Package ‘ggedit’

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Description

add layer from gg object

Usage

```r
## S3 method for class 'gg'
e1 + e2
```

Arguments

- `e1` plot
- `e2` layer

Value

- `gg`
remove layer

Description
remove layer from gg object

Usage

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

Arguments

e1  plot
e2  layer

Value

gg

as.ggedit

Try to coerce a ggplot object into a ggedit object

Description

Applied to ggplot objects to use the plotting function of ggedit.

Usage

as.ggedit(plot)

Arguments

plot  an object

Value

an object of class ggedit

See Also

print.ggedit, ggplot
Examples

```r
p <- ggplot2::ggplot(iris, aes(x = Sepal.Length, y = Sepal.Width))

p1 <- p +
  ggplot2::geom_point(aes(colour = Species)) +
  ggplot2::geom_line()

p2 <- p +
  ggplot2::geom_point() +
  ggplot2::geom_smooth(method = 'loess')

p3 <- list(p1, p2)

p4 <- asNggedit(p3)
```

---

as.gglist recasts to gglist

### Description

adds gglist class to object

### Usage

```r
as.gglist(plot)
```

## S3 method for class 'gglist'
```r
as.gglist(plot)
```

## S3 method for class 'list'
```r
as.gglist(plot)
```

## S3 method for class 'ggplot'
```r
as.gglist(plot)
```

## S3 method for class 'gmatrix'
```r
as.gglist(plot)
```

### Arguments

- **plot**: object

### Value

gglist object
**cloneFacet**

*Clone ggplot facet object*

**Description**

Clone ggplot facet object and return either a gg object or the script to parse and evaluate.

**Usage**

```r
cloneFacet(obj, verbose = FALSE)
```

**Arguments**

- **obj**
  - gg facet wrap or facet_grid object
- **verbose**
  - boolean, toggles to return object or script (TRUE), Default: FALSE

**Value**

- gg object or script

**Examples**

```r
obj <- ggplot2::facet_grid(a+b+c+d, scales = 'free', as.table = FALSE, switch = 'x', shrink = FALSE)

cloneFacet(obj)
cloneFacet(obj, verbose=TRUE)
```

**cloneLayer**

*Creates an independent copy of a ggplot layer object*

**Description**

Creates copies of ggplot layers from within ggplot objects that are independent of the parent object.

**Usage**

```r
cloneLayer(l, verbose = FALSE, showDefaults = TRUE)
```

**Arguments**

- **l**
  - ggplot2 object layer
- **verbose**
  - toggle to control if the output is ggproto object (verbose==FALSE, default) or string of layer call (verbose==TRUE)
- **showDefaults**
  - toggle to control if the verbose output shows all the input arguments passed to the proto object (if verbose==FALSE then ignored)
Details

ggplot objects are comprised of layer objects. Once compiled they are part of the plot object environment and if they are changed internally regardless of where they are in the (ie different environment) it will change the original plot. This function allows to create replicates of the plot layers and edit them independent of the original plot. When setting verbose to TRUE function returns the ggplot2 call as a string to paste in regular ggplot script to generate the layer.

Value

ggproto or string object (conditional on verbose)

Examples

```r
p <- ggplot2::ggplot(iris, ggplot2::aes(x = Sepal.Length, y = Sepal.Width))
p <- p +
ggplot2::geom_point(ggplot2::aes(colour='Species')) +
ggplot2::geom_line()
p$layers[[1]]
newLayer <- cloneLayer(l=p$layers[[1]])
all.equal(p$layers[[1]], newLayer)
(v <- cloneLayer(l=p$layers[[1]], verbose=TRUE))
eval(parse(text=v))
all.equal(p$layers[[1]], eval(parse(text=v)))
```

cloneRoot  

clone root of ggplot object

Description

close root of ggplot object and return either a new gg object or a string to parse and evaluate

Usage

```r
cloneRoot(obj, verbose = FALSE)
```

Arguments

<table>
<thead>
<tr>
<th>obj</th>
<th>ggplot object</th>
</tr>
</thead>
<tbody>
<tr>
<td>verbose</td>
<td>boolean, toggles to return object or script (TRUE), Default: FALSE</td>
</tr>
</tbody>
</table>
cloneScales

Value

gg object or script

See Also

capture.output

Examples

cloneRoot(plist$pointSmooth)
cloneRoot(plist$pointSmooth,verbose=TRUE)

cloneScales  

Clone ggplot2 scales from compiled ggplot objects

Description

Clone ggplot2 scales from compiled ggplot objects returns not the original call but the nested call

Usage

clonescales(pL verbose = FALSE)

Arguments

p             ggplot object
verbose  boolean, if TRUE then returns script to eval(parse) if FALSE returns new compiled object Default: FALSE

Value

ggplot scale or script (depends on verbose)

Examples

# p <- plist$pointSmooth+scale_colour_continuous(low='red')
# p
# plist$pointSmooth+cloneScales(p)
**describe**

Compare differences theme object e1 (new theme) to theme object e2 (old theme)

**usage**

```r
compare(e1, e2, verbose = TRUE)
```

**arguments**

e1: theme object
e2: theme object
verbose: logical to control if the output is a character of script or a theme object (default TRUE)

**value**

theme object or character depending on verbose

**examples**

```r
compare(ggplot2::theme_bw(), ggplot2::theme_get())
```

**dput.ggedit**

Convert ggplot object to a string call

**description**

Convert ggplot object to a string call

**usage**

```r
dput.ggedit(obj, file = "")
```

**arguments**

obj: compiled ggplot object
file: either a character string naming a file or a connection. " " indicates output to the console, Default: " "

```r
dput.ggedit(ggplot2::theme_bw(), ggplot2::theme_get(), verbose=FALSE)
```
**ggedit**

**Value**
character

**Examples**

```r
pList$pointSmooth #original compiled plot
this.gg <- dput.ggedit(pList$pointSmooth) #dput the plot
writeLines(this.gg) #show the output
eval(parse(text=this.gg)) #recompile the plot

#add theme change
p <- pList$pointSmooth+theme(panel.background = element_rect(fill='green'))
this.gg<-dput.ggedit(p) #dput the plot
writeLines(this.gg) #show the output
eval(parse(text=this.gg)) #recompile the plot
```

---

**ggedit**  
*Interactive shiny gadget for editing ggplot layers and themes.*

**Description**

Shiny gadget that takes an input ggplots and populates a user interface with objects that let the user update aesthetics of layers and theme elements.

**Usage**

```r
ggedit(p.in, ...)
```

**Arguments**

- `p.in`: ggplot plot object or list of objects
- `...`: options that are passed to ggedit

**Details**

The user can start the gadget using the console `ggedit(plotobj)` or through the Addins menu in Rstudio.
If you are using the Addin option highlight on the editor window the ggplot object and then click the addin.

**Options to pass to ggedit**
viewer | shiny viewer options. It can be either paneViewer (default with minHeight=1000), dialogViewer, browserViewer

verbose | logical to control if the output includes script for layers and themes calls for parsing to create objects (default, verbose=TRUE)

showDefaults | toggle to control if the verbose output shows all the input arguments passed to the proto object (if verbose==FALSE then ignored)

width, height | dimensions of the renderPlot where the active plot is displayed

Once the gadget is running the list of plots are shown in a grid and a number of objects will appear above them.

Action buttons

Cancel:
Returns a NULL object

Done:
Returns the list decribed below.

Dropdown list

Navigates through the plots in the input list. If the input list is a named list the names will be in the dropdown. The plot chosen is termed as the “active plot”

Radio buttons

The options to choose in the radio buttons are the layer names in the active plot.

Links

Update Plot Layer:

A pop up window will appear and be populated with aesthetic elements found in the layer chosen from the radio buttons. The layer is cloned using cloneLayer creating a layer independent of the original plot. If the aesthetic is a factor the values will be shown in dropdown lists. If it is numeric it will be shown in a slider. If it is a factor colour/fill aesthetic the colourPicker package will allow to choose from the full pallete of colours. If the continuous colour/fill aesthetic a dropdown list will be shown with different palletes

Update Plot Theme:

A popup modal will appear populated with the theme elements found in the active plot. Each element will appear as having a value or empty depending if it was defined or not. The user can change or fill in any element with valid values and any textboxes left empty will use ggplot defaults.

Update Grid Theme:

Copies the theme of the active plot to the other plots in the list

Update Global Theme:

Copies the theme of the active plot to the session theme and all plots created outside of the gadget will have this theme.

View Layer Code:

Opens an ace editor to compare the active layer initial script call and the updated script call.

The ggplot objects returned (layers and themes) can be used on any ggplot object.
Value

List of elements

- **updatedPlots**  list containing updated ggplot objects
- **updatedLayers**  For each plot a list of updated layers (ggproto) objects
- **UpdatedLayersElements**  For each plot a list elements and their values in each layer
- **UpdatedLayerCalls**  For each plot a list of scripts that can be run directly from the console to create a layer
- **updatedScales**  For each plot a list of updated scale objects
- **UpdatedScalesCalls**  For each plot a list of scripts that can be run directly from the console to create a scale object
- **updatedThemes**  For each plot a list of updated theme objects
- **UpdatedThemeCalls**  For each plot a list of scripts that can be run directly from the console to create a theme

See Also

- cloneLayer
- rgg
- ggplot
- colourPicker

Examples

```r
p <- ggplot(irisLggplot::aes(x = Sepal.Length, y = Sepal.Width))

p <- p +
  ggplot2::geom_point(ggplot2::aes(colour = Species)) +
  ggplot2::geom_line()

if(interactive()){  
  ## Not run:
  pnew <- ggedit(p)
  pnew

  ## End(Not run)
}
```

---

**ggedit_opts**

*Default and current ggedit options*

**Description**

Options for functions in the ggedit package. When running R code, the object `ggedit_opts` (default options) is not modified by chunk headers (local chunk options are merged with default options), whereas `ggedit_opts_current` (current options) changes with different chunk headers and it always reflects the options for the current chunk.

Normally we set up the global options once in the first code chunk in a document using `ggedit_opts$set()`, so that all *latter* chunks will use these options. Note the global options set in one chunk will not affect the options in this chunk itself, and that is why we often need to set global options in a separate chunk.
Usage

`ggedit_opts`

`ggedit_opts_current`

Format

An object of class `list` of length 5.

Note

`ggedit_opts_current` is read-only in the sense that it does nothing if you call `ggedit_opts_current$set()`; you can only query the options via `ggedit_opts_current$get()`.

Examples

`ggedit_opts$get('themeDefaultClass')`

gggsave

Description

Wrapper of `ggsave` that saves `ggplot` or list of `ggplot` objects to image or pdf.

Usage

`gggsave(filename = "Rplot.pdf", plot = last_plot(), ...)`

Arguments

- `filename`: a character string giving the name of the file. If it is of the form "lcmd", the output is piped to the command given by cmd. If it is NULL, then no external file is created (effectively, no drawing occurs), but the device may still be queried (e.g., for size of text). For use with onefile = FALSE give a C integer format such as "Rplot%03d.pdf" (the default in that case)
- `plot`: `ggplot` or list of `ggplots` to save, defaults to last plot displayed
- `...`: other arguments passed on to graphics device

Details

default output is to create one pdf regardless of size of list of plots inputted

Value

nothing
gg_session

Examples

## Not run: gggsave(pList)

### gg_session

Retreive all functions that create ggroto layers or stats

**Usage**

`gg_session(gg_pkg = NULL)`

**Arguments**

- `gg_pkg`: character, package names to search in if NULL an internal search will look in loaded packages, Default: NULL

**Value**

data.frame

**Examples**

`gg_session('ggplot2')`

---

gg_vetting

*Backcheck what functions created the layers in a ggplot2 plot object*

**Description**

Validate geoms with their unique attributes

**Usage**

`gg_vetting(p, obj = ggedit_opts$get("session_geoms"))`

**Arguments**

- `p`: gg, compiled ggplot object
- `obj`: data.frame, contains the mapping of layer functions as created in `gg_session()`, Default: `ggedit_opts$get('session_geoms')`
Value

data.frame

Examples

gg_vetting(plist$boxplotWrap)
lapply(plist, gg_vetting)

isNggedit

Is the object of class ggedit

Description

Is the object of class ggedit. Very basic for many functions in the package.

Usage

is.ggedit(x)

Arguments

x an object

Value

logical - is the object of class ggedit

isNgglist
gglist

Description

Checks if object is a gglist object

Usage

is.gglist(x)

Arguments

x object

Value

boolean
layersList

Description
Runs the ggplot_build function on the input and converts the output data objects into a nested list with the unique values of each of the aesthetic columns.

Usage
layersList(obj)

Arguments
obj ggplot2 plot object or list of plot objects

Value
list of aesthetics and their values for each layer in a plot

Examples
p < ggplot(iris, aes(x = Sepal.Length, y = Sepal.Width))
p <- p + geom_point(aes(colour = Species)) + geom_line()
p
p.list <- layersList(p)
p.list

pList

List of plots for ggedit examples

Description
List of ggplot objects for examples in ggedit.

Usage
pList

Format
An object of class "list"

Details
list includes a geom_point, geom_point+facet_wrap, geom_boxplot+facet_wrap, geom_point+geom_line,
geom_point+geom_smooth, geom_point+geom_line+facet_wrap, geom_point+geom_line+facet_grid
## Description

Plots lists of ggplot2 plot objects layout functionality.

## Usage

```r
## S3 method for class 'ggedit'
print(x, layout = NULL, byrow = FALSE, ...)
```

## Arguments

- `x` list of ggplot2 plot objects
- `layout` matrix, layout of plots like in `layout`, if NULL then a default layout is used. Default: NULL
- `byrow` boolean, argument passed to default layout (when layout is NULL), used to transpose the output.
- `...` not used

## Examples

```r
p <- as.gglist(plist[1:2])
p

pl <- p + geom_hline(aes(yintercept=3))
pl
print(pl, byrow=TRUE)
print(pl, layout = matrix(c(2,2,NA,1), ncol=2))
```

---

---

## Description

Extract geom, stat and position protos from a ggplot2 layer.

## Usage

```r
proto_features(l)
```
remove_geom

Arguments

1. ggproto

Value
data.frame

Examples

proto_features(ggplot2::geom_smooth())
proto_features(ggplot2::annotation_logticks())

remove_geom(p, geom, idx = NULL)

Description

Removes specified layers from a ggplot object.

Usage

remove_geom(p, geom, idx = NULL)

Arguments

p : ggplot2 plot object
geom : character string of the name of the layer to remove
idx : numeric of which index of geom to remove, Default: 1

Examples

p <- ggplot2::ggplot(iris, ggplot2::aes(x = Sepal.Length, y = Sepal.Width))
p <- p + ggplot2::geom_point(ggplot2::aes(colour = Species)) + ggplot2::geom_line()
p
pnew <- p操纵 geom('point', 1)
pnew
**rgg**  
*Remove and replace ggplot2 layers.*

**Description**

Removes specified layers from a ggplot object and gives the option to replace them with a new layer. This layer can be either a geom object created from regular ggplot functions or an output from the ggedit gadget. In the latter case the layers are found in the updatedLayers object in the ggedit output.

**Usage**

```r
ggg(p, oldGeom, oldGeomIdx = 1, newLayer = NULL)
```

**Arguments**

- `p`: ggplot2 plot object
- `oldGeom`: character string of the name of the layer to remove
- `oldGeomIdx`: numeric of which index of OldGeom to remove (default is 1)
- `newLayer`: ggplot layer or list of layers

**Examples**

```r
p <- ggplot2::ggplot(iris, ggplot2::aes(x = Sepal.Length, y = Sepal.Width))
p <- p + ggplot2::geom_point(ggplot2::aes(colour = Species)) + ggplot2::geom_line()
p
p$r gg('point', 1)
if( interactive() ){
x <- ggedit(p)
pnew <- p$r gg('point', 1, x$updatedLayers[[1]])
pnew
}
```

**summary.ggedit**  
*Print to console verbose outputs of objects of class ggedit*

**Description**

function to tidy the ggedit output to single script calls for each plot
Usage

```r
## S3 method for class 'ggedit'
summary(object, ...)
```

Arguments

- `object`: gedit object
- `...`: used to pass `show.structure`

Details

use `show.structure` (boolean) to control if the output shows the full `data.frame` structure or just a placeholder `[data.frame]`, Default: FALSE

---

`%>% `  
*magrittr forward-pipe operator*

Description

Pipe an object forward into a function or call expression.

Usage

```r
lhs %>% rhs
```

Arguments

- `lhs`: A value or the magrittr placeholder.
- `rhs`: A function call using the magrittr semantics.

Details

**Using `%>%` with unary function calls**
When functions require only one argument, `x %>% f` is equivalent to `f(x)` (not exactly equivalent; see technical note below.)

**Placing `lhs` as the first argument in `rhs` call**
The default behavior of `%>%` when multiple arguments are required in the `rhs` call, is to place `lhs` as the first argument, i.e. `x %>% f(y)` is equivalent to `f(x, y)`.

**Placing `lhs` elsewhere in `rhs` call**
Often you will want `lhs` to the `rhs` call at another position than the first. For this purpose you can use the dot (.) as placeholder. For example, `y %>% f(x, .)` is equivalent to `f(x, y) and z %>% f(x, y, arg = .)` is equivalent to `f(x, y, arg = z).

**Using the dot for secondary purposes**
Often, some attribute or property of `lhs` is desired in the `rhs` call in addition to the value of `lhs`
itself, e.g. the number of rows or columns. It is perfectly valid to use the dot placeholder several times in the rhs call, but by design the behavior is slightly different when using it inside nested function calls. In particular, if the placeholder is only used in a nested function call, lhs will also be placed as the first argument! The reason for this is that in most use-cases this produces the most readable code. For example, iris %>% subset(1:nrow(.), % 2 == 0) is equivalent to iris %>% subset(., 1:nrow(.) % 2 == 0) but slightly more compact. It is possible to overrule this behavior by enclosing the rhs in braces. For example, 1:10 %>% {c(min(.), max(.))} is equivalent to c(min(1:10), max(1:10)).

**Using %>% with call- or function-producing rhs**

It is possible to force evaluation of rhs before the piping of lhs takes place. This is useful when rhs produces the relevant call or function. To evaluate rhs first, enclose it in parentheses, i.e. a %>% (function(x) x^2), and 1:10 %>% (call("sum")). Another example where this is relevant is for reference class methods which are accessed using the $ operator, where one would do x %>% (rc$f), and not x %>% rc$f.

**Using lambda expressions with %>%**

Each rhs is essentially a one-expression body of a unary function. Therefore defining lambdas in magrittr is very natural, and as the definitions of regular functions: if more than a single expression is needed one encloses the body in a pair of braces, { rhs }. However, note that within braces there are no "first-argument rule": it will be exactly like writing a unary function where the argument name is "." (the dot).

**Using the dot-placeholder as lhs**

When the dot is used as lhs, the result will be a functional sequence, i.e. a function which applies the entire chain of right-hand sides in turn to its input. See the examples.

**Technical notes**

The magrittr pipe operators use non-standard evaluation. They capture their inputs and examines them to figure out how to proceed. First a function is produced from all of the individual right-hand side expressions, and then the result is obtained by applying this function to the left-hand side. For most purposes, one can disregard the subtle aspects of magrittr's evaluation, but some functions may capture their calling environment, and thus using the operators will not be exactly equivalent to the "standard call" without pipe-operators.

Another note is that special attention is advised when using non-magrittr operators in a pipe-chain (+, -, $, etc.), as operator precedence will impact how the chain is evaluated. In general it is advised to use the aliases provided by magrittr.

**See Also**

%<>%, %|%>, %%%

**Examples**

# Basic use:
iris %>% head
# Use with lhs as first argument
iris %>% head(10)

# Using the dot place-holder
"Ceci n'est pas une pipe" %>% gsub("une", "un", .)

# When dot is nested, lhs is still placed first:
sample(1:10) %>% paste0(LETTERS[.])

# This can be avoided:
rnorm(100) %>% c(min(., mean(.), max(., .))) %>% floor

# Lambda expressions:
iris %>%
  {
    size <- sample(1:10, size = 1)
    rbind(head(., size), tail(., size))
  }

# Renaming in lambdas:
iris %>%
  {
    my_data <- .
    size <- sample(1:10, size = 1)
    rbind(head(my_data, size), tail(my_data, size))
  }

# Building unary functions with %>%
trig_fest <- . %>% tan %>% cos %>% sin

1:10 %>% trig_fest
trig_fest(1:10)
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