Package 'gginnards'

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Description Extensions to 'ggplot2' providing low-level debug tools: statistics and geometries reporting data passed to compute_group() and compute_panel() functions and to geometries. Layer manipulation: functions for deletion, insertion, extraction and reordering of layers of ``ggplot" objects. Data manipulation: function for deletion of unused variables from the data object embedded in ``ggplot" objects.

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URL https://www.r4photobiology.info, https://bitbucket.org/aphalo/gginnards

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gginnards-package gginnards: Explore the Innards of 'ggplot2' Objects

Description

Extensions to 'ggplot2' providing low-level debug tools: statistics and geometries reporting data passed to compute_group() and compute_panel() functions and to geometries. Layer manipulation: functions for deletion, insertion, extraction and reordering of layers of "ggplot" objects. Data manipulation: function for deletion of unused variables from the data object embedded in "ggplot" objects.

Details

The new facilities for cleanly defining new stats and geoms added to package 'ggplot2' in version 2.0.0 gave origin to this package. I needed tools to help me learn how layers work and to debug the extensions to 'ggplot2' that I was developing. I share them through this package in the hope that they will help other users of 'ggplot2' understand how this this vry popular graphics package works internally. The vignettes provide examples of how to use these tools both for debugging and learning how ggplots are stored.

Extensions provided:

- "Debug" stats and a "debug" geom that print to the console a summary of their data input.
- Functions for inspecting and manipulating the list of layers of a ggplot object.
- Functions for exploring and manipulating the data embedded in ggplot objects, including dropping unused variables.

Author(s)

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References

Package 'tidyverse' web site at https://www.tidyverse.org/ Package 'ggplot2' documentation at https://ggplot2.tidyverse.org/ Package 'ggplot2' source code at https://github.com/tidyverse/ggplot2

delete_layers

See Also

Useful links:

- https://www.r4photobiology.info
- https://bitbucket.org/aphalo/gginnards
- Report bugs at https://bitbucket.org/aphalo/gginnards/issues

Examples

```
# echo to the R console \code{data} as received by geoms
ggplot(mtcars, aes(cyl, mpg, color = factor(cyl))) +
 geom_point() +
 geom_debug()
# echo to the R console \code{data} as received by geoms
ggplot(mtcars, aes(cyl, mpg, colour = factor(cyl))) +
 stat_summary(fun.data = "mean_se") +
 stat_summary(fun.data = "mean_se", geom = "debug")
# echo to the R console \code{data} received by \code{compute_panel()}
ggplot(mtcars, aes(cyl, mpg, color = factor(cyl))) +
 geom_point() +
 stat_debug_panel()
# echo to the R console \code{data} received by \code{compute_group()}
ggplot(mtcars, aes(cyl, mpg, color = factor(cyl))) +
 geom_point() +
 stat_debug_group()
```

delete_layers Layer manipulation.

Description

Delete, move or append one or more layers in a ggplot object.

Usage

```
delete_layers(x, match_type = NULL, idx = NULL)
append_layers(x, object, position = "top")
move_layers(x, match_type = NULL, position = "top", idx = NULL)
shift_layers(x, match_type = NULL, idx = NULL, shift = 1L)
which_layers(x, match_type = NULL, idx = NULL)
```

```
extract_layers(x, match_type = NULL, idx = NULL)
```

top_layer(x)

bottom_layer(x)

num_layers(x)

Arguments

х	an object of class gg to be operated upon.
<pre>match_type</pre>	The name of the ggproto object class for the geom(s), position(s) or stat(s) matching that of the layers to be operated upon.
idx	integer vector Index into the list of layers used to select the layers to be operated upon.
object	a ggplot layer created by a geom_ or stat_ function or a list of such layers or an empty list.
position	character or interger, the position of the layer immediately above of which to move or append the moved or appended layers.
shift	integer.

Details

These functions must be used with care as they select all layers matching the provided geom, position or stat ggproto object class. Layers added with a stat do use a geom, and vice versa.

One and only one of match_type and idx must be passed a non-null argument.

In plots with several layers, it is possible that more than one layer matches the class name passed to match_type. It is also possible to pass a numeric vector with multiple indexes through parameter idx. In both cases multiple layers will be operated upon, but their relative positions will remain unchanged.

If a numeric vector with multiple position indexes is supplied as argument for position, the topmost position will be used. As indexing in R starts at 1, passing 0 or "bottom" as argument for position puts the moved or appended layer(s) behind all other layers (prepends the layer).

Value

An edited copy of x for delete_layers, append_layers and move_layers. An integer vector of indexes giving the positions of the matching layers in the list of layers contained in x in the case of which_layers.

Note

The functions described here are not expected to be useful in everyday plotting as one can more easily change the order in which layers are added to a ggplot. However, if one uses high level methods or functions that automatically produce a full plot using 'ggplot2' internally, one may need to add, move or delete layers so as to profit from such canned methods and retain enough flexibility.

drop_vars

References

https://stackoverflow.com/questions/13407236/remove-a-layer-from-a-ggplot2-chart

Examples

```
library(ggplot2)
df <- data.frame(</pre>
  gp = factor(rep(letters[1:3], each = 10)),
  y = rnorm(30)
)
p <- ggplot(df, aes(gp, y)) +</pre>
     geom_point() +
     stat_summary(fun.data = "mean_se", colour = "red")
n
delete_layers(p, "GeomPoint")
delete_layers(p, "StatSummary")
move_layers(p, "GeomPoint", position = "top")
move_layers(p, "GeomPointrange", position = "bottom")
move_layers(p, "StatSummary", position = "bottom")
move_layers(p, "GeomPointrange", position = 1L)
append_layers(p, geom_line(colour = "orange"), position = "bottom")
append_layers(p, geom_line(colour = "orange"), position = 1L)
extract_layers(p, "GeomPoint")
which_layers(p, "GeomPoint")
num_layers(p)
top_layer(p)
bottom_layer(p)
num_layers(ggplot())
top_layer(ggplot())
bottom_layer(ggplot())
```

drop_vars

Explore and manipulate the embedded data.

Description

Automatically remove unused variables from the "default" data object embedded in a gg or ggplot object with drop_vars(). Explore data variables and their use with mapped_vars(), data_vars() and data_attributes().

Usage

```
drop_vars(p, keep.vars = character(), guess.vars = TRUE)
mapped_vars(p, invert = FALSE)
data_vars(p)
```

```
data_attributes(p)
```

Arguments

р	ggplot Plot object with embedded data.
keep.vars	character Names of unused variables to be kept.
guess.vars	logical Flag indicating whether to find used variables automatically.
invert	logical If TRUE return indices for elements of data that are not mapped to any aesthetic or facet.

Value

character vector with names of mapped variables in the default data object.

character vector with names of all variables in the default data object.

list containing all attributes of the default data object.

Warning!

The current implementation drops variables only from the default data object. Data objects within layers are not modified.

Note

These functions are under development and not yet thoroughly tested! They are a demonstration of how one can manipulate the internals of ggplot objects creayed with 'ggplot2' versions 3.1.0 and later. These functions may stop working after some future update to the 'ggplot2' package. Although I will maintain this package for use in some of my other packages, there is no guarantee that I will be able to achieve this transparently.

Obviously, rather than using function drop_vars() after creating the ggplot object it is usually more efficient to select the variables of interest and pass a data frame containing only these to the ggplot() constructor.

```
library(ggplot2)
p <- ggplot(mpg, aes(factor(year), (cty + hwy) / 2)) +
  geom_boxplot() +
  facet_grid(. ~ class)
mapped_vars(p) # those in use
mapped_vars(p, invert = TRUE) # those not used
p.dp <- drop_vars(p) # we drop unused vars
# number of columns in the data member
ncol(p$data)
ncol(p.dp$data)</pre>
```

geom_debug

```
# which vars are in the data member
data_vars(p)
data_vars(p.dp)
# which variables in data are used in the plot
mapped_vars(p)
mapped_vars(p.dp)
# the plots identical
p
p.dp
# structure and size of p
str(p, max.level = 0)
str(p.dp, max.level = 0) # smaller in size
# structure and size of p["data"]
str(p, components = "data") # smaller in size
```

geom_debug

```
Geom which prints input data to console.
```

Description

The debug geom is used to print to the console a summary of the data being received by geoms as input data data frame.

Usage

```
geom_debug(
   mapping = NULL,
   data = NULL,
   stat = "identity",
   summary.fun = tibble::as_tibble,
   summary.fun.args = list(),
   print.fun = print,
   print.fun.args = list(),
   position = "identity",
   na.rm = FALSE,
   show.legend = FALSE,
   inherit.aes = TRUE,
   ...
)
```

Arguments

mapping	Set of aesthetic mappings created by aes or aes If specified and inherit.aes = TRUE (the default), is combined with the default mapping at the top level of the plot. You only need to supply mapping if there isn't a mapping defined for the plot.
data	A data frame. If specified, overrides the default data frame defined at the top level of the plot.
stat	The statistical transformation to use on the data for this layer, as a string.
summary.fun	A function used to print the data object received as input.
summary.fun.arg	zs
	A list of additional arguments to be passed to summary.fun.
print.fun	A function used to print the value returned by summary.fun.
print.fun.args	A list of additional arguments to be passed to print.fun.
position	Position adjustment, either as a string, or the result of a call to a position adjust- ment function.
na.rm	If FALSE (the default), removes missing values with a warning. If TRUE silently removes missing values.
show.legend	logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes.
inherit.aes	If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders.
	other arguments passed on to layer. There are three types of arguments you can use here:
	• Aesthetics: to set an aesthetic to a fixed value, like color = "red" or size = 3.
	• Other arguments to the layer, for example you override the default stat associated with the layer.
	• Other arguments passed on to the stat

• Other arguments passed on to the stat.

Details

It can be useful when debugging the code of statistics or to learn how the stats and geoms work in 'ggplot2' (>= 2.0.0).

Note

This _geom_ is very unusual in that it does not produce visible graphic output. It only returns a grid.null() grob (graphical object).

```
# echo to the R console \code{data} as received by geoms
ggplot(mtcars, aes(cyl, mpg, color = factor(cyl))) +
geom_point() +
```

geom_null

geom_debug()

```
# echo to the R console \code{data} as received by geoms
ggplot(mtcars, aes(cyl, mpg, colour = factor(cyl))) +
   stat_summary(fun.data = "mean_se") +
   stat_summary(fun.data = "mean_se", geom = "debug")
```

geom_null

A null geom or 'no-op' geom.

Description

The null geom can be used to silence graphic output from a stat, such as stat_debug_group() and stat_debug_panel() defined in this same package. No visible graphical output is returned. An invisible grid::grid_null() grob is returned instead.

Usage

```
geom_null(
  mapping = NULL,
  data = NULL,
  stat = "identity",
  position = "identity",
  na.rm = FALSE,
  show.legend = FALSE,
  inherit.aes = TRUE,
  ...
)
```

Arguments

mapping	Set of aesthetic mappings created by aes. If specified and inherit.aes = TRUE (the default), are combined with the default mapping at the top level of the plot. You only need to supply mapping if there isn't a mapping defined for the plot.
data	A data frame. If specified, overrides the default data frame defined at the top level of the plot.
stat	The statistical transformation to use on the data for this layer, as a string.
position	Position adjustment, either as a string, or the result of a call to a position adjustment function.
na.rm	If FALSE (the default), removes missing values with a warning. If TRUE silently removes missing values.
show.legend	logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes.
inherit.aes	If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders.

other arguments passed on to layer. There are three types of arguments you can use here:

- Aesthetics: to set an aesthetic to a fixed value, like color = "red" or size
 = 3.
- Other arguments to the layer, for example you override the default stat associated with the layer.
- Other arguments passed on to the stat.

Note

This _geom_ is very unusual in that it does not produce visible graphic output. It only returns a grid.null grob (graphical object). However, it accepts for consistency all the same parameters as normal geoms, which have no effect on the graphical output, except for show.legend.

Examples

```
ggplot(mtcars, aes(cyl, mpg)) +
geom_null()
```

stat_debug_group *Print to console data received by the compute group function.*

Description

stat_debug reports all distinct values in group and PANEL, and nrow, ncol and the names of the columns or variables, and the class of x and y for each group in a ggplot as passed to the compute_group function in the ggproto object.

Usage

```
stat_debug_group(
  mapping = NULL,
  data = NULL,
  geom = "null",
  summary.fun = tibble::as_tibble,
  summary.fun.args = list(),
  position = "identity",
  na.rm = FALSE,
  show.legend = FALSE,
  inherit.aes = TRUE,
  ...
)
```

. . .

Arguments

mapping	The aesthetic mapping, usually constructed with aes or aes Only needs to be set at the layer level if you are overriding the plot defaults.
data	A layer specific dataset - only needed if you want to override the plot defaults.
geom	The geometric object to use display the data
summary.fun	A function used to print the data object received as input.
summary.fun.arg	gs
	A list.
position	The position adjustment to use for overlapping points on this layer
na.rm	a logical value indicating whether NA values should be stripped before the com- putation proceeds.
show.legend	logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes.
inherit.aes	If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders.
	other arguments passed on to layer. This can include aesthetics whose values you want to set, not map. See layer for more details.

Computed variables

x x at centre of range
y y at centre of range
nrow nrow() of data object
ncol ncol() of data object
colnames colnames() of data object
colclasses class() of x and y columns in data object
group all distinct values in group as passed in data object
PANEL all distinct values in PANEL as passed in data object

See Also

Other diagnosis functions: stat_debug_panel()

```
ggplot(my.df, aes(x,y, colour = group)) +
geom_point() +
stat_debug_group()
ggplot(my.df, aes(x,y)) +
geom_point() +
facet_wrap(~group) +
stat_debug_group()
```

stat_debug_panel Print to console data received by the compute panel function.

Description

stat_debug reports all distinct values in group and PANEL, and nrow, ncol and the names of the columns or variables, and the class of x and y for each panel in a ggplot as passed to the compute_panel function in the ggproto object.

Usage

```
stat_debug_panel(
  mapping = NULL,
  data = NULL,
  geom = "null",
  summary.fun = tibble::as_tibble,
  summary.fun.args = list(),
  position = "identity",
  na.rm = FALSE,
  show.legend = FALSE,
  inherit.aes = TRUE,
  ...
)
```

Arguments

mapping	The aesthetic mapping, usually constructed with aes or aes Only needs to be set at the layer level if you are overriding the plot defaults.	
data	A layer specific dataset - only needed if you want to override the plot defaults.	
geom	The geometric object to use display the data	
summary.fun	A function used to print the data object received as input.	
summary.fun.args		
	A list.	
position	The position adjustment to use for overlapping points on this layer	
na.rm	a logical value indicating whether NA values should be stripped before the com- putation proceeds.	

show.legend	logical. Should this layer be included in the legends? NA, the default, includes if any aesthetics are mapped. FALSE never includes, and TRUE always includes.
inherit.aes	If FALSE, overrides the default aesthetics, rather than combining with them. This is most useful for helper functions that define both data and aesthetics and shouldn't inherit behaviour from the default plot specification, e.g. borders.
	other arguments passed on to layer. This can include aesthetics whose values you want to set, not map. See layer for more details.

Computed variables

x x at centre of range
y y at centre of range
nrow nrow() of data object
ncol ncol() of data object
colnames colnames() of data object
colclasses class() of x and y columns in data object
group all distinct values in group as passed in data object
PANEL all distinct values in PANEL as passed in data object

See Also

Other diagnosis functions: stat_debug_group()

Description

A str() method tailored to objects of class "ggplot". It adds to the output the size of the object, and the ability to subset individual components.

Usage

```
## S3 method for class 'ggplot'
str(
    object,
    ...,
    max.level = 1,
    components = TRUE,
    vec.len = 2,
    list.len = 99,
    give.attr = FALSE,
    comp.str = "$ ",
    nest.lev = 0,
    indent.str = paste(rep.int(" ", max(0, nest.lev + 1)), collapse = ".."),
    size = TRUE
)
```

Arguments

object	ggplot Plot object with embedded data.
	accept additional parameter arguments
<pre>max.level</pre>	integer Maximum depth of recursion (of lists within lists) to be printed.
components	Vector of components to print, as indexes into object.
vec.len	integer Approximate maximum length allowed when showing the first few values of a vector.
list.len	integer Maximum number of components to show of any list that will be described.
give.attr	logical Flag, determining whether a description of attributes will be shown.
comp.str	character String to be used for separating list components.
nest.lev	numeric current nesting level in the recursive calls to str().
indent.str	character String used for each level of indentation.
size	logical Flag, should the size of the object in bytes be printed?

str

Value

A NULL is returned invisibly. While a description of the structure of p or its components will be printed in outline form as a "side-effect", with indentation for each level of recursion, showing the internal storage mode, class(es) if any, attributes, and first few elements of each data vector. By default each level of list recursion is indicated and attributes enclosed in angle brackets.

Note

In the case of objects with a nested structure str() is called recursively and dispatched according to the class of each nested member.

See Also

A summary method for class ggplot is defined by package 'ggplot2'. Method summary() provides a more compact description of "ggplot" objects than method str(). Here we provide a wrapper on R's str() with different default arguments. A summary does not directly describe how the different components of an R object are stored, while the structure does.

```
p <- ggplot(mpg, aes(factor(year), (cty + hwy) / 2)) +</pre>
 geom_boxplot() +
 geom_point(color = "red") +
 facet_grid(. ~ class) +
 ggtitle("Example plot")
р
# str(p) vs. summary(p)
str(p)
summary(p) # from pacakge 'ggplot2'
# structure of p at 2 levels of nesting
str(p, max.level = 2, size = FALSE)
# top level structure and size of p
str(p, max.level = 0)
# names of ggplot members
names(p)
# structure and size of p["data"]
str(p, max.level = 2, components = "data")
# structure and size of p["layers"]
str(p, max.level = 1, components = "layers")
```

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