

Is a data table grouped?

Description

Determines if the input data frame or tibble is grouped (using dplyr::group_by())

Usage

is_grouped(data)

Arguments

data input data.frame or tibble

Value

boolean

Examples

df <- data.frame(x = c(1:5), g = c(1, 1, 2, 2, 2))
is_grouped(df)
is_grouped(dplyr::group_by(df, g))

make_fill_df

Make a data frame of all a single value

Description

Makes a data frame with the same columns of df and n_rows number of rows and all values fill_val

Usage

make_fill_df(df, n_rows = 1, fill_val = NA)

Arguments

df a data.frame (or tibble) object
n_rows number of rows for the final data frame
fill_val value to fill all cells of the data frame

Value

a data frame (or tibble) with the desired number of rows filled with fill_val
Examples

df <- data.frame(
    col_a = c("A", "B"),
    col_b = c("C", "D")
)
df

make_fill_df(df, 5)

organize_levels

Decides on the levels of factors x and y

Description

Organizes the levels to use for the two inputs. This is useful for when one wants to merge two vectors that are factors. Ideally, they have the same levels, in which case those are returned. If they have overlapping levels, then the levels are merged and sorted (using sort()). Otherwise, the levels are dropped (returning NULL)

Usage

organize_levels(x, y, …)

Arguments

  x, y
  Two factor vectors

  …
  passed to sort; see ?sort for options

Value

vector of levels or NULL for no levels

Examples

set.seed(0)
a <- factor(sample(LETTERS, 5), levels = LETTERS)
b <- factor(sample(LETTERS, 5), levels = LETTERS)
a
b
organize_levels(a, b)
**prepare_data**

*Prepares the input data into asymmetrise_stats*

**Description**

Tries to make the data ready for use in the asymmetrise_stats() function using broom::tidy()

**Usage**

prepare_data(df)

**Arguments**

- **df**
  
in input data of either a tibble, data.frame, or results from a statistical test

**Value**

a tibble data table

**Warning**

If you repeatedly get errors, try preparing the data before-hand using broom::tidy(df)

**Examples**

```r
a <- rnorm(10, mean = 1, sd = 1)
b <- rnorm(10, mean = 1.5, sd = 1)
prepare_data(t.test(a, b))
```

---

**scale_gradient**

*Gradient colour scales geom_asymmat*

**Description**

This dictates a gradient colour scheme for the top-left (tl), bottom_right (br), or diagonal (diag) of a geom_asymmat() geom. scale_*_tl/br_gradient() creates a two colour gradient (low-high), scale_*_tl/br_gradient2() creates a diverging colour gradient (low-mid-high), scale_*_tl/br_gradientn() creates a n-colour gradient.
Usage

```r
scale_fill_tl_gradient(
  ..., 
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_tl"
)

scale_fill_br_gradient(
  ..., 
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_br"
)

scale_fill_diag_gradient(
  ..., 
  low = "#132B43",
  high = "#56B1F7",
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_diag"
)

scale_fill_tl_gradient2(
  ..., 
  low = scales::muted("red"),
  mid = "white",
  high = scales::muted("blue"),
  midpoint = 0,
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_tl"
)

scale_fill_br_gradient2(
  ..., 
  low = scales::muted("red"),
  mid = "white",
  high = scales::muted("blue"),
scale_gradient

midpoint = 0,
space = "Lab",
na.value = "grey50",
guide = "colourbar",
aesthetics = "fill_br"
)

scale_fill_diag_gradient2(
...
low = scales::muted("red"),
mid = "white",
high = scales::muted("blue"),
midpoint = 0,
space = "Lab",
na.value = "grey50",
guide = "colourbar",
aesthetics = "fill_diag"
)

scale_fill_tl_gradientn(
...
colours,
values = NULL,
space = "Lab",
na.value = "grey50",
guide = "colourbar",
aesthetics = "fill_tl",
colors
)

scale_fill_br_gradientn(
...
colours,
values = NULL,
space = "Lab",
na.value = "grey50",
guide = "colourbar",
aesthetics = "fill_br",
colors
)

scale_fill_diag_gradientn(
...
colours,
values = NULL,
space = "Lab",
na.value = "grey50",
guide = "colourbar",

scale_gradient

aesthetics = "fill_diag",
  colors
)

scale_fill_tl_distiller(
  ...,  
  type = "seq",
  palette = 1,
  direction = -1,
  values = NULL,
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_tl"
)

scale_fill_br_distiller(
  ...,  
  type = "seq",
  palette = 1,
  direction = -1,
  values = NULL,
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_br"
)

scale_fill_diag_distiller(
  ...,  
  type = "seq",
  palette = 1,
  direction = -1,
  values = NULL,
  space = "Lab",
  na.value = "grey50",
  guide = "colourbar",
  aesthetics = "fill_diag"
)

Arguments

  ...  arguments passed on to continuous_scale_asym()
  low, high  the colors to represent low and high values
  space  colour space in which to calculate gradient. Must be "Lab" - other values are deprecated.
  na.value  color of missing (NA) values
**guide**
Type of legend. Use "colourbar" for continuous colour bar, or "legend" for discrete colour legend.

**aesthetics**
Character string or vector of character strings listing the name(s) of the aesthetic(s) that this scale works with. For now, leave the default alone, though I plan to reinstate the standard 'ggplot2' system here, eventually.

**mid**
color for mid point (see ?scales::div_gradient_pal for more documentation of how colors are calculated)

**midpoint**
The midpoint (in data value) of the diverging scale. Defaults to 0.

**colours, colors**
Vector of colours to use for n-colour gradient.

**values**
if colours should not be evenly positioned along the gradient this vector gives the position (between 0 and 1) for each colour in the colours vector. See rescale() for a convenience function to map an arbitrary range to between 0 and 1.

**type**
One of "seq" (sequential), "div" (diverging) or "qual" (qualitative)

**palette**
If a string, will use that named palette. If a number, will index into the list of palettes of appropriate type

**direction**
Sets the order of colours in the scale. If 1, the default, colours are as output by RColorBrewer::brewer_pal(). If -1, the order of colours is reversed.

---

**Examples**

```r
library(tibble)
library(ggplot2)
set.seed(0)

tib <- tibble(
val_1 = c(1:10),
val_2 = sample(-10:10, 10),
val_3 = c(rep(NA, 6), 1, 2, 3, 4)
)
tib <- asymmetrise(tib, g1, g2)
g <- ggplot(tib, aes(x = g1, y = g2)) +
  geom_asymmat(aes(fill_tl = val_1, fill_br = val_2, fill_diag = val_3))
g + scale_fill_tl_gradient(low = "lightpink", high = "tomato") +
  scale_fill_br_gradient(low = "lightblue", high = "dodgerblue") +
  scale_fill_diag_gradient(low = "yellow", high = "orange3")
g + scale_fill_tl_gradient2(
  low = "dodgerblue",
  mid = "white", midpoint = 5,
  high = "tomato"
) +
  scale_fill_br_gradient2(
  low = "seagreen4",
  mid = "white", midpoint = 0,
  high = "orange"
```

swap_cols

**Description**

Swap columns `.x` and `.y` in `df`.

**Usage**

```r
swap_cols(df, .x, .y)
```

**Arguments**

- `df` a data.frame (or tibble) object
- `.x`, `.y` column names to switch

**Value**

a data.frame (or tibble) object with `.x` and `.y` swapped

**Examples**

```r
df <- data.frame(
a = c("A", "B"),
b = c("C", "D"),
untouched = c(1, 2)
)
df

swap_cols(df, a, b)
```
which_level

Determine the level of a value in a vector of type factor

Description

Determine the level of a value in a vector of type factor

Usage

which_level(x)

Arguments

x vectors of type factor

Value

a vector holding the corresponding level of the input factors

Examples

first <- factor(c("J", "O", "S", "H"), LETTERS)
which_level(first)
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