Package ‘patchwork’

December 1, 2019

**Type** Package
**Title** The Composer of Plots
**Version** 1.0.0
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**Description**
The ‘ggplot2’ package provides a strong API for sequentially building up a plot, but does not concern itself with composition of multiple plots. ‘patchwork’ is a package that expands the API to allow for arbitrarily complex composition of plots by, among others, providing mathematical operators for combining multiple plots. Other packages that try to address this need (but with a different approach) are 'gridExtra' and 'cowplot'.

**License** MIT + file LICENSE
**Encoding** UTF-8
**LazyData** true
**Imports** ggplot2 (>= 3.0.0), gtable, grid, stats, grDevices, utils, graphics
**RoxygenNote** 6.1.1


**BugReports** https://github.com/thomasp85/patchwork/issues
**Suggests** knitr, rmarkdown, gridGraphics, gridExtra, ragg, testthat (>= 2.1.0), vdiffr, covr

**VignetteBuilder** knitr

**NeedsCompilation** no
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**Repository** CRAN
**Date/Publication** 2019-12-01 12:40:02 UTC
area

Specify a plotting area in a layout

Description

This is a small helper used to specify a single area in a rectangular grid that should contain a plot. Objects constructed with `area()` can be concatenated together with `c()` in order to specify multiple areas.

Usage

```r
area(t, l, b = t, r = l)
```

Arguments

- `t, b` The top and bottom bounds of the area in the grid
- `l, r` The left and right bounds of the area in the grid

Details

The grid that the areas are specified in reference to enumerate rows from top to bottom, and columns from left to right. This means that `t` and `l` should always be less or equal to `b` and `r` respectively. Instead of specifying area placement with a combination of `area()` calls, it is possible to instead pass in a single string.

```r
areas <- c(area(1, 1, 2, 1),
           area(2, 3, 3, 3))
```

is equivalent to

```r
areas <- "A##
         A#B
         ##B"
```

For an example of this, see the `plot_layout()` examples.
**guide_area**

Add an area to hold collected guides

**Description**

Using the guides argument in `plot_layout()` you can collect and collapse guides from plots. By default these guides will be put on the side like with regular plots, but by adding a `guide_area()` to the plot you can tell patchwork to place the guides in that area instead. If guides are not collected or no guides exists to collect it behaves as a standard `plot_spacer()` instead.

**Usage**

`guide_area()`

**Examples**

```r
library(ggplot2)

p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp, colour = factor(gear)))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))
p3 <- ggplot(mtcars) + geom_bar(aes(gear)) + facet_wrap(~cyl)

layout <- c(
  area(1, 1),
  area(1, 3, 3),
  area(3, 1, 3, 2)
)

# Show the layout to make sure it looks as it should
plot(layout)

# Apply it to a patchwork
p1 + p2 + p3 + plot_layout(design = layout)
```

```r
# Guides are by default kept beside their plot
p1 + p2 + p3

# They can be collected and placed on the side (according to the patchwork
```
Align plots across multiple pages

Description

Sometimes it is necessary to make sure that separate plots are aligned, with each other, but still exists as separate plots. That could e.g. be if they need to be part of a slideshow and you don’t want titles and panels jumping around as you switch between slides. patchwork provides a range of utilities to achieve that. Currently it is only possible to align ggplots, but aligning patchworks will be supported in the future.

Usage

get_dim(plot)

set_dim(plot, dim)

get_max_dim(...)

align_plots(...)

Arguments

plot A ggplot object
dim A plot_dimension object as created by get_dim()...
ggplot objects or a single list of them

Value

get_dim() and get_max_dim() return a plot_dimension object. set_dim() returns a modified ggplot object with fixed outer dimensions and align_plots() return a list of such. The modified ggplots still behaves like a standard ggplot and new layers, scales, etc can be added to them.

Examples

library(ggplot2)
p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp)) + ggtitle('Plot 1')
p2 <- ggplot(mtcars) +
```
geom_boxplot(aes(gear, disp, group = gear)) +
ggtitle('Plot 2')

p3 <- ggplot(mtcars) +
  geom_point(aes(hp, wt, colour = mpg)) +
  ggtitle('Plot 3')

p4 <- ggplot(mtcars) +
  geom_bar(aes(gear)) +
  facet_wrap(~cyl) +
  ggtile('Plot 4')

# Align a plot to p4
p4_dim <- get_dim(p4)
set_dim(p1, p4_dim)

# Align a plot to the maximum dimensions of a list of plots
max_dims <- get_max_dim(p1, p2, p3, p4)
set_dim(p2, max_dims)

# Align a list of plots with each other
aligned_plots <- align_plots(p1, p2, p3, p4)
aligned_plots[[3]]

# Aligned plots still behave like regular ggplots
aligned_plots[[3]] + theme_bw()
```

---

**plot_annotation**

Annotate the final patchwork

---

**Description**

The result of this function can be added to a patchwork using + in the same way as `plot_layout()`, but unlike `plot_layout()` it will only have an effect on the top level plot. As the name suggests it controls different aspects of the annotation of the final plot, such as titles and tags.

**Usage**

```
plot_annotation(title = NULL, subtitle = NULL, caption = NULL,
               tag_levels = NULL, tag_prefix = NULL, tag_suffix = NULL,
               tag_sep = NULL, theme = NULL)
```

**Arguments**

- `title`, `subtitle`, `caption`  
  Text strings to use for the various plot annotations.

- `tag_levels`  
  A character vector defining the enumeration format to use at each level. Possible values are 'a' for lowercase letters, 'A' for uppercase letters, '1' for numbers, 'i' for lowercase Roman numerals, and 'I' for uppercase Roman numerals.
plot_annotation

`tag_prefix`, `tag_suffix`  
Strings that should appear before or after the tag.

`tag_sep`  
A separator between different tag levels

`theme`  
A ggplot theme specification to use for the plot. Only elements related to the titles as well as plot margin and background is used.

Details

Tagging of subplots is done automatically and following the order of the plots as they are added. When the plot contains nested layouts the `tag_level` argument in the nested `plot_layout` will define whether enumeration should continue as usual or add a new level. The format of the levels are defined with `tag_levels` argument in `plot_annotation`.

Value

A `plot_annotation` object

Examples

```
library(ggplot2)

p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))
p3 <- ggplot(mtcars) + geom_bar(aes(gear)) + facet_wrap(~cyl)

# Add title, etc. to a patchwork
p1 + p2 + plot_annotation('This is a title', caption = 'made with patchwork')

# Change styling of patchwork elements
p1 + p2 +
plot_annotation(
  title = 'This is a title',
  caption = 'made with patchwork',
  theme = theme(plot.title = element_text(size = 16))
)

# Add tags to plots
p1 / (p2 | p3) +
plot_annotation(tag_levels = 'A')

# Add multilevel tagging to nested layouts
p1 / (p2 | p3 + plot_layout(tag_level = 'new')) +
plot_annotation(tag_levels = c('A', '1'))
```
Description

In addition to the + operator known in ggplot2, patchwork defines logic for some of the other operators that aids in building up your plot composition and reduce code-reuse.

Usage

```r
## S3 method for class 'ggplot'
# e1 - e2

## S3 method for class 'ggplot'
e1 / e2

## S3 method for class 'ggplot'
e1 | e2

## S3 method for class 'gg'
e1 * e2

## S3 method for class 'gg'
e1 & e2
```

Arguments

e1 A ggplot or patchwork object
e2 A ggplot or patchwork object in case of /, or a gg object such as a geom or theme specification in case of * and &

Details

patchwork augment the + operator from ggplot2 and allows the user to add full ggplot objects together in order to compose them into the same view. The last added plot is always the active one where new geoms etc. are added to. Another operator that is much like it, but not quite, is -. It also adds plots together but instead of adding the right hand side to the patchwork defined in the left hand side, it puts the left hand side besides the right hand side in a patchwork. This might sound confusing, but in essence - ensures that the right and left side are put in the same nesting level (+ puts the right side into the left side). Using - might seem unintuitive if you think of the operator as "subtract", but look at it as a hyphen instead (the underlying reason is that - is the only operator in the same precedence group as +).

Often you are interested in creating single column or single row layouts. patchwork provides | (besides) and / (over) operators to support stacking and packing of plots. See the examples for their use.
In order to reduce code repetition, `patchwork` provides two operators for adding `ggplot` elements (geoms, themes, facets, etc.) to multiple/all plots in a `patchwork`. `*` will add the element to all plots in the current nesting level, while `&` will recurse into nested patches.

**Value**

A `patchwork` object

**Examples**

```r
library(ggplot2)

p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))
p3 <- ggplot(mtcars) + geom_bar(aes(gear)) + facet_wrap(~cyl)
p4 <- ggplot(mtcars) + geom_bar(aes(carb))

# Standard addition vs division
p1 + p2 + p3 + plot_layout(ncol = 1)
p1 + p2 - p3 + plot_layout(ncol = 1)

# Stacking and packing
(p1 | p2 | p3) /
   p4

# Add elements to the same nesting level
(p1 + (p2 + p3) + p4 + plot_layout(ncol = 1)) * theme_bw()

# Recurse into nested plots as well
(p1 + (p2 + p3) + p4 + plot_layout(ncol = 1)) & theme_bw()
```

---

**plot_layout**

*Define the grid to compose plots in*

**Description**

In order to control how different plots are laid out, you need to add a layout specification. If you are nesting grids, the layout is scoped to the current nesting level.

**Usage**

```r
plot_layout(ncol = NULL, nrow = NULL, byrow = NULL, widths = NULL, heights = NULL, guides = NULL, tag_level = NULL, design = NULL)
```
**plot_layout**

Arguments

- **ncol, nrow**
  
The dimensions of the grid to create - if both are NULL it will use the same logic as `facet_wrap()` to set the dimensions.

- **byrow**
  
  Analogous to `byrow` in `matrix()`. If FALSE the plots will be filled in in column-major order.

- **widths, heights**
  
  The relative widths and heights of each column and row in the grid. Will get repeated to match the dimensions of the grid.

- **guides**
  
  A string specifying how guides should be treated in the layout. `collect` will collect guides below to the given nesting level, removing duplicates. `keep` will stop collection at this level and let guides be placed alongside their plot. auto will allow guides to be collected if a upper level tries, but place them alongside the plot if not.

- **tag_level**
  
  A string (‘keep’ or ‘new’) to indicate how auto-tagging should behave. See `plot_annotation()`.

- **design**
  
  Specification of the location of areas in the layout. Can either be specified as a text string or by concatenating calls to `area()` together. See the examples for further information on use.

Value

A `plot_layout` object to be added to a `ggassemble` object

Examples

```r
library(ggplot2)

p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))
p3 <- ggplot(mtcars) + geom_bar(aes(gear)) + facet_wrap(~cyl)
p4 <- ggplot(mtcars) + geom_bar(aes(carb))
p5 <- ggplot(mtcars) + geom_violin(aes(cyl, mpg, group = cyl))

# The plots are layed out automatically by default
p1 + p2 + p3 + p4 + p5

# Use byrow to change how the grid is filled out
p1 + p2 + p3 + p4 + p5 + plot_layout(byrow = FALSE)

# Change the grid dimensions
p1 + p2 + p3 + p4 + p5 + plot_layout(ncol = 2, widths = c(1, 2))

# Define layout at different nesting levels
p1 +
  p2 +
  (p3 +
    p4 +
    plot_layout(ncol = 1)
  ) +
```

p5 + plot_layout(widths = c(2, 1))

# Complex layouts can be created with the `design` argument
design <- c(
  area(1, 1, 2),
  area(1, 2, 1, 3),
  area(2, 3, 3),
  area(3, 1, 3, 2),
  area(2, 2)
)
p1 + p2 + p3 + p4 + p5 + plot_layout(design = design)

# The same can be specified as a character string:
design <- "122
  153
  443"
p1 + p2 + p3 + p4 + p5 + plot_layout(design = design)

# When using strings to define the design `#` can be used to denote empty
# areas
design <- "1##
  123
  ##3"
p1 + p2 + p3 + plot_layout(design = design)

---

**plot_spacer**  
*Add a completely blank area*

**Description**

This simple wrapper creates an empty transparent patch that can be added to push your other plots apart. The patch responds to adding `theme()` specifications, but only `plot.background` will have an effect.

**Usage**

`plot_spacer()`

**Value**

A `ggplot` object containing an empty plot
Examples

```r
library(ggplot2)

p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))

p1 + plot_spacer() + p2

# To have more control over spacing, you can use the `plot.margin`
# parameter for `theme()` on each individual plot.

(p1 + theme(plot.margin = unit(c(0,30,0,0), "pt"))) +
(p2 + theme(plot.margin = unit(c(0,0,0,30), "pt")))
```

Description

In order to add non-ggplot2 element to a patchwork they can be converted to a compliant representation using the `wrap_elements()` function. This allows you to position either grobs, ggplot objects, patchwork objects, or even base graphics (if passed as a formula) in either the full area, the full plotting area (anything between and including the axis label), or the panel area (only the actual area where data is drawn). Further you can still add title, subtitle, tag, and caption using the same approach as with normal ggplots (using `ggtitle()` and `labs()`) as well as styling using `theme()`. For the latter, only the theme elements targeting plot margins and background as well as title, subtitle, etc styling will have an effect. If a patchwork or ggplot object is wrapped, it will be fixated in its state and will no longer respond to addition of styling, geoms, etc.. When grobs and formulas are added directly, they will implicitly be converted to `wrap_elements(full = x)`.

Usage

```r
wrap_elements(panel = NULL, plot = NULL, full = NULL, clip = TRUE, ignore_tag = FALSE)
```

Arguments

- `panel, plot, full` A grob, ggplot, patchwork, or formula object to add to the respective area.
- `clip` Should the grobs be clipped if expanding outside its area
- `ignore_tag` Should tags be ignored for this patch. This is relevant when using automatic tagging of plots and the content of the patch does not qualify for a tag.

Value

A wrapped_patch object
Examples

```r
library(ggplot2)
library(grid)

# Combine grobs with each other
wrap_elements(panel = textGrob('Here are some text')) +
  wrap_elements(
    panel = rectGrob(gp = gpar(fill = 'steelblue'),
    full = rectGrob(gp = gpar(fill = 'goldenrod'))
  )

# wrapped elements can still get titles etc like ggplots
wrap_elements(panel = textGrob('Here are some text')) +
  wrap_elements(
    panel = rectGrob(gp = gpar(fill = 'steelblue'),
    full = rectGrob(gp = gpar(fill = 'goldenrod'))
  ) +
  ggtitle('Title for the amazing rectangles')

# You can also pass in ggplots or patchworks to e.g. have it fill out the
# panel area
p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p1 + wrap_elements(panel = p1 + ggtitle('Look at me shrink'))

# You can even add base graphics if you pass it as a formula
p1 + wrap_elements(full = ~ plot(mtcars$mpg, mtcars$disp))

# Adding a grob or formula directly is equivalent to placing it in 'full'
p1 + ~ plot(mtcars$mpg, mtcars$disp)
```

Description

This function converts a gtable, as produced by `ggplot2::ggplotGrob()` and makes it ready to be added to a patchwork. In contrast to passing the gtable to `wrap_elements()`, `wrap_ggplot_grob()` ensures proper alignment as expected. On the other hand, major restructuring of the gtable will result in an object that doesn’t work properly with `wrap_ggplot_grob()`.

Usage

`wrap_ggplot_grob(x)`

Arguments

- `x` A gtable as produced by `ggplot2::ggplotGrob()`
wrap_plots

Wrap plots into a patchwork

Description

While the use of + is a natural way to add plots together, it can be difficult to string together multiple plots programmatically if the number of plots is not known beforehand. wrap_plots makes it easy to take a list of plots and add them into one composition, along with layout specifications.

Usage

wrap_plots(..., ncol = NULL, nrow = NULL, byrow = NULL, 
  widths = NULL, heights = NULL, guides = NULL, tag_level = NULL, 
  design = NULL)

Arguments

... multiple ggplots or a list containing ggplot objects

ncol The dimensions of the grid to create - if both are NULL it will use the same logic as facet_wrap() to set the dimensions
The dimensions of the grid to create - if both are NULL it will use the same logic as `facet_wrap()` to set the dimensions

Analogous to `byrow` in `matrix()`. If FALSE the plots will be filled in in column-major order

The relative widths and heights of each column and row in the grid. Will get repeated to match the dimensions of the grid.

The relative widths and heights of each column and row in the grid. Will get repeated to match the dimensions of the grid.

A string specifying how guides should be treated in the layout. 'collect' will collect guides below to the given nesting level, removing duplicates. 'keep' will stop collection at this level and let guides be placed alongside their plot. auto will allow guides to be collected if a upper level tries, but place them alongside the plot if not.

A string ('keep' or 'new') to indicate how auto-tagging should behave. See `plot_annotation()`.

Specification of the location of areas in the layout. Can either be specified as a text string or by concatenating calls to `area()` together. See the examples for further information on use.

If `design` is specified as a text string and the plots are named (e.g. `wrap_plots(A = p1, ...)` and all plot names are single characters represented in the design layout string, the plots will be matched to their respective area by name. Otherwise the areas will be filled out sequentially in the same manner as using the `+` operator. See the examples for more.

A patchwork object

Examples

```r
library(ggplot2)

p1 <- ggplot(mtcars) + geom_point(aes(mpg, disp))
p2 <- ggplot(mtcars) + geom_boxplot(aes(gear, disp, group = gear))
p3 <- ggplot(mtcars) + geom_bar(aes(gear)) + facet_wrap(~cyl)
p4 <- ggplot(mtcars) + geom_bar(aes(carb))
p5 <- ggplot(mtcars) + geom_violin(aes(cyl, mpg, group = cyl))

# Either add the plots as single arguments
wrap_plots(p1, p2, p3, p4, p5)

# Or add them as a list...
plots <- list(p1, p2, p3, p4, p5)
wrap_plots(plots)

# Match plots to areas by name
design <- "#BB"
```
wrap_plots(B = p1, A = p2, design = design)

# Compare to not using named plot arguments
wrap_plots(p1, p2, design = design)
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