

Package: ggtilematrix (via r-universe)

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Title Create customisable ggplot2 matrix

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Description More about what it does (maybe more than one line) Use
four spaces when indenting paragraphs within the Description.

License GPL (>= 3)

Encoding UTF-8

LazyData true

Suggests testthat (>= 3.0.0)

Config/testthat.edition 3

Depends R (>= 2.10)

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Imports dplyr, ggplot2, tibble, tidyverse, tidyselect

Repository https://cynthiahqy.r-universe.dev

RemoteUrl https://github.com/cynthiahqy/ggtilematrix

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`gg_tilematrix.matrix` *Plot a matrix as a ggplot2 object*

Description

Plot a matrix as a ggplot2 object

Usage

```
gg_tilematrix.matrix(
  .matrix,
  .geom = list(geom_tile(color = "pink", fill = "blue"), geom_text()),
  .scale_coord = list(scale_y_discrete(limits = base::rev), scale_x_discrete(position =
    "top"), coord_fixed()),
  .theme = list(theme_bw()),
  .layers = list(labs(x = NULL, y = NULL))
)
```

Arguments

.matrix	A matrix to plot
.layers	

Examples

```
library(ggplot2)
matrices$unnamed |>
  gg_tilematrix.matrix()
```

`matrix_to_triples` *Pivots a matrix into (bi-graph) triples*

Description

Converts a named (adjacency) matrix into (graph) triples, dropping any NA cells by default.

Usage

```
matrix_to_triples(
  matrix,
  x_names_to = "x_name",
  y_names_to = "y_name",
  values_to = "value",
  drop_na = TRUE
)
```

```
triples_from_matrix(
  matrix,
  x_names_to = "x_name",
  y_names_to = "y_name",
  values_to = "value",
  drop_na = TRUE
)
```

Arguments

<code>matrix</code>	A matrix to pivot
<code>x_names_to</code>	A string specifying the new column for the names of the first/x-dimension of <code>matrix</code>
<code>y_names_to</code>	A string specifying the new column for the names of the second/y-dimension <code>matrix</code>
<code>values_to</code>	A character vector specifying the new column to create from the cell values of <code>matrix</code>
<code>drop_na</code>	Boolean option to drop x-y pairs with missing values.

Details

Uses `pivot_longer` and reverses `triples_to_matrix()`. Setting `drop_na=FALSE` retain the complete set of graph edges. If `matrix` is unnamed, default names are generated by `base::as.data.frame()`

Examples

```
## named matrices
blx_mtx <- triples$blx_tbl |>
  triples_to_matrix(from, to, weighted)
return_triples <- blx_mtx |>
  triples_from_matrix("from", "to", "weighted")

identical(triples$blx_tbl, return_triples)

## Setting `drop_na=FALSE` retains the `x`-`y` pairs with missing values,
return_with_na <- blx_mtx |>
  triples_from_matrix("from", "to", "weighted", drop_na = FALSE)

## which is equivalent to using `tidyr::complete()` on the original triples
complete_triples <- triples$blx_tbl |>
  tidyr::complete(from, to)

identical(return_with_na, complete_triples)

## names are generated for unnamed matrices
matrices$unnamed |>
  matrix_to_triples(drop_na = FALSE)
```

triples_to_matrix *Pivot (bi-graph) triples into (bi-adjacency) matrix*

Description

Converts a data frame of triples (`x_names`, `y_names`, `values`) into a matrix with `x_names` and `y_names` respectively for row and column names and cell values from `values`.

Usage

```
triples_to_matrix(
  triples,
  x_names_from = x_name,
  y_names_from = y_name,
  values_from = value,
  values_fill = NA
)

matrix_from_triples(
  triples,
  x_names_from = x_name,
  y_names_from = y_name,
  values_from = value,
  values_fill = NA
)
```

Arguments

<code>triples</code>	A data frame to pivot. Unused columns are dropped.
<code>x_names_from</code>	column to get matrix row names from.
<code>y_names_from</code>	column to get matrix column names from.
<code>values_from</code>	column to get cell values from.
<code>values_fill</code>	Optionally, a (scalar) value that specifies what each value should be filled in with when missing. This can be a named list if you want to apply different fill values to different value columns.

Details

The triples define graph edges between `x_names` and `y_names` with attribute `value`. `x_names` and `y_names` are treated as disjoint nodes sets of the bi-partite graph, and the resultant matrix corresponds to the bi-adjacency matrix.

Uses [pivot_wider](#) and makes explicit all possible pairs of `x_names` and `y_names`.

Examples

```
triples$blx_tbl  
triples$blx_tbl |>  
  triples_to_matrix(from, to, weighted)
```

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