

# Package ‘aplotExtra’

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**Title** Creating Composite Plots using 'aplot'

**Version** 0.0.4

**Description** Many complex plots are actually composite plots, such as 'oncoplot', 'funkyheatmap', 'upsetplot', etc. We can produce subplots using 'ggplot2' and combine them to create composite plots using 'aplot'. In this way, it is easy to customize these complex plots, by adding, deleting or modifying subplots in the final plot. This package provides a set of utilities to help users to create subplots and complex plots.

**Depends** R (>= 4.1.0)

**Imports** aplot (>= 0.2.3), dplyr,forcats, ggfun (>= 0.1.1), ggplot2, grid, maftools, purrr, rlang, stats, tibble, tidyR, utils, ggstar, yulab.utils (>= 0.0.8)

**Suggests** ggtree, data.table, RColorBrewer, R.utils

**URL** <https://github.com/YuLab-SMU/aplotExtra>

**License** Artistic-2.0

**Encoding** UTF-8

**RoxygenNote** 7.3.2

**NeedsCompilation** no

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funky\_bar                    *funky\_bar*

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**Description**

create bar plot for funkyheatmap

**Usage**

```
funky_bar(data, cols)
```

**Arguments**

data	data frame
cols	selected columns

**Value**

ggplot object

**Author(s)**

Guangchuang Yu

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funky\_heatmap                    *funky\_heatmap*

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**Description**

create a funkyheatmap

**Usage**

```
funky_heatmap(..., data = NULL, widths = NULL, options = NULL)
```

**Arguments**

...	funky plots (e.g., outputs of <code>funky_point</code> , <code>funky_bar</code> , etc.)
<code>data</code>	If data is provided, create a funkyheatmap from it. Otherwise, create composite plot from ...
<code>widths</code>	relative widths of the plots
<code>options</code>	any ggplot component that can be added to the plots

**Value**

gglister object

**Author(s)**

Guangchuang Yu

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`funky_point`

*funky\_point*

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**Description**

create dot plot for funkyheatmap

**Usage**

`funky_point(data, cols, cols2 = NULL, ...)`

**Arguments**

<code>data</code>	data frame
<code>cols</code>	selected columns
<code>cols2</code>	selected columns to keep names
...	additional parameters, passing to <code>geom_star</code>

**Value**

ggplot object

**Author(s)**

Guangchuang Yu

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funky_text	<i>funky_text</i>
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**Description**

create text plot (i.e., rownames) for funkyheatmap

**Usage**

```
funky_text(data, cols = 1, hjust = 0)
```

**Arguments**

data	data frame
cols	selected column
hjust	text alignment adjustment

**Value**

ggplot object

**Author(s)**

Guangchuang Yu

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get_all_subsets	<i>Get the items/names/ids of subsets from a named list</i>
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**Description**

Get the items/names/ids of subsets from a named list

**Usage**

```
get_all_subsets(list, name_separator = "/")
```

**Arguments**

list	a named list
name_separator	default is /

**Value**

a tibble

**Examples**

```
list = list(A = sample(LETTERS, 20),
            B = sample(LETTERS, 22),
            C = sample(LETTERS, 24),
            D = sample(LETTERS, 30, replace = TRUE))
get_all_subsets(list)
```

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**oncoplot***ploting oncoplot with a plot***Description**

ploting oncoplot with a plot

**Usage**

```
oncoplot(maf, genes = 20)
```

**Arguments**

maf	MAF object.
genes	the gene names or the number, default is 20.

**Value**

oncoplot object, which is also a aplot object

**Examples**

```
laml.maf <- system.file("extdata", "tcga_laml.maf.gz", package = "maftools")
laml.clin <- system.file('extdata', 'tcga_laml_annot.tsv', package = 'maftools')
laml <- maftools:::read.maf(maf = laml.maf, clinicalData = laml.clin)
oncoplot(maf = laml, genes = 20)
```

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**upset\_plot***upsetplot2***Description**

Plot a upset plot

**Usage**

```
upset_plot(
  list,
  nintersects = NULL,
  order.intersect.by = c("size", "name"),
  order.set.by = c("size", "name"),
  color.intersect.by = "none",
  color.set.by = "none",
  remove_empty_intersects = TRUE
)
```

**Arguments**

list	a list of sets
nintersects	number of intersects. If NULL, all intersections will show.
order.intersect.by	one of 'size' or 'name'
order.set.by	one of 'size' or 'name'
color.intersect.by	color scheme for 'intersect' bars (e.g., "Set2"), default is "none"
color.set.by	color scheme for 'set' bars (e.g., "Set3"), default is "none"
remove_empty_intersects	remove the intersects which have zero elements. Default is TRUE.

**Details**

This function generate a upset plot by creating a composite plot which contains subplots generated by ggplot2.

**Value**

an upset plot

**Examples**

```
list = list(A = sample(LETTERS, 20),
           B = sample(LETTERS, 22),
           C = sample(LETTERS, 14),
           D = sample(LETTERS, 30, replace = TRUE))
upset_plot(list)
upset_plot(list, remove_empty_intersects = TRUE)
upset_plot(list, order.intersect.by = "name")
upset_plot(list, order.set.by = "name")
upset_plot(list, nintersects = 6)
```

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