

Package: varnisch (via r-universe)

August 25, 2024

Title What the Package Does (One Line, Title Case)

Version 0.0.0.9000

Description What the package does (one paragraph).

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URL <https://igordot.github.io/varnisch/>

Depends ggplot2, R (>= 4.1)

Imports cowplot, FNN, ggsci, methods, RColorBrewer, rlang,
scattermore, tibble

Suggests SeuratObject, SingleCellExperiment, tidyseurat,
tidySingleCellExperiment

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.3.1

Repository <https://igordot.r-universe.dev>

RemoteUrl <https://github.com/igordot/varnisch>

RemoteRef HEAD

RemoteSha 484a14eacc5142a881d4f11d101a90d9ca53a579

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|----------------|--|
| flatten_object | <i>Convert a specialized single-cell data object to a data frame</i> |
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Description

Turn a specialized single-cell data object ([Seurat](#) or [SingleCellExperiment](#)) into a more simplified "tidy" data frame. In tidy data, each column is a variable and each observation is a row.

Usage

```
flatten_object(x, features = NULL, assay = "RNA", slot = "data", n_dims = 3)
```

Arguments

| | |
|----------|--|
| x | An object. |
| features | A vector of features/genes to include in the data frame. |
| assay | The object assay name for feature abundance. |
| slot | The object slot name for feature abundance. |
| n_dims | An integer scalar specifying the maximum number of dimensions to return. |

Value

A tibble, which is a data frame with class `tbl_df`.

| | |
|--------------|--|
| plot_scatter | <i>Generate a generic scatter plot</i> |
|--------------|--|

Description

Create a scatter plot to display the relationship between two continuous variables. This is essentially a wrapper for `ggplot2::geom_point()`.

Usage

```
plot_scatter(  
  data,  
  x,  
  y,  
  color_by,  
  smooth = FALSE,  
  range = c(0.01, 0.99),  
  title = "",  
  aspect_ratio = 1  
)
```

Arguments

| | |
|---------------------------|--|
| <code>data</code> | A data frame. |
| <code>x</code> | . |
| <code>y</code> | . |
| <code>color_by</code> | Column metadata field(s) or feature(s) to color by. |
| <code>smooth</code> | A logical scalar. Smooth values. Helps to visualize expression patterns in a plot with many overlapping points. |
| <code>range</code> | A vector of 2 values indicating the minimum and maximum percentiles for the color range. Helps to visualize expression patterns when extreme outliers are present. For example, <code>c(0, 0.99)</code> will not expand the color scale above 99th percentile. |
| <code>title</code> | Plot title. |
| <code>aspect_ratio</code> | Aspect ratio of the panel. |

Value

A ggplot object.

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