

# Package ‘aplotExtra’

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

**Version** 0.0.3

**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, 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	<i>funky_bar</i>
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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	<i>funky_heatmap</i>
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### Description

create a funkyheatmap

### Usage

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

### Arguments

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

**Value**

gglist 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
<code>...</code>	additional parameters, passing to <a href="#">geom_star</a>

**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 aplot*

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**Description**

ploting oncoplot with aplot

**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*

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**Description**

Plot a upset plot

**Usage**

```
upset_plot(  
  list,  
  nintersects = NULL,  
  order.intersect.by = c("size", "name"),  
  order.set.by = c("size", "name")  
)
```

**Arguments**

<code>list</code>	a list of sets
<code>nintersects</code>	number of intersects. If NULL, all intersections will show.
<code>order.intersect.by</code>	one of 'size' or 'name'
<code>order.set.by</code>	one of 'size' or 'name'

**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, order.intersect.by = "name")  
upset_plot(list, nintersects = 6)
```

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