Package ‘ggformula’

April 11, 2023

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Description  Provides a formula interface to ‘ggplot2’ graphics.
Type  Package
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License  MIT + file LICENSE
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LazyLoad  TRUE
Depends  R (>= 3.1), ggplot2 (>= 3.3), ggstance (>= 0.3.4), scales, ggridges
Imports  mosaicCore (>= 0.7.0), rlang, magrittr, tibble, stringr, ggridges, grid, labelled
Suggests  tidyr, mosaicData, dplyr, lattice, mosaic, palmerpenguins, testthat, vdiffr, knitr, rmarkdown, lubridate, survival, broom, maps, maptools, rgeos, sf, purrr, ggthemes, covr, ggplot2movies, interp, quantreg
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   'layer_factory.R' 'gf_function2d.R' 'gf_functions.R'
   'gf_plot.R' 'ggstance.R' 'ggridges.R' 'ggstrings.R'
   'newplots.R' 'reexports.R' 'scales.R' 'utils.R' 'relabel.R'
   'vdiffr.R' 'zzz.R'
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discrete_breaks

Description

Creates a function that can be passed to scales for creating discrete breaks at multiples of resolution.

Usage

discrete_breaks(resolution = 1)

Arguments

resolution Resolution of the breaks

Value

A function that can be passed to scales functions as the breaks argument.

Examples

```r
x <- rbinom(100, 100, 0.4)
p <- gf_bar(~ x)
p %>% gf_refine(scale_x_continuous(breaks = discrete_breaks()))
p %>% gf_refine(scale_x_continuous(breaks = discrete_breaks(5)))
p %>% gf_refine(scale_x_continuous(breaks = discrete_breaks(2)))
```
get_variable_labels

Set and extract labels from a labeled object

Description

Some packages like expss provide mechanisms for providing longer labels to R objects. These labels can be used when labeling plots and tables, for example, without requiring long or awkward variable names. This is an experimental feature and currently only supports expss or any other system that stores a label in the label attribute of a vector.

Usage

get_variable_labels(...)

Arguments

... passed to labelled::var_label()

Details

get_variable_labels() is a synonym of labelled::var_label().

See Also

labelled::var_label(), labelled::set_variable_labels()

Examples

KF <-
mosaicData::KidsFeet %>%
  set_variable_labels(
    length = 'foot length (cm)',
    width  = 'foot width (cm)',
    birthmonth = 'birth month',
    birthyear = 'birth year',
    biggerfoot = 'bigger foot',
    domhand  = 'dominant hand'
  )
KF %>%
gf_point(length ~ width, color = ~ domhand)
get_variable_labels(KF)
Reference lines – horizontal, vertical, and diagonal.

Description
These functions create layers that display lines described in various ways. Unlike most of the plotting functions in ggformula, these functions do not take a formula as input for describing positional attributes of the plot.

Usage
gf_abline(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...
  slope,
  intercept,
  color,
  linetype,
  linewidth,
  alpha,
  xlab,
  ylab,
  title,
  subtitle,
  show.legend = NA,
  show.help = NULL,
  inherit = FALSE,
  environment = parent.frame()
)

gf_hline(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...
  yintercept,
  color,
  linetype,
  linewidth,
  alpha,
  xlab,
  ylab,
  title,
  subtitle,
caption,
show.legend = NA,
show.help = NULL,
inherit = FALSE,
environment = parent.frame()
)

gf_vline(
  object = NULL,
gformula = NULL,
data = NULL,
  ...,xintercept,
color,
linetype,
linewidth,
alpha,
xlab,
ylab,
title,
subtitle,
caption,
show.legend = NA,
show.help = NULL,
inherit = FALSE,
environment = parent.frame()
)

gf_coefline(object = NULL, coef = NULL, model = NULL, ...)

Arguments

object  When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula  Must be NULL.
data  The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
...

Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
\textbf{gf_abline}

\begin{itemize}
  \item \textbf{color} A color or a formula used for mapping color.
  \item \textbf{linetype} A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
  \item \textbf{linewidth} A numerical line width or a formula used for mapping linewidth.
  \item \textbf{alpha} Opacity (0 = invisible, 1 = opaque).
  \item \textbf{xlab} Label for x-axis. See also \texttt{gf_labs()}.
  \item \textbf{ylab} Label for y-axis. See also \texttt{gf_labs()}.
  \item \textbf{title}, \textbf{subtitle}, \textbf{caption}
    Title, sub-title, and caption for the plot. See also \texttt{gf_labs()}.
  \item \textbf{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. It can also be a named logical vector to finely select the aesthetics to display.
  \item \textbf{show.help} If TRUE, display some minimal help.
  \item \textbf{inherit} A logical indicating whether default attributes are inherited.
  \item \textbf{environment} An environment in which to look for variables not found in \texttt{data}.
  \item \textbf{xintercept}, \textbf{yintercept}, \textbf{slope}, \textbf{intercept}
    Parameters that control the position of the line. If these are set, \texttt{data}, \texttt{mapping} and \texttt{show.legend} are overridden.
  \item \textbf{coef} A numeric vector of coefficients.
  \item \textbf{model} A model from which to extract coefficients.
\end{itemize}

\section*{See Also}
\texttt{ggplot2::geom_abline()}, \texttt{ggplot2::geom_vline()}, \texttt{ggplot2::geom_hline()}

\section*{Examples}
\begin{verbatim}
mtcars2 <- df_stats(wt ~ cyl, data = mtcars, median_wt = median)
gf_point(wt ~ hp, size = ~wt, color = ~cyl, data = mtcars) %>%
gf_abline(slope = ~0, intercept = ~median_wt, color = ~cyl, data = mtcars2)

# avoid warnings by using formulas:
gf_point(wt ~ hp, size = ~wt, color = ~cyl, data = mtcars) %>%
gf_abline(slope = ~0, intercept = ~3, color = "green")

gf_point(wt ~ hp, size = ~wt, color = ~cyl, data = mtcars) %>%
gf_hline(yintercept = ~median_wt, color = ~cyl, data = mtcars2)

gf_point(mpg ~ hp, color = ~cyl, size = ~wt, data = mtcars) %>%
gf_abline(color = "red", slope = ~ - 0.10, intercept = ~ 35)

gf_point(mpg ~ hp, color = ~cyl, size = ~wt, data = mtcars) %>%
gf_abline(
\end{verbatim}
color = "red", slope = ~slope, intercept = ~intercept,
  data = data.frame(slope = -0.10, intercept = 33:35)
)

# We can set the color of the guidelines while mapping color in other layers
gf_point(mpg ~ hp, color = ~cyl, size = ~ wt, data = mtcars) %>%
gf_hline(color = "navy", yintercept = ~ c(20, 25), data = NA) %>%
gf_vline(color = "brown", xintercept = ~ c(200, 300), data = NA)

# If we want to map the color of the guidelines, it must work with the
# scale of the other colors in the plot.
gf_point(mpg ~ hp, size = ~ wt, data = mtcars, alpha = 0.3) %>%
gf_hline(color = ~ "horizontal", yintercept = ~ c(20, 25), data = NA) %>%
gf_vline(color = ~ "vertical", xintercept = ~ c(100, 200, 300), data = NA)

gf_point(mpg ~ hp, size = ~ wt, color = ~ factor(cyl), data = mtcars, alpha = 0.3) %>%
gf_hline(color = ~ "orange", yintercept = ~ 20) %>%
gf_vline(color = ~ c("4", "6", "8"), xintercept = ~ c(80, 120, 250), data = NA)

gf_point(mpg ~ hp, size = ~ wt, color = ~ factor(cyl), data = mtcars, alpha = 0.3) %>%
gf_hline(color = ~ "orange", yintercept = ~ 20) %>%
gf_vline(color = c("green", "red", "blue"), xintercept = ~ c(80, 120, 250),
  data = NA)

# reversing the layers requires using inherit = FALSE

gf_hline(color = ~ "orange", yintercept = ~ 20) %>%
gf_vline(color = ~ c("4", "6", "8"), xintercept = ~ c(80, 120, 250), data = NA) %>%
gf_point(mpg ~ hp,
  size = ~ wt, color = ~ factor(cyl), data = mtcars, alpha = 0.3,
  inherit = FALSE)

---

gf_area  

Formula interface to geom_area()

Description

For each x value, geom_ribbon() displays a y interval defined by ymin and ymax. geom_area() is a special case of geom_ribbon(), where the ymin is fixed to 0 and y is used instead of ymax.

Usage

gf_area(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
gf_area

fill,
group,
linetype,
linewidth,
xlab,
ylab,
title,
subtitle,
caption,
geom = "area",
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
    If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
    A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
    A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
...
    Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
xlab Label for x-axis. See also gf_labs().
gf_ash

**Description**

An ASH plot is the average over all histograms of a fixed bin width. `geom_ash()` and `gf_ash()` provide ways to create ASH plots using `ggplot2` or `ggformula`.

**Examples**

```r
if (require(dplyr) && require(mosaicData)) {
  Temps <- Weather %>%
    filter(city == "Chicago", year == 2016, month <= 4)
  gf_linerange(low_temp + high_temp ~ date, color = ~high_temp, data = Temps)
  gf_ribbon(low_temp + high_temp ~ date, data = Temps, color = "navy", alpha = 0.3)
  gf_area(high_temp ~ date, data = Temps, color = "navy", alpha = 0.3)

  gf_ribbon(low_temp + high_temp ~ date, data = Weather, alpha = 0.3) %>%
    gf_facet_grid(city ~ .)

  gf_linerange(low_temp + high_temp ~ date, color = ~high_temp, data = Weather) %>%
    gf_facet_grid(city ~ .) %>%
    gf_refine(scale_colour_gradientn(colors = rev(rainbow(5))))
}
```
Usage

gf_ash(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., alpha, color, group, linetype, linewidth, xlab, ylab, title, subtitle, caption, geom = "line", stat = "ash", position = "identity", show.legend = NA, show.help = NULL, inherit = TRUE, environment = parent.frame()
)

stat_ash(
  mapping = NULL, data = NULL, geom = "line", position = "identity", na.rm = FALSE, show.legend = NA, inherit.aes = TRUE, binwidth = NULL, adjust = 1, ...
)

geom_ash(
  mapping = NULL, data = NULL, stat = "ash", position = "identity", na.rm = FALSE, show.legend = NA, inherit.aes = TRUE, binwidth = NULL, adjust = 1,
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape \(~x\) or \(y \sim x\). \(y\) may be \text{stat(density)} or \text{stat(count)} or \text{stat(ndensity)} or \text{stat(ncount)}. Faceting can be achieved by including \(|\) in the formula.

data
A data frame with the variables to be plotted.

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with \text{attribute} = \text{value}, (b) ggplot2 aesthetics to be mapped with \text{attribute} = \~ \text{expression}, or (c) attributes of the layer as a whole, which are set with \text{attribute} = \text{value}.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

xlab
Label for x-axis. See also \text{gf_labs()}.

ylab
Label for y-axis. See also \text{gf_labs()}.

title, subtitle, caption
Title, sub-title, and caption for the plot. See also \text{gf_labs()}.

geom
A character string naming the geom used to make the layer.

stat
A character string naming the stat used to make the layer.

position
Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend
A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.

mapping
set of aesthetic mappings created by \text{aes()} or \text{aes()}.

na.rm
If FALSE (the default), removes missing values with a warning. If TRUE silently removes missing values.

inherit.aes
A logical indicating whether default aesthetics are inherited.

binwidth
the width of the histogram bins. If \text{NULL} (the default) the binwidth will be chosen so that approximately 10 bins cover the data. \text{adjust} can be used to to increase or decrease \text{binwidth}.

adjust
a numeric adjustment to \text{binwidth}. Primarily useful when \text{binwidth} is not specified. Increasing \text{adjust} makes the plot smoother.
Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

geom_histogram(), link(gf_histogram()).

Examples

data(penguins, package = "palmerpenguins")
gf_ash(~bill_length_mm, color = ~species, data = penguins)
gf_ash(~bill_length_mm, color = ~species, data = penguins, adjust = 2)
gf_ash(~bill_length_mm, color = ~species, data = penguins, binwidth = 1)
gf_ash(~bill_length_mm, color = ~species, data = penguins, binwidth = 1, adjust = 2)
ggplot(faithful, aes(x = eruptions)) +
  geom_histogram(aes(y = stat(density)),
      fill = "lightskyblue", colour = "gray50", alpha = 0.2
    ) +
  geom_ash(colour = "red") +
  geom_ash(colour = "forestgreen", adjust = 2) +
  geom_ash(colour = "navy", adjust = 1 / 2) +
  theme_minimal()

---

gf_bar

Formula interface to geom_bar()

Description

There are two types of bar charts: geom_bar() and geom_col(). geom_bar() makes the height of the bar proportional to the number of cases in each group (or if the weight aesthetic is supplied, the sum of the weights). If you want the heights of the bars to represent values in the data, use geom_col() instead. geom_bar() uses stat_count() by default: it counts the number of cases at each x position. geom_col() uses stat_identity(): it leaves the data as is.
Usual

gf_bar(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  width = NULL,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "bar",
  stat = "count",
  position = "stack",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_counts(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  width = NULL,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "bar",
  stat = "count",
  position = "stack",
  show.legend = NA,
`gf_bar`

```
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_props(
  object = NULL,
gformula = NULL,
data = NULL,
...,
alpha,
color,
fill,
group,
linetype,
linewidth,
xlab,
ylab = "proportion",
title,
subtitle,
caption,
geom = "bar",
stat = "count",
position = "stack",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame(),
denom = ~PANEL
)

gf_percents(
  object = NULL,
gformula = NULL,
data = NULL,
..., 
alpha,
color,
fill,
group,
linetype,
linewidth,
xlab,
ylab = "percent",
title,
subtitle,
caption,
geom = "bar",
```
\texttt{gf\_countsh(}
\begin{verbatim}
    object = \text{NULL},
    gformula = \text{NULL},
    data = \text{NULL},
    ..., 
    alpha,
    color,
    fill,
    group,
    linetype,
    linewidth,
    width = \text{NULL},
    xlab,
    ylab,
    title,
    subtitle,
    caption,
    geom = "barh",
    stat = "count",
    position = "stackv",
    show.legend = \text{NA},
    show.help = \text{NULL},
    inherit = \text{TRUE},
    environment = \text{parent.frame()},
    denom = \sim \text{PANEL}
\end{verbatim}
\texttt{)}

\texttt{gf\_colh(}
\begin{verbatim}
    object = \text{NULL},
    gformula = \text{NULL},
    data = \text{NULL},
    ..., 
    alpha,
    color,
    fill,
    group,
    linetype,
    linewidth,
    width = \text{NULL},
    xlab,
\end{verbatim}
\texttt{)}
gf_bar

ylab,
title,
subtitle,
caption,
geom = "colh",
stat = "identity",
position = "stackv",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_propsh(
  object = NULL,
gformula = NULL,
data = NULL,
...,
alpha,
color,
fill,
group,
linetype,
linewidth,
xlab = "proportion",
ylab,
title,
subtitle,
caption,
geom = "barh",
stat = "counth",
position = "stackv",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame(),
denom = ~PANEL
)

gf_percentsh(
  object = NULL,
gformula = NULL,
data = NULL,
...,
alpha,
color,
fill,
group,
linetype, linewidth, xlab = "percent", ylab, title, subtitle, caption, geom = "barh", stat = "counth", position = "stackv", show.legend = NA, show.help = NULL, inherit = TRUE, environment = parent.frame(), denom = ~PANEL)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula, typically with shape ~ x. (y ~ x is also possible, but typically using one of gf_col(), gf_props(), or gf_percents() is preferable to using this formula shape.) Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
width Width of the bars.
gf_bar

xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom, stat Override the default connection between geom_bar() and stat_count().
position Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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. It can also be a named logical vector to finely select the aesthetics to display.
show.help If TRUE, display some minimal help.
inherit A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.
denom A formula, the right hand side of which describes the denominators used for computing proportions and percents. These are computed after the stat has been applied to the data and should refer to variables available at that point. See the examples.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_bar()
Examples

```
gf_bar(~substance, data = mosaicData::HELPrct)
gf_bar(~substance, data = mosaicData::HELPrct, fill = ~sex)  
gf_bar(~substance,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge()  
)  
# gf_counts() is another name for gf_bar()  
gf_counts(~substance,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge()  
)  
# gf_props() and gf_percents() use proportions or percentages instead of counts  
# use denom to control which denominators are used.  
gf_props(~substance,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge()  
)  
gf_props(substance ~ .,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge(),  
orientation = 'y'  
)  
gf_propsh(substance ~ .,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge(),  
)  
gf_percents(~substance,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge()  
)  
gf_percents(~substance,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge(),  
denom = ~x  
)  
gf_percents(~substance,  
data = mosaicData::HELPrct, fill = ~sex,  
position = position_dodge(),  
denom = ~fill  
)  
gf_percents(~substance | sex,  
data = mosaicData::HELPrct, fill = ~homeless,  
position = position_dodge()  
)  
gf_percents(~substance | sex,  
data = mosaicData::HELPrct, fill = ~homeless,  
denom = ~fill,  
position = position_dodge()  
)  
```
gf_barh

gf_percents(~substance | sex,
  data = mosaicData::HELPrct,
  fill = ~homeless,
  denom = ~interaction(fill, PANEL),
  position = position_dodge()
)
if (require(scales)) {
  gf_percents(~substance,
    data = mosaicData::HELPrct, fill = ~sex,
    position = position_dodge(),
    denom = ~ x,
  )
  gf_refine(scale_y_continuous(labels = scales::percent))
}

---

gf_barh

Formula interface to geom_barh()

Description

Horizontal version of geom_bar().

Usage

gf_barh(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  width = NULL,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "barh",
  stat = "counth",
  position = "stackv",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)`
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula, typically with shape \(~ x\). \((y \sim x\) is also possible, but typically using one of \(gf\_col()\), \(gf\_props()\), or \(gf\_percents()\) is preferable to using this formula shape.) Faceting can be achieved by including \( | \) in the formula.

data
The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to \(ggplot()\).
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See \(fortify()\) for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. \(~ \text{head}(.x, 10)\)).

\[\ldots\]
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with \(\text{attribute} = \text{value}\), (b) ggplot2 aesthetics to be mapped with \(\text{attribute} = \text{expression}\), or (c) attributes of the layer as a whole, which are set with \(\text{attribute} = \text{value}\).

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

fill
A color for filling, or a formula used for mapping fill.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

width
Width of the bars.

xlab
Label for x-axis. See also \(gf\_labs()\).

ylab
Label for y-axis. See also \(gf\_labs()\).
title, subtitle, caption
Title, sub-title, and caption for the plot. See also \(gf\_labs()\).

geom
A character string naming the geom used to make the layer.

stat
Override the default connection between \(geom\_bar()\) and \(stat\_count()\).

position
Position adjustment, either as a string, or the result of a call to a position adjustment function.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.
```
Value

a gg object

Horizontal Geoms

There are two ways to obtain "horizontal" geoms: (1) The ggstance package provides a set of "horizontal" geoms and positions; (2) Thee ggplot2 now provides an orientation argument for "native" horizontal geoms and positions. ggformula supports both.

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A ~ B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggstance::geom_barh()`

Examples

```r
gf_barh(~Diet, data = ChickWeight)
gf_bar(Diet ~ ., data = ChickWeight, orientation = 'y')
gf_barh(~substance, data = mosaicData::HELPrct, fill = ~sex)
gf_bar(substance ~ ., data = mosaicData::HELPrct, fill = ~sex, orientation = 'y')
gf_barh(~substance, data = mosaicData::HELPrct, fill = ~sex, position = position_dodgev())
# gf_countsh() is another name for gf_barh()
gf_countsh(~Diet, data = ChickWeight)

# gf_propsh() and gf_percentsh() use proportions or percentages instead of counts
gf_propsh(substance ~ ., data = mosaicData::HELPrct, fill = ~sex, position = position_dodgev())
gf_props(substance ~ ., data = mosaicData::HELPrct, fill = ~sex, position = position_dodgev(), orientation = 'y')
gf_props(~substance, data = mosaicData::HELPrct, fill = ~sex, position = position_dodgev())
gf_percents(~substance, data = mosaicData::HELPrct, fill = ~sex,
```

position = position_dodge()

if (require(scales)) {
  gf_props(~substance, data = mosaicData::HELPrct, fill = ~sex,
           position = position_dodge()) %>%
    gf_refine(scale_y_continuous(labels = scales::percent))
}

---

**gf_bin2d**

*Formula interface to geom_bin2d()*

---

**Description**

`geom_bin2d()` uses `ggplot2::stat_bin2d()` to bin the data before using `gf_tile()` to display the results.

**Usage**

```r
gf_bin2d(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "tile",
  stat = "bin2d",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)
```

**Arguments**

- **object**
  - When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
A formula with shape $y \sim x$. Faceting can be achieved by including $|$ in the formula.

data
A data frame with the variables to be plotted.

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha Opacity ($0 = $invisible, $1 = opaque$).

color A color or a formula used for mapping color.

fill A color for filling, or a formula used for mapping fill.

group Used for grouping.

linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth A numerical line width or a formula used for mapping linewidth.

xlab Label for x-axis. See also gf_labs().

ylab Label for y-axis. See also gf_labs().

title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().

geom A character string naming the geom used to make the layer.

stat A character string naming the stat used to make the layer.

position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help If TRUE, display some minimal help.

inherit A logical indicating whether default attributes are inherited.

environment An environment in which to look for variables not found in data.

Value a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to g_facet_wrap() and g_facet_grid() that is terser and may feel more familiar to users of lattice.
Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_bin2d(), gf_tile()

Examples

gf_bin2d(eruptions ~ waiting, data = faithful, bins = 15) %>%
gf_refine(scale_fill_viridis_c(begin = 0.1, end = 0.9))

---

gf_blank

Formula interface to geom_blank()

Description

The blank geom draws nothing, but can be a useful way of ensuring common scales between different plots. See expand_limits() for more details.

Usage

gf_blank(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "blank",
)
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape y ~ x. Faceting can be achieved by including | in the formula.

data
A data frame with the variables to be plotted.

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

xlab
Label for x-axis. See also gf_labs().

ylab
Label for y-axis. See also gf_labs().
title, subtitle, caption
Title, sub-title, and caption for the plot. See also gf_labs().

geom
A character string naming the geom used to make the layer.

stat
A character string naming the stat used to make the layer.

position
Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend: A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help: If TRUE, display some minimal help.

inherit: A logical indicating whether default attributes are inherited.

environment: An environment in which to look for variables not found in data.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_blank()

Examples

gf_point((c(0, 1)) ~ (c(0, 5)))
gf_frame((c(0, 1)) ~ (c(0, 5)))
gf_blank((c(0, 1)) ~ (c(0, 5)))
# gf_blank() can be used to expand the view
gf_point((c(0, 1)) ~ (c(0, 5))) %>%
gf_blank((c(0, 3)) ~ (c(-2, 7)))
gf_boxplot

Formula interface to geom_boxplot()

Description

The boxplot compactly displays the distribution of a continuous variable. It visualises five summary statistics (the median, two hinges and two whiskers), and all "outlying" points individually.

Usage

gf_boxplot(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha, 
  color, 
  fill, 
  group, 
  linetype, 
  linewidth, 
  coef, 
  outlier.color = NULL, 
  outlier.fill = NULL, 
  outlier.shape = 19, 
  outlier.size = 1.5, 
  outlier.stroke = 0.5, 
  outlier.alpha = NULL, 
  notch = FALSE, 
  notchwidth = 0.5, 
  varwidth = FALSE, 
  xlab, 
  ylab, 
  title, 
  subtitle, 
  caption, 
  geom = "boxplot", 
  stat = "boxplot", 
  position = "dodge", 
  show.legend = NA, 
  show.help = NULL, 
  inherit = TRUE, 
  environment = parent.frame()
)
Arguments

- **object**: When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

- **gformula**: A formula with shape y ~ x. Faceting can be achieved by including | in the formula.

- **data**: The data to be displayed in this layer. There are three options:
  - If NULL, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.
  - A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.
  - A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. `~ head(.x, 10)`).

- ... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

- **alpha**: Opacity (0 = invisible, 1 = opaque).

- **color**: A color or a formula used for mapping color.

- **fill**: A color for filling, or a formula used for mapping fill.

- **group**: Used for grouping.

- **linetype**: A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

- **linewidth**: A numerical line width or a formula used for mapping linewidth.

- **coef**: Length of the whiskers as multiple of IQR. Defaults to 1.5.

- **outlier.color, outlier.fill, outlier.shape, outlier.size, outlier.stroke, outlier.alpha**: Default aesthetics for outliers. Set to NULL to inherit from the aesthetics used for the box. In the unlikely event you specify both US and UK spellings of colour, the US spelling will take precedence. Sometimes it can be useful to hide the outliers, for example when overlaying the raw data points on top of the boxplot. Hiding the outliers can be achieved by setting outlier.shape = NA. Importantly, this does not remove the outliers, it only hides them, so the range calculated for the y-axis will be the same with outliers shown and outliers hidden.

- **notch**: If FALSE (default) make a standard box plot. If TRUE, make a notched box plot. Notches are used to compare groups; if the notches of two boxes do not overlap, this suggests that the medians are significantly different.

- **notchwidth**: For a notched box plot, width of the notch relative to the body (defaults to `notchwidth = 0.5`).

- **varwidth**: If FALSE (default) make a standard box plot. If TRUE, boxes are drawn with widths proportional to the square-roots of the number of observations in the groups (possibly weighted, using the weight aesthetic).
**gf_boxplot**

- **xlab**
  Label for x-axis. See also `gf_labs()`.

- **ylab**
  Label for y-axis. See also `gf_labs()`.

- **title, subtitle, caption**
  Title, sub-title, and caption for the plot. See also `gf_labs()`.

- **geom, stat**
  Use to override the default connection between `geom_boxplot()` and `stat_boxplot()`.

- **position**
  Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

- **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. It can also be a named logical vector to finely select the aesthetics to display.

- **show.help**
  If TRUE, display some minimal help.

- **inherit**
  A logical indicating whether default attributes are inherited.

- **environment**
  An environment in which to look for variables not found in data.

**Value**

A gg object

**Specifying plot attributes**

Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

**Evaluation**

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

**References**


**See Also**

`ggplot2::geom_boxplot()`, `fivenum()`, `df_stats()`
Examples

```r
gf_boxplot(age ~ substance, data = mosaicData::HELPrct)
gf_boxplot(age ~ substance, data = mosaicData::HELPrct, varwidth = TRUE)
gf_boxplot(age ~ substance, data = mosaicData::HELPrct, color = ~sex)
gf_boxplot(age ~ substance, 
  data = mosaicData::HELPrct, 
  color = ~sex, outlier.color = "gray50"
)
# longer whiskers
gf_boxplot(age ~ substance, 
  data = mosaicData::HELPrct, 
  color = ~sex, coef = 2
)

# Note: width for boxplots is full width of box.
# For jittering, it is the half-width.
gf_boxplot(age ~ substance | sex, 
  data = mosaicData::HELPrct, 
  coef = 5, width = 0.4
) %>%
  gf_jitter(width = 0.2, alpha = 0.3)
# move boxplots away a bit by adjusting dodge
gf_boxplot(age ~ substance, 
  data = mosaicData::HELPrct, 
  color = ~sex, position = position_dodge(width = 0.9)
)
```

---

**gf_boxplot**  
*Formula interface to geom_boxplot()*

Description

Horizontal version of `geom_boxplot()`.

Usage

```r
gf_boxplot(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha, 
  color, 
  fill, 
  group, 
  linetype, 
  linewidth, 
  coef,
```
outlier.color = NULL,
outlier.fill = NULL,
outlier.shape = 19,
outlier.size = 1.5,
outlier.stroke = 0.5,
outlier.alpha = NULL,
notch = FALSE,
notchwidth = 0.5,
varwidth = FALSE,
xlab,
ylab,
title,
subtitle,
caption,
geom = "boxploth",
stat = "boxploth",
position = "dodgev",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain.
Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape y ~ x. Faceting can be achieved by including | in the
formula.

data
The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the
call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be
fortified to produce a data frame. See fortify() for which variables will be
created.
A function will be called with a single argument, the plot data. The return
value must be a data.frame, and will be used as the layer data. A function
can be created from a formula (e.g. ~ head(.x, 10)).

...
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with
attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with
attribute = value.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

fill
A color for filling, or a formula used for mapping fill.
group
Used for grouping.
linetype  A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth  A numerical line width or a formula used for mapping linewidth.

coef  Length of the whiskers as multiple of IQR. Defaults to 1.5.

outlier.color, outlier.size, outlier.stroke, outlier.shape  Default aesthetics for outliers. Set to NULL to inherit from the aesthetics used for the box. In the unlikely event you specify both US and UK spellings of colour, the US spelling will take precedence. Sometimes it can be useful to hide the outliers, for example when overlaying the raw data points on top of the boxplot. Hiding the outliers can be achieved by setting outlier.shape = NA. Importantly, this does not remove the outliers, it only hides them, so the range calculated for the y-axis will be the same with outliers shown and outliers hidden.

outlier.fill  Default aesthetics for outliers. Set to NULL to inherit from the aesthetics used for the box. In the unlikely event you specify both US and UK spellings of colour, the US spelling will take precedence. Sometimes it can be useful to hide the outliers, for example when overlaying the raw data points on top of the boxplot. Hiding the outliers can be achieved by setting outlier.shape = NA. Importantly, this does not remove the outliers, it only hides them, so the range calculated for the y-axis will be the same with outliers shown and outliers hidden.

outlier.alpha  Default aesthetics for outliers. Set to NULL to inherit from the aesthetics used for the box. In the unlikely event you specify both US and UK spellings of colour, the US spelling will take precedence. Sometimes it can be useful to hide the outliers, for example when overlaying the raw data points on top of the boxplot. Hiding the outliers can be achieved by setting outlier.shape = NA. Importantly, this does not remove the outliers, it only hides them, so the range calculated for the y-axis will be the same with outliers shown and outliers hidden.

notch  If FALSE (default) make a standard box plot. If TRUE, make a notched box plot. Notches are used to compare groups; if the notches of two boxes do not overlap, this suggests that the medians are significantly different.

notchwidth  For a notched box plot, width of the notch relative to the body (defaults to notchwidth = 0.5).

varwidth  If FALSE (default) make a standard box plot. If TRUE, boxes are drawn with widths proportional to the square-roots of the number of observations in the groups (possibly weighted, using the weight aesthetic).

xlab  Label for x-axis. See also gf_labs().

ylab  Label for y-axis. See also gf_labs().

title, subtitle, caption  Title, sub-title, and caption for the plot. See also gf_labs().

geom  A character string naming the geom used to make the layer.
stat
Position adjustment, either as a string, or the result of a call to a position adjustment function.
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. It can also be a named logical vector to finely select the aesthetics to display.
show.help
If TRUE, display some minimal help.
inherit
A logical indicating whether default attributes are inherited.
environment
An environment in which to look for variables not found in data.

Value

a gg object

Horizontal Geoms

There are two ways to obtain "horizontal" geoms: (1) The ggstance package provides a set of "horizontal" geoms and positions; (2) Thee ggplot2 now provides an orientation argument for "native" horizontal geoms and positions. ggformula supports both.

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggstance::geom_boxploth(), fivenum(), df_stats()

Examples

gf_boxploth(sex ~ age, data = mosaicData::HELPrcr, varwidth = TRUE)
gf_boxplot(sex ~ age, data = mosaicData::HELPrcr, varwidth = TRUE, orientation = 'y')
gf_boxploth(substance ~ age, data = mosaicData::HELPrcr, color = ~sex)
# move boxplots away a bit by adjusting dodge
gf_boxploth(substance ~ age, data = mosaicData::HELPrcr, color = ~sex,
position = position_dodgev(height = 0.9)
)
# gf_boxplot guesses horizontal because substance is categorical
gf_boxplot(substance ~ age,
  data = mosaicData::HELPrct, color = ~sex,
  position = position_dodgev(height = 0.9)
)
gf_boxploth(substance ~ age, data = mosaicData::HELPrct, color = ~sex, outlier.color = "gray50")
# longer whiskers
gf_boxploth(substance ~ age, data = mosaicData::HELPrct, color = ~sex, coef = 2)
# Note: height for boxplots is full width of box.
# For jittering, it is the half-height.
gf_boxploth(substance ~ age | sex, data = mosaicData::HELPrct, coef = 5, height = 0.4) %>%
gf_jitter(height = 0.2, alpha = 0.3)

# combining boxplots and histograms
gf_histogram(~eruptions, data = faithful) %>%
gf_boxploth(0 ~ eruptions, alpha = 0, width = 2)
gf_histogram(~eruptions, data = faithful) %>%
gf_boxploth(-2 ~ eruptions, alpha = 0, width = 2)
gf_histogram(~eruptions, data = faithful) %>%
gf_boxploth(32 ~ eruptions, alpha = 0, width = 2)

---

**gf_col**  
*Formula interface to geom_col()*

**Description**

There are two types of bar charts: `geom_bar()` and `geom_col()`. `geom_bar()` makes the height of the bar proportional to the number of cases in each group (or if the weight aesthetic is supplied, the sum of the weights). If you want the heights of the bars to represent values in the data, use `geom_col()` instead. `geom_bar()` uses `stat_count()` by default: it counts the number of cases at each x position. `geom_col()` uses `stat_identity()`: it leaves the data as is.

**Usage**

```r
gf_col(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  xlab,
  ylab,
```

title, subtitle, caption,
geom = "col",
stat = "identity",
position = "stack",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption
geom A character string naming the geom used to make the layer.
stat A character string naming the stat used to make the layer.
position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.
show.help If TRUE, display some minimal help.
inherit A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.
Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_col()`

Examples

```r
SomeData <- data.frame(
  group = LETTERS[1:3],
  count = c(20, 25, 18)
)
gf_col(count ~ group, data = SomeData)

# A Pareto chart

if (require(dplyr) && require(mosaicData)) {
  HELPrct %>%
    group_by(substance) %>%
    summarise(count = n()) %>%
    ungroup() %>%
    dplyr::arrange(-count) %>%
    mutate(
      cumcount = cumsum(count),
      substance = reorder(substance, -count)
    ) %>%
    gf_col(count ~ substance, fill = "skyblue") %>%
    gf_point(cumcount ~ substance) %>%
    gf_line(cumcount ~ substance, group = 1) %>%
    gf_refine(
      scale_y_continuous(sec.axis = sec_axis(~ . / nrow(HELPrct)))
    )
}
```
gf_contour  

Formula interface to geom_contour() and geom_contour_filled()

Description

ggplot2 can not draw true 3D surfaces, but you can use geom_contour(), geom_contour_filled(), and geom_tile() to visualise 3D surfaces in 2D.

These functions require regular data, where the x and y coordinates form an equally spaced grid, and each combination of x and y appears once. Missing values of z are allowed, but contouring will only work for grid points where all four corners are non-missing. If you have irregular data, you’ll need to first interpolate on to a grid before visualising, using interp::interp(), akima::bilinear(), or similar.

Usage

gf_contour(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "contour",
  stat = "contour",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_contour_filled(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "contour_filled",
  stat = "contour_filled",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
...
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
xlab Label for x-axis. See also `gf_labs()`.
ylab Label for y-axis. See also `gf_labs()`.
title, subtitle, caption Title, sub-title, and caption for the plot. See also `gf_labs()`.
geom The geometric object to use to display the data, either as a ggproto Geom subclass or as a string naming the geom stripped of the geom_ prefix (e.g. "point" rather than "geom_point")
stat The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")
position Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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. It can also be a named logical vector to finely select the aesthetics to display.
show.help If TRUE, display some minimal help.
inherit A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.
gf_count

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_contour(), gf_density_2d()

Examples

gf_density_2d(eruptions ~ waiting, data = faithful, alpha = 0.5, color = "navy") %>%
gf_contour(density ~ waiting + eruptions, data = faithful, bins = 10, color = "red")
gf_contour_filled(density ~ waiting + eruptions, data = faithful, bins = 10,
    show.legend = FALSE) %>%
gf_jitter(eruptions ~ waiting, data = faithful, color = "white", alpha = 0.5,
    inherit = FALSE)

gf_count

Formula interface to geom_count()

Description

This is a variant geom_point() that counts the number of observations at each location, then maps the count to point area. It useful when you have discrete data and overplotting.

Usage

gf_count(
    object = NULL,
    gformula = NULL,
    data = NULL,
    ...,
Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.  
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
group Used for grouping.
shape An integer or letter shape or a formula used for mapping shape.
size A numeric size or a formula used for mapping size.
stroke A numeric size of the border or a formula used to map stroke.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom A character string naming the geom used to make the layer.
specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and
mapping of additional attributes can be done through the use of additional arguments. Attributes
can be set can be set using arguments of the form attribute = value or mapped using arguments
of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid().
This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel
more familiar to users of lattice.

evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the
right thing when formulas are created on the fly, but might not be the right thing if formulas created
in one environment are used to create plots in another.

See Also

ggplot2::geom_count()

Examples

# Best used in conjunction with scale_size_area which ensures that
# counts of zero would be given size 0. This doesn't make much difference
# here because the smallest count is already close to 0.

gf_count(hwy ~ cty, data = mpg, alpha = 0.3) %>%
gf_refine(scale_size_area())
**gf_crossbar**

*Formula interface to geom_crossbar()*

**Description**

Various ways of representing a vertical interval defined by \( x, \) \( y_{\text{min}} \) and \( y_{\text{max}} \). Each case draws a single graphical object.

**Usage**

```r
gf_crossbar(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha, 
  color, 
  group, 
  linetype, 
  linewidth, 
  fatten = 2.5, 
  xlab, 
  ylab, 
  title, 
  subtitle, 
  caption, 
  geom = "crossbar", 
  stat = "identity", 
  position = "identity", 
  show.legend = NA, 
  show.help = NULL, 
  inherit = TRUE, 
  environment = parent.frame()
)

gf_crossbarh(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha, 
  color, 
  group, 
  linetype, 
  linewidth, 
  fatten = 2.5, 
  xlab,
```


Arguments

object

When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula

A formula with shape y + ymin + ymax ~ x. Faceting can be achieved by including | in the formula.

data

The data to be displayed in this layer. There are three options:

If NULL, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.

A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha

Opacity (0 = invisible, 1 = opaque).

color

A color or a formula used for mapping color.

group

Used for grouping.

linetype

A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth

A numerical line width or a formula used for mapping linewidth.

fatten

A multiplicative factor used to increase the size of the middle bar in `geom_crossbar()` and the middle point in `geom_pointrange()`.

xlab

Label for x-axis. See also `gf_labs()`.

ylab

Label for y-axis. See also `gf_labs()`.

title, subtitle, caption

Title, sub-title, and caption for the plot. See also `gf_labs()`.
gf_crossbar

<table>
<thead>
<tr>
<th>Argument</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>geom</td>
<td>A character string naming the geom used to make the layer.</td>
</tr>
<tr>
<td>stat</td>
<td>The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. &quot;count&quot; rather than &quot;stat_count&quot;)</td>
</tr>
<tr>
<td>position</td>
<td>Position adjustment, either as a string naming the adjustment (e.g. &quot;jitter&quot; to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.</td>
</tr>
<tr>
<td>show.legend</td>
<td>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. It can also be a named logical vector to finely select the aesthetics to display.</td>
</tr>
<tr>
<td>show.help</td>
<td>If TRUE, display some minimal help.</td>
</tr>
<tr>
<td>inherit</td>
<td>A logical indicating whether default attributes are inherited.</td>
</tr>
<tr>
<td>environment</td>
<td>An environment in which to look for variables not found in data.</td>
</tr>
</tbody>
</table>

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_crossbar()

Examples

```r
if (require(mosaicData) && require(dplyr)) {
  HELP2 <- HELPrt %>%
    group_by(substance, sex) %>%
    summarise(
      mean.age = mean(age),
      median.age = median(age),
      max.age = max(age),
      min.age = min(age),
```


gf_jitter(age ~ substance, data = HELPrct,
    alpha = 0.7, width = 0.2, height = 0, color = "skyblue") %>%
gf_pointrange(mean.age + lo + hi ~ substance, data = HELP2) %>%
gf_facet_grid(~sex)

gf_jitter(age ~ substance, data = HELPrct,
    alpha = 0.7, width = 0.2, height = 0, color = "skyblue") %>%
gf_errorbar(lo + hi ~ substance, data = HELP2, inherit = FALSE) %>%
gf_facet_grid(~sex)

gf_jitter(age ~ substance, data = HELPrct,
    alpha = 0.7, width = 0.2, height = 0, color = "skyblue") %>%
gf_crossbar(mean.age + lo + hi ~ substance, data = HELP2,
    fill = "transparent") %>%
gf_facet_grid(~sex)

gf_jitter(substance ~ age, data = HELPrct,
    alpha = 0.7, height = 0.2, width = 0, color = "skyblue") %>%
gf_crossbarh(substance ~ mean.age + lo + hi, data = HELP2,
    fill = "transparent", color = "red") %>%
gf_facet_grid(~sex)

---------------------------------------------------------------
gf_curve

Formula interface to geom_curve()

Description

gf_curve() draws a curved line. See the underlying drawing function grid::curveGrob() for the parameters that control the curve.

Usage

gf_curve(
    object = NULL,
    gformula = NULL,
    data = NULL,
    ...,
    alpha,
    color,
    group,
    linetype,
Arguments

**object**
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

**gformula**
A formula with shape \( y + y_{\text{end}} \sim x + x_{\text{end}} \).

**data**
The data to be displayed in this layer. There are three options:
- If `NULL`, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.
- A `data.frame`, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.
- A function will be called with a single argument, the plot data. The return value must be a data frame, and will be used as the layer data. A function can be created from a formula (e.g. \(~ \text{head}(.x, 10)\)).

**...**
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with `attribute = value`, (b) ggplot2 aesthetics to be mapped with `attribute = ~ expression`, or (c) attributes of the layer as a whole, which are set with `attribute = value`.

**alpha**
Opacity (0 = invisible, 1 = opaque).

**color**
A color or a formula used for mapping color.

**group**
Used for grouping.

**linetype**
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

**linewidth**
A numerical line width or a formula used for mapping linewidth.

**curvature**
A numeric value giving the amount of curvature. Negative values produce left-hand curves, positive values produce right-hand curves, and zero produces a straight line.
**gf_curve**

<table>
<thead>
<tr>
<th>Attribute</th>
<th>Description</th>
</tr>
</thead>
<tbody>
<tr>
<td>angle</td>
<td>A numeric value between 0 and 180, giving an amount to skew the control points of the curve. Values less than 90 skew the curve towards the start point and values greater than 90 skew the curve towards the end point.</td>
</tr>
<tr>
<td>ncp</td>
<td>The number of control points used to draw the curve. More control points creates a smoother curve.</td>
</tr>
<tr>
<td>arrow</td>
<td>Specification for arrow heads, as created by <code>grid::arrow()</code>.</td>
</tr>
<tr>
<td>lineend</td>
<td>Line end style (round, butt, square).</td>
</tr>
<tr>
<td>xlab</td>
<td>Label for x-axis. See also <code>gf_labs()</code>.</td>
</tr>
<tr>
<td>ylab</td>
<td>Label for y-axis. See also <code>gf_labs()</code>.</td>
</tr>
<tr>
<td>title, subtitle, caption</td>
<td>Title, sub-title, and caption for the plot. See also <code>gf_labs()</code>.</td>
</tr>
<tr>
<td>geom</td>
<td>A character string naming the geom used to make the layer.</td>
</tr>
<tr>
<td>stat</td>
<td>The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the <code>stat_</code> prefix (e.g. &quot;count&quot; rather than &quot;stat_count&quot;)</td>
</tr>
<tr>
<td>position</td>
<td>Position adjustment, either as a string naming the adjustment (e.g. &quot;jitter&quot; to use <code>position_jitter</code>), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.</td>
</tr>
<tr>
<td>show.legend</td>
<td>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. It can also be a named logical vector to finely select the aesthetics to display.</td>
</tr>
<tr>
<td>show.help</td>
<td>If TRUE, display some minimal help.</td>
</tr>
<tr>
<td>inherit</td>
<td>A logical indicating whether default attributes are inherited.</td>
</tr>
<tr>
<td>environment</td>
<td>An environment in which to look for variables not found in data.</td>
</tr>
</tbody>
</table>

**Value**

a gg object

**Specifying plot attributes**

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

**Evaluation**

Evaluation of the `ggplot2` code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.
gf_density

See Also

ggplot2::geom_curve()

Examples

D <- data.frame(x1 = 2.62, x2 = 3.57, y1 = 21.0, y2 = 15.0)
gf_point(mpg ~ wt, data = mtcars) %>%
gf_curve(y1 + y2 ~ x1 + x2, data = D, color = "navy") %>%
gf_segment(y1 + y2 ~ x1 + x2, data = D, color = "red")

gf_density

Description

Computes and draws a kernel density estimate, which is a smoothed version of the histogram and
is a useful alternative when the data come from an underlying smooth distribution. The only differ-
ence between gf_dens() and gf_density() is the default geom used to show the density curve:
gf_density() uses an area geom (which can be filled). gf_dens() using a line geom (which
cannot be filled).

Usage

gf_density(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha = 0.5,
  color,
  fill,
  group,
  linetype,
  linewidth,
  kernel = "gaussian",
  n = 512,
  trim = FALSE,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "area",
  stat = "density",
  position = "identity",
  show.legend = NA,
  show.help = NULL,)
gf_density

    inherit = TRUE,
    environment = parent.frame()
  )

gf_dens(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha = 0.5,
  color,
  fill = NA,
  group,
  linetype,
  linewidth,
  kernel = "gaussian",
  n = 512,
  trim = FALSE,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "line",
  stat = "density",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
  )

gf_dens2(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha = 0.5,
  color,
  fill = NA,
  group,
  linetype,
  linewidth,
  kernel = "gaussian",
  n = 512,
  trim = FALSE,
  xlab,
  ylab,
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape ~ x. Faceting can be achieved by including ~ in the formula.

data
The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

fill
A color for filling, or a formula used for mapping fill.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

kernel
Kernel. See list of available kernels in density().

n
number of equally spaced points at which the density is to be estimated, should be a power of two, see density() for details

trim
If FALSE, the default, each density is computed on the full range of the data. If TRUE, each density is computed over the range of that group: this typically means the estimated x values will not line-up, and hence you won’t be able to stack density values. This parameter only matters if you are displaying multiple densities in one plot or if you are manually adjusting the scale limits.
gf_density

xlab  Label for x-axis. See also gf_labs().
ylab  Label for y-axis. See also gf_labs().
title, subtitle, caption
   Title, sub-title, and caption for the plot. See also gf_labs().
geom, stat
   Use to override the default connection between geom_density() and stat_density().
position
   Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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. It can also be a named logical vector to finely select the aesthetics to display.
show.help
   If TRUE, display some minimal help.
inherit
   A logical indicating whether default attributes are inherited.
environment
   An environment in which to look for variables not found in data.

Value
   a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also
   gf_ash(), ggplot2::geom_density()

Examples
   gf_dens()
data(penguins, package = "palmerpenguins")
gf_density(~bill_length_mm, fill = ~species, data = penguins)
gf_dens(~bill_length_mm, color = ~species, data = penguins)
gf_dens2(~bill_length_mm, color = ~species, fill = ~species, data = penguins)
gf_freqpoly(~bill_length_mm, color = ~species, data = penguins, bins = 15)
# Chaining in the data

data(penguins, package = "palmerpenguins")
penguins %>% gf_dens(~bill_length_mm, color = ~species)
# horizontal orientation
penguins %>% gf_dens(bill_length_mm ~ ., color = ~species)

gf_density_2d  

*Formula interface to geom_density_2d() and geom_density_2d_filled()*

**Description**

Perform a 2D kernel density estimation using `MASS::kde2d()` and display the results with contours. This can be useful for dealing with overplotting. This is a 2D version of `geom_density()`. `geom_density_2d()` draws contour lines, and `geom_density_2d_filled()` draws filled contour bands.

**Usage**

```r
gf_density_2d(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  contour = TRUE,
  n = 100,
  h = NULL,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "density_2d",
  stat = "density_2d",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)```
```
gf_density_2d

)

gf_density_2d_filled(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
  color,
  group,
  linetype,
  linewidth,
  contour = TRUE,
  n = 100,
  h = NULL,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "density_2d_filled",
  stat = "density_2d_filled",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_density2d(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
  color,
  group,
  linetype,
  linewidth,
  contour = TRUE,
  n = 100,
  h = NULL,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
Arguments

object  When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
A formula with shape \( y \sim x \). Faceting can be achieved by including | in the formula.

The data to be displayed in this layer. There are three options:

- If NULL, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.
- A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.
- A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. \(~ \text{head}(x, 10)\)).

Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with `attribute = value`, (b) ggplot2 aesthetics to be mapped with `attribute = ~ expression`, or (c) attributes of the layer as a whole, which are set with `attribute = value`.

Opacity \((0 = \text{invisible}, 1 = \text{opaque})\).

A color or a formula used for mapping color.

Used for grouping.

A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

A numerical line width or a formula used for mapping linewidth.

If TRUE, contour the results of the 2d density estimation.

Number of grid points in each direction.

Bandwidth (vector of length two). If NULL, estimated using `MASS::bandwidth.nrd()`.

Line end style (round, butt, square).

Line join style (round, mitre, bevel).

Line mitre limit (number greater than 1).

Label for x-axis. See also `gf_labs()`.

Label for y-axis. See also `gf_labs()`.

Title, sub-title, and caption for the plot. See also `gf_labs()`.

Use to override the default connection between `geom_density_2d()` and `stat_density_2d()`.

Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

If TRUE, display some minimal help.

A logical indicating whether default attributes are inherited.

An environment in which to look for variables not found in data.
Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_density_2d()`

Examples

```r
gf_jitter(avg_drinks ~ age,
  alpha = 0.2, data = mosaicData::HELPrct,
  width = 0.4, height = 0.4
) %>%
gf_density_2d(avg_drinks ~ age, data = mosaicData::HELPrct)
gf_density_2d_filled(avg_drinks ~ age, data = mosaicData::HELPrct, show.legend = FALSE) %>%
gf_jitter(avg_drinks ~ age,
  alpha = 0.3, data = mosaicData::HELPrct,
  width = 0.4, height = 0.4,
  color = "white"
)
gf_jitter(avg_drinks ~ age,
  alpha = 0.2, data = mosaicData::HELPrct,
  width = 0.4, height = 0.4
) %>%
gf_density2d(avg_drinks ~ age, data = mosaicData::HELPrct)
gf_density2d_filled(avg_drinks ~ age, data = mosaicData::HELPrct, show.legend = FALSE) %>%
gf_jitter(avg_drinks ~ age,
  alpha = 0.4, data = mosaicData::HELPrct,
  width = 0.4, height = 0.4,
  color = "white"
)```
Description

Create a layer displaying a probability distribution.

Usage

gf_dist(
  object = ggplot(),
  dist,
  ...,
  xlim = NULL,
  kind = c("density", "cdf", "qq", "qqstep", "histogram"),
  resolution = 5000L,
  eps = 1e-06,
  params = NULL
)

Arguments

- object: a gg object.
- dist: A character string providing the name of a distribution. Any distribution for which the functions with names formed by prepending "d", "p", or "q" to dist exist can be used.
- ...: additional arguments passed both to the distribution functions and to the layer. Note: Possible ambiguities using params or by preceding plot argument with plot_.
- xlim: A numeric vector of length 2 providing lower and upper bounds for the portion of the distribution that will be displayed. The default is to attempt to determine reasonable bounds using quantiles of the distribution.
- kind: One of "density", "cdf", "qq", "qqstep", or "histogram" describing what kind of plot to create.
- resolution: An integer specifying the number of points to use for creating the plot.
- eps: a (small) numeric value. When other defaults are not available, the distribution is processed from the eps to 1 - eps quantiles.
- params: a list of parameters for the distribution.

Examples

gf_dhistogram(~ rnorm(100), bins = 20) %>%
gf_dist("norm", color = "red")

# shading tails -- but see pdist() for this
gf_dist("norm", fill = ~ (abs(x) <= 2), geom = "area")
gf_dist("norm", color = "red", kind = "cdf")
gf_dist("norm", fill = "red", kind = "histogram")
gf_dist("norm", color = "red", kind = "qqstep", resolution = 25) %>%
gf_dist("norm", color = "black", kind = "qq", resolution = 25, linewidth = 2, alpha = 0.5)
# size is used as parameter for binomial distribution
gf_dist("binom", size = 20, prob = 0.25)
# If we want to adjust size argument for plots, we have two choices:
gf_dist("binom", size = 20, prob = 0.25, plot_size = 2)
gf_dist("binom", params = list(size = 20, prob = 0.25), size = 2)

gf_dotplot                      Formula interface to geom_dotplot()

Description

Scatterplots in ggformula.

Usage

gf_dotplot(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  fill,
  group,
  binwidth = NULL,
  binaxis = "x",
  method = "dotdensity",
  binpositions = "bygroup",
  stackdir = "up",
  stackratio = 1,
  dotsize = 1,
  stackgroups = FALSE,
  origin = NULL,
  right = TRUE,
  width = 0.9,
  drop = FALSE,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  position = "identity",
  show.legend = NA,
Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
group Used for grouping.
binwidth When method is "dotdensity", this specifies maximum bin width. When method is "histodot", this specifies bin width. Defaults to 1/30 of the range of the data.
binaxis The axis to bin along, "x" (default) or "y"
method "dotdensity" (default) for dot-density binning, or "histodot" for fixed bin widths (like stat_bin)
binpositions When method is "dotdensity", "bygroup" (default) determines positions of the bins for each group separately. "all" determines positions of the bins with all the data taken together; this is used for aligning dot stacks across multiple groups.
stackdir which direction to stack the dots. "up" (default), "down", "center", "centerwhole" (centered, but with dots aligned)
stackratio how close to stack the dots. Default is 1, where dots just touch. Use smaller values for closer, overlapping dots.
dotsize The diameter of the dots relative to binwidth, default 1.
stackgroups should dots be stacked across groups? This has the effect that position = "stack" should have, but can’t (because this geom has some odd properties).
origin When method is "histodot", origin of first bin
right When method is "histodot", should intervals be closed on the right (a, b], or not [a, b)
width When binaxis is "y", the spacing of the dot stacks for dodging.
drop If TRUE, remove all bins with zero counts
xlab Label for x-axis. See also gf_labs()
ylab Label for y-axis. See also gf_labs()
Title, sub-title, and caption
Title, sub-title, and caption for the plot. See also gf_labs().

position
Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend
A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.

Details
There are two basic approaches: dot-density and histodot. With dot-density binning, the bin positions are determined by the data and binwidth, which is the maximum width of each bin. See Wilkinson (1999) for details on the dot-density binning algorithm. With histodot binning, the bins have fixed positions and fixed widths, much like a histogram.

When binning along the x axis and stacking along the y axis, the numbers on y axis are not meaningful, due to technical limitations of ggplot2. You can hide the y axis, as in one of the examples, or manually scale it to match the number of dots.

Value
a gg object

Warning
Dotplots in ggplot2 (and hence in ggformula) often require some fiddling because the default y-axis is meaningless and the ideal size of the dots depends on the aspect ratio of the plot.

Specifying plot attributes
Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation
Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.
**gf_ecdf**  

**References**  

**See Also**  
ggplot2::geom_dotplot()

**Examples**  
```r  
data(penguins, package = "palmerpenguins")  
gf_dotplot(~bill_length_mm, fill = ~species, data = penguins)  
```
Arguments

object  When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data  The data to be displayed in this layer. There are three options:
  If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
  A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
  A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
...
  Other arguments passed on to layer(). These are often aesthetics, used to set an aesthetic to a fixed value, like colour = "red" or size = 3. They may also be parameters to the paired geom/stat.
group  Used for grouping.
pad  If TRUE, pad the ecdf with additional points (-Inf, 0) and (Inf, 1)
n  if NULL, do not interpolate. If not NULL, this is the number of points to interpolate with.
xlab  Label for x-axis. See also gf_labs().
ylab  Label for y-axis. See also gf_labs().
title, subtitle, caption  Title, sub-title, and caption for the plot. See also gf_labs().
geom  The geometric object to use to display the data, either as a ggproto Geom subclass or as a string naming the geom stripped of the geom_ prefix (e.g. "point" rather than "geom_point")
stat  The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")
position  Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
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. It can also be a named logical vector to finely select the aesthetics to display.
show.help  If TRUE, display some minimal help.
inherit  A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.
Examples

Data <- data.frame(
  x = c(rnorm(100, 0, 1), rnorm(100, 0, 3), rt(100, df = 3)),
  g = gl(3, 100, labels = c("N(0, 1)", "N(0, 3)", "T(df = 3)")
)
gf_ecdf(~ x, data = Data)
  # Don't go to positive/negative infinity
  gf_ecdf(~ x, data = Data, pad = FALSE)
  # Multiple ECDFs
  gf_ecdf(~ x, data = Data, color = ~ g)

---

gf_ellipse  Function interface to stat_ellipse()

Description

Formula interface to ggplot2::stat_ellipse().

Usage

gf_ellipse(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  type = "t",
  level = 0.95,
  segments = 51,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "path",
  stat = "ellipse",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)
Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
group Used for grouping.
type The type of ellipse. The default "t" assumes a multivariate t-distribution, and "norm" assumes a multivariate normal distribution. "euclid" draws a circle with the radius equal to level, representing the euclidean distance from the center. This ellipse probably won’t appear circular unless coord_fixed() is applied.
level The level at which to draw an ellipse, or, if type="euclid", the radius of the circle to be drawn.
segments The number of segments to be used in drawing the ellipse.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom Geom for drawing ellipse. Note: "polygon" allows fill; "path" does not; on the other hand, "path" allows alpha to be applied to the border, while "polygon" applies alpha only to the interior.
stat A character string naming the stat used to make the layer.
position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.
show.help If TRUE, display some minimal help.
inherit A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.

See Also

ggplot2::stat_ellipse()
gf_empty

Create an "empty" plot

Description

This is primarily useful as a way to start a sequence of piped plot layers.

Usage

gf_empty(environment = parent.frame())

Arguments

  environment An environment passed to ggplot2::ggplot()

Value

A plot with now layers.
Examples

gf_empty()
data(penguins, package = "palmerpenguins")
gf_empty() %>%
  gf_point(bill_length_mm ~ bill_depth_mm, data = penguins, color = ~species)

gf_errorbar

Formula interface to geom_errorbar()

Description

For each x value, geom_ribbon() displays a y interval defined by ymin and ymax. geom_area() is a special case of geom_ribbon(), where the ymin is fixed to 0 and y is used instead of ymax.

Usage

gf_errorbar(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "errorbar",
  stat = "identity",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula A formula with shape ymin + ymax ~ x. Faceting can be achieved by including | in the formula.
The data to be displayed in this layer. There are three options:

If `NULL`, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.

A `data.frame`, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a `data.frame`, and will be used as the layer data. A function can be created from a formula (e.g. `~ head(.x, 10)`).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with `attribute = value`, (b) ggplot2 aesthetics to be mapped with `attribute = ~ expression`, or (c) attributes of the layer as a whole, which are set with `attribute = value`.

- `alpha` Opacity (0 = invisible, 1 = opaque).
- `color` A color or a formula used for mapping color.
- `group` Used for grouping.
- `linetype` A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
- `linewidth` A numerical line width or a formula used for mapping linewidth.
- `xlab` Label for x-axis. See also `gf_labs()`.
- `ylab` Label for y-axis. See also `gf_labs()`.
- `title, subtitle, caption` Title, sub-title, and caption for the plot. See also `gf_labs()`.
- `geom` The geometric object to use to display the data, either as a ggproto Geom subclass or as a string naming the geom stripped of the geom_ prefix (e.g. "point" rather than "geom_point").
- `stat` The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count").
- `position` Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
- `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. It can also be a named logical vector to finely select the aesthetics to display.
- `show.help` If `TRUE`, display some minimal help.
- `inherit` A logical indicating whether default attributes are inherited.
- `environment` An environment in which to look for variables not found in data.

**See Also**

`ggplot2::geom_errorbar()`
Examples

```r
if (require(mosaicData) && require(dplyr)) {
HELP2 <- HELPrct %>%
group_by(substance, sex) %>%
summarise(
  mean.age = mean(age),
  median.age = median(age),
  max.age = max(age),
  min.age = min(age),
  sd.age = sd(age),
  lo = mean.age - sd.age,
  hi = mean.age + sd.age
)

gf_jitter(age ~ substance, data = HELPrct,
  alpha = 0.5, width = 0.2, height = 0, color = "skyblue") %>%
gf_pointrange(mean.age + lo + hi ~ substance, data = HELP2,
  inherit = FALSE) %>%
gf_facet_grid(~sex)

gf_jitter(age ~ substance, data = HELPrct,
  alpha = 0.5, width = 0.2, height = 0, color = "skyblue") %>%
gf_errorbar(lo + hi ~ substance, data = HELP2, inherit = FALSE) %>%
gf_facet_grid(~sex)

gf_jitter(age ~ substance, data = HELPrct,
  alpha = 0.5, width = 0.2, height = 0, color = "skyblue") %>%
gf_boxplot(age ~ substance, data = HELPrct, color = "red") %>%
gf_crossbar(mean.age + lo + hi ~ substance, data = HELP2) %>%
gf_facet_grid(~sex)
}
```

---

gf_errorbarh  

*Formula interface to geom_errorbarh()*

**Description**

A rotated version of `geom_errorbar()`.  

**Usage**

```r
gf_errorbarh(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  linetype,
```
gf_errorbarh

```r
  linewidth,
xlab,
ylab,
title,
subtitle,
caption,
geom = "errorbarh",
stat = "identity",
position = "identity",
show.legend = "identity",
show.help = NA,
inherit = TRUE,
environment = parent.frame()
)
```

**Arguments**

- `object` When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

- `gformula` A formula with shape `y ~ x ± xmin + xmax`. Faceting can be achieved by including `|` in the formula.

- `data` The data to be displayed in this layer. There are three options:
  - If `NULL`, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.
  - A `data.frame`, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.
  - A function will be called with a single argument, the plot data. The return value must be a `data.frame`, and will be used as the layer data. A function can be created from a formula (e.g. `~ head(.x, 10)`).

- `...` Additional arguments. Typically these are (a) `ggplot2` aesthetics to be set with `attribute = value`, (b) `ggplot2` aesthetics to be mapped with `attribute = ~ expression`, or (c) attributes of the layer as a whole, which are set with `attribute = value`.

- `alpha` Opacity (0 = invisible, 1 = opaque).

- `color` A color or a formula used for mapping color.

- `group` Used for grouping.

- `linetype` A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

- `linewidth` A numerical line width or a formula used for mapping linewidth.

- `xlab` Label for x-axis. See also `gf_labs()`.

- `ylab` Label for y-axis. See also `gf_labs()`.

- `title, subtitle, caption` Title, sub-title, and caption for the plot. See also `gf_labs()`.

- `geom` A character string naming the geom used to make the layer.
stat

The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")

position

Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help

If TRUE, display some minimal help.

inherit

A logical indicating whether default attributes are inherited.

environment

An environment in which to look for variables not found in data.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a. aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_errorbarh()

Examples

if (require(dplyr)) {
  HELP2 <- mosaicData::HELPrct %>%
    group_by(substance, sex) %>%
    summarise(
      mean.age = mean(age),
      median.age = median(age),
      max.age = max(age),
      min.age = min(age),
      sd.age = sd(age),
    }
gf.facet_wrap

\[
\begin{align*}
lo &= \text{mean.age} - \text{sd.age}, \\
hi &= \text{mean.age} + \text{sd.age}
\end{align*}
\]

gf.jitter(substance ~ age, data = mosaicData::HELPct, 
alpha = 0.5, height = 0.2, width = 0, color = "skyblue")
gf.errorbarh(substance ~ lo + hi, data = HELP2, inherit = FALSE) 
gf.facet.grid(~sex)

gf.jitter(age ~ substance, data = mosaicData::HELPct, 
alpha = 0.5, width = 0.2, height = 0, color = "skyblue")
gf.errorbar(lo + hi ~ substance, data = HELP2, inherit = FALSE) 
gf.facet.grid(~sex)

---

gf.facet_wrap  Add facets to a plot

Description

These functions provide more control over faceting than is possible using the formula interface.

Usage

\[
gf\text{-facet}\text{\_wrap}(\text{object}, \ldots)
\]

\[
gf\text{-facet}\text{\_grid}(\text{object}, \ldots)
\]

Arguments

object  A ggplot object

\[
\ldots
\]

Additional arguments passed to \texttt{facet\_wrap()} or \texttt{facet\_grid()}. This typically includes an unnamed formula argument describing the facets. scales and space are additional useful arguments. See the examples.

See Also

\texttt{ggplot2::facet\_grid()}, \texttt{ggplot2::facet\_wrap()}.

Examples

\[
gf\text{-histogram}(\text{avg.drink} ~ \text{data} = \text{mosaicData::HELPct}) \\
gf\text{-facet}\_grid(~\text{substance})
\]

\[
gf\text{-histogram}(\text{avg.drink} ~ \text{data} = \text{mosaicData::HELPct}) \\
gf\text{-facet}\_grid(~\text{substance}, \text{scales} = "free")
\]

\[
gf\text{-histogram}(\text{avg.drink} ~ \text{data} = \text{mosaicData::HELPct}) \\
gf\text{-facet}\_grid(~\text{substance}, \text{scales} = "free", \text{space} = "free")
\]

\[
gf\text{-line}(\text{birth} ~ \text{date} ~ \text{data} = \text{mosaicData::Births}, \text{color} = \sim\text{wday}) \\
gf\text{-facet}\_wrap(~\text{year}, \text{scales} = "free\_x", \text{nrow} = 5)
\]
MASS::fitdistr() is used to fit coefficients of a specified family of distributions and the resulting density curve is displayed.

Usage

gf_fitdistr(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  dist = "dnorm",
  start = NULL,
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  size,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "path",
  stat = "fitdistr",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = FALSE,
  environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See examples.
gf_fitdistr

| gformula | A formula with shape ~ x used to specify the data to be fit to a family of distributions. |
| data | A data frame containing the variable to be fitted. |
| ... | Additional arguments |
| dist | A quoted name of a distribution function. See mosaicCore::fit_distr_fun() for more details about allowable distributions. |
| start | Starting value(s) for the search for MLE. (See MASS::fitdistr.) |
| alpha | Opacity (0 = invisible, 1 = opaque). |
| color | A color or a formula used for mapping color. |
| fill | A color for filling, or a formula used for mapping fill. |
| group | Used for grouping. |
| linetype | A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype. |
| linewidth | A numerical line width or a formula used for mapping linewidth. |
| size | Size aesthetic for dots in pmf plots. |
| xlab | Label for x-axis. See also gf_labs(). |
| ylab | Label for y-axis. See also gf_labs(). |
| title,subtitle,caption | Title, sub-title, and caption for the plot. See also gf_labs(). |
| geom | A character string naming the geom used to make the layer. |
| stat | A character string naming the stat used to make the layer. |
| position | Either a character string naming the position function used for the layer or a position object returned from a call to a position function. |
| show.legend | A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped. |
| show.help | If TRUE, display some minimal help. |
| inherit | A logical indicating whether default attributes are inherited. |
| environment | An environment in which to look for variables not found in data. |

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.
Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`mosaicCore::fit_distr_fun()`

Examples

```r
gf_fitdistr(~length, data = mosaicData::KidsFeet, inherit = FALSE) %>%
gf_dhistogram(~length, data = mosaicData::KidsFeet, binwidth = 0.5, alpha = 0.25)

gf_dhistogram(~length, data = mosaicData::KidsFeet, binwidth = 0.5, alpha = 0.25) %>%
gf_fitdistr()

set.seed(12345)
Dat <- data.frame(
  f = rf(500, df1 = 3, df2 = 47),
  g = rgamma(500, 3, 10)
)
gf_dhistogram(~g, data = Dat) %>%
gf_fitdistr(dist = "dgamma", linewidth = 1.4)

gf_dhistogram(~g, data = Dat) %>%
gf_fun(mosaicCore::fit_distr_fun(~g, data = Dat, dist = "dgamma"))

gf_dhistogram(~f, data = Dat) %>%
gf_fitdistr(dist = "df", start = list(df1 = 2, df2 = 50))

# fitted parameters are default argument values
args(
  mosaicCore::fit_distr_fun(~f,
    data = Dat, dist = "df",
    start = list(df1 = 2, df2 = 50)
  )
)
args(mosaicCore::fit_distr_fun(~g, data = Dat, dist = "dgamma"))
```

---

gf_freqpoly

**Formula interface to geom_freqpoly()**

Description

Visualise the distribution of a single continuous variable by dividing the x axis into bins and counting the number of observations in each bin. Histograms (geom_histogram()) display the counts with bars; frequency polygons (geom_freqpoly()) display the counts with lines. Frequency polygons are more suitable when you want to compare the distribution across the levels of a categorical variable.
Usage

gf_freqpoly(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha, 
  color, 
  group, 
  linetype, 
  linewidth, 
  binwidth, 
  bins, 
  center, 
  boundary, 
  xlab, 
  ylab, 
  title, 
  subtitle, 
  caption, 
  geom = "path", 
  stat = "bin", 
  position = "identity", 
  show.legend = NA, 
  show.help = NULL,
  inherit = TRUE, 
  environment = parent.frame()
)

Arguments

object      When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula   A formula with shape ~ x or y ~ x. Faceting can be achieved by including | in the formula.

data        The data to be displayed in this layer. There are three options:
             If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
             A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
             A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

...        Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute =
~ expression, or (c) attributes of the layer as a whole, which are set with 
attribute = value.

alpha       Opacity (0 = invisible, 1 = opaque).

color       A color or a formula used for mapping color.

group       Used for grouping.

linetype     A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping 
linetype.

linewidth    A numerical line width or a formula used for mapping linewidth.

binwidth     The width of the bins. Can be specified as a numeric value or as a function that 
calculates width from unscaled x. Here, "unscaled x" refers to the original x val-
ues in the data, before application of any scale transformation. When specifying 
a function along with a grouping structure, the function will be called once per 
group. The default is to use the number of bins in bins, covering the range of 
the data. You should always override this value, exploring multiple widths to 
find the best to illustrate the stories in your data.

The bin width of a date variable is the number of days in each time; the bin 
width of a time variable is the number of seconds.

bins        Number of bins. Overridden by binwidth. Defaults to 30.

center, boundary 
bin position specifiers. Only one, center or boundary, may be specified for a 
single plot. center specifies the center of one of the bins. boundary specifies 
the boundary between two bins. Note that if either is above or below the range 
of the data, things will be shifted by the appropriate integer multiple of binwidth. 
For example, to center on integers use binwidth = 1 and center = 0, even if 0 is 
outside the range of the data. Alternatively, this same alignment can be specified 
with binwidth = 1 and boundary = 0.5, even if 0.5 is outside the range of the 
data.

xlab        Label for x-axis. See also gf_labs().

ylab        Label for y-axis. See also gf_labs().
title, subtitle, caption 
Title, sub-title, and caption for the plot. See also gf_labs().

geom, stat  Use to override the default connection between geom_histogram()/geom_freqpoly() 
and stat_bin().

position    Position adjustment, either as a string naming the adjustment (e.g. "jitter" to 
use position_jitter), or the result of a call to a position adjustment function. 
Use the latter if you need to change the settings of the adjustment.

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. It 
can also be a named logical vector to finely select the aesthetics to display.

show.help   If TRUE, display some minimal help.

inherit     A logical indicating whether default attributes are inherited.

environment An environment in which to look for variables not found in data.
Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_freqpoly()

Examples

data(penguins, package = "palmerpenguins")
gf_histogram(~ bill_length_mm | species, alpha = 0.2, data = penguins, bins = 20) %>%
gf_freqpoly(~bill_length_mm, data = penguins, color = ~species, bins = 20)
gf_freqpoly(~bill_length_mm, color = ~species, data = penguins, bins = 20)
gf_dens(~bill_length_mm, data = penguins, color = "navy") %>%
gf_freqpoly(after_stat(density) ~ bill_length_mm, data = penguins, color = "red", bins = 20)

---

gf_function

Layers displaying graphs of functions

Description

These functions provide two different interfaces for creating a layer that contains the graph of a function.

Usage

gf_function(object = NULL, fun, xlim, ..., inherit = FALSE)
gf_fun(object = NULL, formula, xlim, ..., inherit = FALSE)
Arguments

object

When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

fun

A function.

xlim

A numeric vector providing the extent of the x-axis when creating the first layer in a plot. Ignored when creating a subsequent layer.

... Other arguments such as position="dodge".

inherit

A logical indicating whether default attributes are inherited.

formula

A formula describing a function. See examples and mosaicCore::makeFun().

Examples

gf_function(fun = sqrt, xlim = c(0, 10))
gf_dhistogram(~age, data = mosaicData::HELPct, binwidth = 3, alpha = 0.6) %>%
gf_function(
  fun = stats::dnorm,
  args = list(mean = mean(mosaicData::HELPct$age), sd = sd(mosaicData::HELPct$age)),
  color = "red"
)
gf_fun(5 + 3 * cos(10 * x) ~ x, xlim = c(0, 2))
# Utility bill is quadratic in month?
  f <- makeFun(lm(totalbill ~ poly(month, 2), data = mosaicData::Utilities))
gf_point(totalbill ~ month, data = mosaicData::Utilities, alpha = 0.6) %>%
gf_fun(f(m) ~ m, color = "red")

gf_function_2d

Plot functions of two variables

Description

Plot functions of two variables as tile and/or contour plots.

Usage

gf_function_2d(
  object = NULL,
  fun = identity,
  xlim = NULL,
  ylim = NULL,
  ...,
  tile = TRUE,
  contour = TRUE,
  resolution = 50
)

gf_function2d(
gf_function_2d

object = NULL,
fun = identity,
xlim = NULL,
ylim = NULL,
...,
tile = TRUE,
contour = TRUE,
resolution = 50
)

gf_function_contour(
object = NULL,
fun = identity,
xlim = NULL,
ylim = NULL,
...,
resolution = 50
)

gf_function_tile(
object = NULL,
fun = identity,
xlim = NULL,
ylim = NULL,
...,
resolution = 50
)

gf_fun_2d(
object = NULL,
formula = NULL,
xlim = NULL,
ylim = NULL,
tile = TRUE,
contour = TRUE,
...,
resolution = 50
)

gf_fun2d(
object = NULL,
formula = NULL,
xlim = NULL,
ylim = NULL,
tile = TRUE,
contour = TRUE,
...,
resolution = 50
)
Arguments

- **object**: An R object, typically of class "gg".
- **fun**: A function of two variables to be plotted.
- **xlim**: x limits for generating points to be plotted.
- **ylim**: y limits for generating points to be plotted.
- **...**: additional arguments passed to `gf_tile()` or `gf_contour()`.
- **tile**: A logical indicating whether the tile layer should be drawn.
- **contour**: A logical indicating whether the contour layer should be drawn.
- **resolution**: A numeric vector of length 1 or 2 specifying the number of grid points at which the function is evaluated (in each dimension).
- **formula**: A formula describing a function of two variables to be plotted. See `mosaic::makeFun()` for details regarding the conversion from a formula to a function.

Value

A gg plot.

Examples

```r
tHEME_SET(theme_bw())
gf_function_2d(fun = function(x, y) sin(2 * x * y), xlim = c(-pi, pi), ylim = c(-pi, pi)) %>%
gf_refine(scale_fill_viridis_c())
gf_function_2d(fun = function(x, y) x + y, contour = FALSE)
gf_function_tile(fun = function(x, y) x * y) %>%
gf_function_contour(fun = function(x, y) x * y, color = "white") %>%
gf_refine(scale_fill_viridis_c())
```
gf_hex

gf_fun_tile(x*y ~ x + y, xlim = c(-3, 3), ylim = c(-2, 2)) %>%
gf_fun_contour(x*y ~ x + y, color = "white") %>%
gf_refine(scale_fill_viridis_c()) %>%
gf_labs(fill = "product")

---

**Description**

Line plots in ggformula. `gf_path()` differs from `gf_line()` in that points are connected in the order in which they appear in data.

**Usage**

```r
gf_hex(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...
  bins,
  binwidth,
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "hex",
  stat = "binhex",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)
```

**Arguments**

- **object**: When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

- **gformula**: A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data  The data to be displayed in this layer. There are three options:
   If NULL, the default, the data is inherited from the plot data as specified in the call to `ggplot()`.
   A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.
   A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. `~ head(.x, 10)`).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with `attribute = value`, (b) ggplot2 aesthetics to be mapped with `attribute = ~ expression`, or (c) attributes of the layer as a whole, which are set with `attribute = value`.

bins  numeric vector giving number of bins in both vertical and horizontal directions. Set to 30 by default.

binwidth  Numeric vector giving bin width in both vertical and horizontal directions. Overrides bins if both set.

alpha  Opacity (0 = invisible, 1 = opaque).

color  A color or a formula used for mapping color.

fill  A color for filling, or a formula used for mapping fill.

group  Used for grouping.

linetype  A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth  A numerical line width or a formula used for mapping linewidth.

xlab  Label for x-axis. See also `gf_labs()`.

ylab  Label for y-axis. See also `gf_labs()`.

title, subtitle, caption  Title, sub-title, and caption for the plot. See also `gf_labs()`.

geom, stat  Override the default connection between `geom_hex()` and `stat_binhex()`.

position  Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help  If TRUE, display some minimal help.

inherit  A logical indicating whether default attributes are inherited.

environment  An environment in which to look for variables not found in data.

Value

a gg object
Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_hex()

Examples

```r
gf_hex(avg_drinks ~ age, data = mosaicData::HELPrct, bins = 15) %>%
gf_density2d(avg_drinks ~ age, data = mosaicData::HELPrct, color = "red", alpha = 0.5)
```

gf_histogram

Formula interface to geom_histogram()

Description

Count and density histograms in ggformula.

Usage

```r
gf_histogram(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  bins = 25,
  binwidth,
  alpha = 0.5,
  color,
  fill,
  group,
  linetype,
  linewidth,
  xlab,
```
gf_histogram

ylab,
title,
subtitle,
caption,
geom = "bar",
stat = "bin",
position = "stack",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_dhistogram(
  object = NULL,
gformula = NULL,
data = NULL,
...

  bins = 25,
  binwidth,
  alpha = 0.5,
  color,
  fill,
  group,
  linetype,
  linewidth,
  xlab,
ylab,
title,
subtitle,
caption,
  geom = "bar",
  stat = "bin",
  position = "stack",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_histogramh(
    object = NULL,
gformula = NULL,
data = NULL,
...

  bins = 25,
  binwidth,
  alpha = 0.5,
gf_histogram

color,
fill,
group,
linetype,
linewidth,
xlab,
ylab,
title,
subtitle,
caption,
geom = "barh",
stat = "binh",
position = "stackv",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_dhistogramh(
object = NULL,
gformula = NULL,
data = NULL,
...
)

bins = 25,
binwidth,
alpha = 0.5,
color,
fill,
group,
linetype,
linewidth,
xlab,
ylab,
title,
subtitle,
caption,
geom = "barh",
stat = "binh",
position = "stackv",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape \texttt{x ~ y} (or \texttt{y ~ x}, but this shape is not generally needed).

data
The data to be displayed in this layer. There are three options:
- If \texttt{NULL}, the default, the data is inherited from the plot data as specified in the call to \texttt{ggplot()}. A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See \texttt{fortify()} for which variables will be created.
- A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. \texttt{~ head(x, 10)}).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = \texttt{~ expression}, or (c) attributes of the layer as a whole, which are set with attribute = value.

bins
Number of bins. Overridden by binwidth. Defaults to 30.

binwidth
The width of the bins. Can be specified as a numeric value or as a function that calculates width from unscaled x. Here, "unscaled x" refers to the original x values in the data, before application of any scale transformation. When specifying a function along with a grouping structure, the function will be called once per group. The default is to use the number of bins in bins, covering the range of the data. You should always override this value, exploring multiple widths to find the best to illustrate the stories in your data.

The bin width of a date variable is the number of days in each time; the bin width of a time variable is the number of seconds.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

fill
A color for filling, or a formula used for mapping fill.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

xlab
Label for x-axis. See also \texttt{gf_labs()}.

ylab
Label for y-axis. See also \texttt{gf_labs()}.

title, subtitle, caption
Title, sub-title, and caption for the plot. See also \texttt{gf_labs()}.

geom, stat
Use to override the default connection between \texttt{geom_histogram()}/\texttt{geom_freqpoly()} and \texttt{stat_bin()}.

position
Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
gf_histogram

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help  If TRUE, display some minimal help.

inherit  A logical indicating whether default attributes are inherited.

environment  An environment in which to look for variables not found in data.

Value

a gg object

Horizontal Geoms

There are two ways to obtain "horizontal" geoms: (1) The ggstance package provides a set of "horizontal" geoms and positions; (2) Thee ggplot2 now provides an orientation argument for "native" horizontal geoms and positions. ggformula supports both.

Specifying plot attributes

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In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_histogram()

Examples

x <- rnorm(1000)
gf_histogram(~x, bins = 30)
gf_dhistogram(~x, bins = 30)
gf_dhistogram(~x, binwidth = 0.5, center = 0, color = "black")
gf_dhistogram(~x, binwidth = 0.5, boundary = 0, color = "black")
gf_dhistogram(~x, bins = 30) %>%
gf_fitdistr(dist = "dnorm") # see help for gf_fitdistr() for more info.

gf_histogram(~x, fill = ~ (abs(x) <= 2), boundary = 2, binwidth = 0.25)
data(penguins, package = "palmerpenguins")
gf_histogram(~ bill_length_mm | species, data = penguins, binwidth = 0.25)
gf_histogram(~ age,
  data = mosaicData::HELPrct, binwidth = 5,
  fill = "skyblue", color = "black"
)
# bins can be adjusted left/right using center or boundary
gf_histogram(~ age,
  data = mosaicData::HELPrct,
  binwidth = 5, fill = "skyblue", color = "black", center = 42.5
)
gf_histogram(~ age,
  data = mosaicData::HELPrct,
  binwidth = 5, fill = "skyblue", color = "black", boundary = 40
)
gf_histogram(~ age,
  data = mosaicData::HELPrct,
  binwidth = 5, fill = "skyblue", color = "black", boundary = 40
)
gf_histogramh(~ x, bins = 30)
gf_histogram(~ ., bins = 30)
gf_histogramh(~ ., bins = 30)
gf_histogramh(~ after_stat(density), bins = 30)
gf_dhistogramh(~ x, bins = 30)
gf_dhistogram(~ ., bins = 30)
gf_dhistogramh(~ ., bins = 30)

---

gf_jitter

Formula interface to geom_jitter()

Description

Jittered scatter plots in ggformula.

Usage

gf_jitter(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  size,
  shape,
  fill,
  width,
  height,
Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
size A numeric size or a formula used for mapping size.
shape An integer or letter shape or a formula used for mapping shape.
fill A color for filling, or a formula used for mapping fill.
width Amount of horizontal jitter.
height Amount of vertical jitter.
group Used for grouping.
stroke A numeric size of the border or a formula used to map stroke.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom A character string naming the geom used to make the layer.
stat A character string naming the stat used to make the layer.
position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend  A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help    If TRUE, display some minimal help.

inherit      A logical indicating whether default attributes are inherited.

evironment   An environment in which to look for variables not found in data.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_jitter(), gf_point()

Examples

gf_jitter()
# without jitter
gf_point(age ~ sex, alpha = 0.25, data = mosaicData::HELPct)
# jitter only horizontally
gf_jitter(age ~ sex, alpha = 0.25, data = mosaicData::HELPct, width = 0.2, height = 0)
# alternative way to get jitter
gf_point(age ~ sex,
         alpha = 0.25, data = mosaicData::HELPct,
         position = "jitter", width = 0.2, height = 0
)
gf_labs

Non-layer functions for gf plots

Description

These functions modify things like labels, limits, scales, etc. for plots ggplot2 plots. They are wrappers around functions in ggplot2 that allow for chaining syntax.

Usage

gf_labs(object, ...)
gf_lims(object, ...)
gf_refine(object, ...)

Arguments

object a gg object
...
additional arguments passed through to the similarly named function in ggplot2.

Details

gf_refine() provides a mechanism to replace + with the chaining operator from magrittr. Each of its \dots arguments is added in turn to the base plot in object. The other functions are thin wrappers around specific ggplot2 refinement functions and pass their \dots arguments through to the similarly named ggplot2 functions.

Value

a modified gg object

Examples

gf_dens(~cesd, color = ~substance, linewidth = 1.5, data = mosaicData::HELPrct) %>%
gf_labs(
  title = "Center for Epidemiologic Studies Depression measure",
  subtitle = "(at baseline)",
  color = "Abused substance: ",
  x = "CESD score",
  y = "",
  caption = "Source: HELPrct"
) %>%
gf_theme(theme_classic()) %>%
gf_theme(
  axis.text.y = element_blank(),
  legend.position = "top",
  plot.title = element_text(hjust = 0.5, color = "navy"),
)
gf_point(eruptions ~ waiting, data = faithful, alpha = 0.5)
gf_point(eruptions ~ waiting, data = faithful, alpha = 0.5) %>%
gf_lims(x = c(65, NA), y = c(3, NA))

# modify scales using gf_refine()
data(penguins, package = "palmerpenguins")
gf_jitter(bill_length_mm ~ bill_depth_mm, color = ~species, data = penguins) %>%
gf_refine(scale_color_brewer(type = "qual", palette = 3)) %>%
gf_theme(theme_bw())

gf_jitter(bill_length_mm ~ bill_depth_mm, color = ~species, data = penguins) %>%
gf_refine(scale_color_manual(values = c("red", "navy", "limegreen"))) %>%
gf_theme(theme_bw())

---

**gf_line**  
*Formula interface to geom_line() and geom_path()*

**Description**

Line plots in ggformula. `gf_path()` differs from `gf_line()` in that points are connected in the order in which they appear in data.

**Usage**

```r
gf_line(
  object = NULL,  
gformula = NULL,  
data = NULL,  
...,  
alpha,  
color,  
fill,  
group,  
linetype,  
linewidth,  
lineend,  
linejoin,  
linemitre,  
arrow,  
xlab,  
ylab,  
title,  
textsubtitle,  
textcaption,  
geom = "line",  
)`
```
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_path(
object = NULL,
gformula = NULL,
data = NULL,
..., alpha, color, group, linetype, linewidth, lineend = "butt", linejoin = "round", linemitre = 1, arrow = NULL, xlab, ylab, title, subtitle, caption, geom = "path", stat = "identity", position = "identity", show.legend = NA, show.help = NULL, inherit = TRUE, environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
lineend Line end style (round, butt, square).
linejoin Line join style (round, mitre, bevel).
linemitre Line mitre limit (number greater than 1).
arrow Arrow specification, as created by grid::arrow().
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom A character string naming the geom used to make the layer.
stat A character string naming the stat used to make the layer.
position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.
show.help If TRUE, display some minimal help.
inherit A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.

Value
a gg object

Specifying plot attributes
Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation
Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.
gf_linerange

See Also

ggplot2::geom_line(), gf_point()

Examples

gf_line()
gf_point(age ~ sex, alpha = 0.25, data = mosaicData::HELPrct)
gf_point(births ~ date, color = ~wday, data = mosaicData::Births78)
# lines make the exceptions stand out more prominently
gf_line(births ~ date, color = ~wday, data = mosaicData::Births78)
gf_path()
if (require(dplyr)) {
  data.frame(t = seq(1, 10 * pi, length.out = 400)) %>%
    mutate(x = t * cos(t), y = t * sin(t)) %>%
    gf_path(y ~ x, color = ~t)
}

---

gf_linerange  

Formula interface to geom_linerange() and geom_pointrange()

Description

Various ways of representing a vertical interval defined by x, ymin and ymax. Each case draws a single graphical object.

Usage

gf_linerange(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,  
  alpha,  
  color,  
  group,  
  linetype,  
  linewidth,  
  xlab,  
  ylab,  
  title,  
  subtitle,  
  caption,  
  geom = "linerange",  
  stat = "identity",  
  position = "identity",  
  show.legend = NA,  
  show.help = NULL,  
  inherit = TRUE,
gf_linerange

environment = parent.frame()
)
gf_pointrange(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ....,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  size,
  fatten = 2,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "pointrange",
  stat = "identity",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)
gf_summary(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ....,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  size,
  fun.y = NULL,
  fun.ymax = NULL,
  fun.ymin = NULL,
  fun.args = list(),
  fatten = 2,
  xlab,
  ylab,
  title,
gf_linerange

subtitle,
caption,
geom = "pointrange",
stat = "summary",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_linerangeh(
object = NULL,
gformula = NULL,
data = NULL,
...,
alpha,
color,
group,
linetype,
linewidth,
xlab,
ylab,
title,
subtitle,
caption,
geom = "linerangeh",
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_pointrangeh(
object = NULL,
gformula = NULL,
data = NULL,
...,
alpha,
color,
group,
linetype,
linewidth,
xlab,
ylab,
title,
subtitle, caption, geom = "pointrangeh", stat = "identity", position = "identity", show.legend = NA, show.help = NULL, inherit = TRUE, environment = parent.frame()

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape ymin + ymax ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
   If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot()
   A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
   A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption
   Title, sub-title, and caption for the plot. See also gf_labs().
geom The geometric object to use to display the data, either as a ggproto Geom subclass or as a string naming the geom stripped of the geom_prefix (e.g. "point" rather than "geom_point")
stat
The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")

position
Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.

size
size aesthetic for points (gf_pointrange()).

fatten
A multiplicative factor used to increase the size of the middle bar in geom_crossbar() and the middle point in geom_pointrange().

fun.ymin, fun.y, fun.ymax
[Deprecated] Use the versions specified above instead.

fun.args
Optional additional arguments passed on to the functions.

See Also
ggplot2::geom_linerange()
ggplot2::geom_pointrange()
ggplot2::geom_pointrange(), ggplot2::stat_summary()

Examples
gf_linerange()
gf_ribbon(low_temp + high_temp ~ date, 
  data = mosaicData::Weather, 
  fill = ~city, alpha = 0.4 
) %>%
gf_theme(theme = theme_minimal())
gf_linerange(
  low_temp + high_temp ~ date | city ~ ., 
  data = mosaicData::Weather, 
  color = ~ ((low_temp + high_temp) / 2) 
) %>%
gf_refine(scale_colour_gradientn(colors = rev(rainbow(5)))) %>%
gflabs(color = "mid-temp")
gf_ribbon(low_temp + high_temp ~ date | city ~ ., data = mosaicData::Weather)

# Chaining in the data
mosaicData::Weather %>%
gf_ribbon(low_temp + high_temp ~ date, alpha = 0.4) %>%
if (require(mosaicData) && require(dplyr)) {
  HELP2 <- HELPrct %>%
    group_by(substance, sex) %>%
    summarise(
      age = NA,
      mean.age = mean(age),
      median.age = median(age),
      max.age = max(age),
      min.age = min(age),
      sd.age = sd(age),
      lo = mean.age - sd.age,
      hi = mean.age + sd.age
    )
  gf_jitter(age ~ substance, data = HELPrct, alpha = 0.5, width = 0.2, height = 0, color = "skyblue") %>%
    gf_pointrange(mean.age + lo + hi ~ substance, data = HELP2) %>%
    gf_facet_grid(~sex)
  gf_jitter(age ~ substance, data = HELPrct, alpha = 0.5, width = 0.2, height = 0, color = "skyblue") %>%
    gf_errorbar(lo + hi ~ substance, data = HELP2, inherit = FALSE) %>%
    gf_facet_grid(~sex)
}

p <- gf_jitter(mpg ~ cyl, data = mtcars, height = 0, width = 0.15); p
p %>% gf_summary(fun.data = "mean_cl_boot", color = "red", size = 2, linewidth = 1.3)
# You can supply individual functions to summarise the value at # each x:
p %>% gf_summary(fun.y = "median", color = "red", size = 3, geom = "point")
p %>%
  gf_summary(fun.y = "mean", color = "red", size = 3, geom = "point") %>%
  gf_summary(fun.y = mean, geom = "line")
p %>%
  gf_summary(fun.y = mean, fun.ymin = min, fun.ymax = max, color = "red")
## Not run:
p %>%
  gf_summary(fun.ymin = min, fun.ymax = max, color = "red", geom = "linerange")
## End(Not run)
gf_bar(~ cut, data = diamonds)
gf_col(price ~ cut, data = diamonds, stat = "summary_bin", fun.y = "mean")
# Don't use gf_lims() to zoom into a summary plot - this throws the data away
p <- gf_summary(mpg ~ cyl, data = mtcars, fun.y = "mean", geom = "point")
p
p %>% gf_lims(y = c(15, 30))
# Instead use coord_cartesian()
p %>% gf_refine(coord_cartesian(ylim = c(15, 30)))
# A set of useful summary functions is provided from the Hmisc package.
## Not run:
p <- gf_jitter(mpg ~ cyl, data = mtcars, width = 0.15, height = 0); p
p %>% gf_summary(fun.data = mean_cl_boot, color = "red")
p %>% gf_summary(fun.data = mean_cl_boot, color = "red", geom = "crossbar")
p %>% gf_summary(fun.data = mean_sdl, group = ~ cyl, color = "red", geom = "crossbar", width = 0.3)
p %>% gf_summary(group = ~ cyl, color = "red", geom = "crossbar", width = 0.3, fun.data = mean_sdl, fun.args = list(mult = 1))
p %>% gf_summary(fun.data = median_hilow, group = ~ cyl, color = "red", geom = "crossbar", width = 0.3)
## End(Not run)

# An example with highly skewed distributions:
if (require("ggplot2movies")) {
  set.seed(596)
  Mov <- movies[sample(nrow(movies), 1000), ]
  m2 <- gf_jitter(votes ~ factor(round(rating)), data = Mov, width = 0.15, height = 0, alpha = 0.3)
  m2 <- m2 %>%
    gf_summary(fun.data = "mean_cl_boot", geom = "crossbar",
               colour = "red", width = 0.3) %>%
    gf_labs(x = "rating")

  m2
  # Notice how the overplotting skews off visual perception of the mean
  # supplementing the raw data with summary statistics is _very_ important

  # Next, we'll look at votes on a log scale.

  # Transforming the scale means the data are transformed
  # first, after which statistics are computed:
  m2 %>% gf_refine(scale_y_log10())
  # Transforming the coordinate system occurs after the statistic has been computed. This means we're calculating the summary on the raw data
  # and stretching the geoms onto the log scale. Compare the widths of the standard errors.
  m2 %>% gf_refine(coord_trans(y="log10"))
}
## Not run:
gf_linerangeh(date ~ low_temp + high_temp | ~city, data = mosaicData::Weather,
               color = ~avg_temp)
%>%
gf_refine(scale_color_viridis_c(begin = 0.1, end = 0.9, option = "C"))
gf_linerange(date ~ low_temp + high_temp | ~city, data = mosaicData::Weather,
gf_plot

Formula interface to ggplot()

decription
Create a new ggplot and (optionally) set default dataset aesthetics mapping.

Usage

gf_plot(...)

Arguments

... arguments that can include data (a data frame or something that can be ggplot2::fortify()ed to become one) and aesthetics specified using the following formula notation: aesthetic = ~ expression. See examples.

Value

a gg object

Examples

gf_plot(mtcars, x = ~ wt, y = ~ mpg, color = ~ factor(cyl)) %>%
gf_density_2d() %>%
gf_point()
gf_point

Formula interface to geom_point()

Description

Scatterplots in ggformula.

Usage

gf_point(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  size,
  shape,
  fill,
  group,
  stroke,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "point",
  stat = "identity",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data A data frame with the variables to be plotted.
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, (c) attributes of the layer as a whole, which are set with attribute = value, or (d) arguments for the geom, stat, or position function.
alpha   Opacity (0 = invisible, 1 = opaque).
color   A color or a formula used for mapping color.
size    A numeric size or a formula used for mapping size.
shape   An integer or letter shape or a formula used for mapping shape.
fill    A color for filling, or a formula used for mapping fill.
group   Used for grouping.
stroke  A numeric size of the border or a formula used to map stroke.
xlab    Label for x-axis. See also \texttt{gf_labs()}.
ylab    Label for y-axis. See also \texttt{gf_labs()}.
title, subtitle, caption
        Title, sub-title, and caption for the plot. See also \texttt{gf_labs()}.
geom    A character string naming the geom used to make the layer.
stat    A character string naming the stat used to make the layer.
position Either a character string naming the position function used for the layer or a
            position object returned from a call to a position function.
show.legend A logical indicating whether this layer should be included in the legends. \texttt{NA},
            the default, includes layer in the legends if any of the attributes of the layer are
            mapped.
show.help If \texttt{TRUE}, display some minimal help.
inherit  A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.

Value

a \texttt{gg} object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in \texttt{gformula}. Setting and
mapping of additional attributes can be done through the use of additional arguments. Attributes
can be set can be set using arguments of the form \texttt{attribute = value} or mapped using arguments
of the form \texttt{attribute = \sim expression}.

In formulas of the form \texttt{A | B, B} will be used to form facets using \texttt{facet_wrap()} or \texttt{facet_grid()}. This
provides an alternative to \texttt{gf_facet_wrap()} and \texttt{gf_facet_grid()} that is terser and may feel
more familiar to users of \texttt{lattice}.

Evaluation

Evaluation of the \texttt{ggplot2} code occurs in the environment of \texttt{gformula}. This will typically do
the right thing when formulas are created on the fly, but might not be the right thing if formulas created
in one environment are used to create plots in another.

See Also

\texttt{ggplot2::geom_point(), gf_line(), gf_jitter()}
Examples

gf_point()
gf_point((10 * ((1:25) %% 10)) ~ ((1:25) %% 10),
    shape = 1:25,
    fill = "skyblue", color = "navy", size = 4, stroke = 1, data = NA
)  
gf_point(mpg ~ hp, color = ~cyl, size = ~wt, data = mtcars)
  # faceting -- two ways
  gf_point(mpg ~ hp, data = mtcars) %>%
    gf_facet_wrap(~am)
  gf_point(mpg ~ hp | am, group = ~cyl, data = mtcars)
  gf_point(mpg ~ hp | ~am, group = ~cyl, data = mtcars)
  gf_point(mpg ~ hp | am ~ ., group = ~cyl, data = mtcars)
  # Chaining in the data
  mtcars %>% gf_point(mpg ~ wt)

  # short cuts for main labels in the plot
  gf_point(births ~ date,
    color = ~wday, data = mosaicData::Births78,
    xlab = "Date", ylab = "Number of Live Births",
    title = "Interesting Patterns in the Number of Births",
    subtitle = "(United States, 1978)",
    caption = "Source: mosaicData::Births78"
  )

---

gf_polygon

GF_Polygon

 Formula Interface to geom_polygon()

Description

Line plots in ggformula. gf_path() differs from gf_line() in that points are connected in the order in which they appear in data.

Usage

gf_polygon(
    object = NULL,
    gformula = NULL,
    data = NULL,
    ..., alpha, color, linewidth, shape, fill, group, stroke, xlab,
Arguments

object  When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula  A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data  A data frame with the variables to be plotted.
...  Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha  Opacity (0 = invisible, 1 = opaque).
color  A color or a formula used for mapping color.
linewidth  A numerical line width or a formula used for mapping linewidth.
shape, stroke  Aesthetics for polygons.
fill  A color for filling, or a formula used for mapping fill.
group  Used for grouping.
xlab  Label for x-axis. See also gf_labs().
ylab  Label for y-axis. See also gf_labs().
title, subtitle, caption  Title, sub-title, and caption for the plot. See also gf_labs().
geom  A character string naming the geom used to make the layer.
stat  A character string naming the stat used to make the layer.
position  Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend  A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.
show.help  If TRUE, display some minimal help.
inherit  A logical indicating whether default attributes are inherited.
environment  An environment in which to look for variables not found in data.
**gf_polygon**

**Value**

a gg object

**Specifying plot attributes**

Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

**Evaluation**

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

**See Also**

`ggplot2::geom_line()`, `gf_point()`

**Examples**

```r
gf_polygon()
if (require(maps) && require(ggthemes) && require(dplyr)) {
  US <- map_data("state") %>%
    dplyr::mutate(name_length = nchar(region))
  States <- US %>%
    dplyr::group_by(region) %>%
    dplyr::summarise(lat = mean(range(lat)), long = mean(range(long))) %>%
    dplyr::mutate(name = abbreviate(region, 3))
  gf_polygon(lat ~ long,
            data = US, group = ~group,
            fill = ~name_length, color = "white"
          ) %>%
  gf_text(lat ~ long,
          label = ~name, data = States,
          color = "gray70", inherit = FALSE
        ) %>%
  gf_refine(ggthemes::theme_map())
}
```
gf_qq

Formula interface to geom_qq()

Description

gf_qq() and gf_qqstep() both create quantile-quantile plots. They differ in how they display the qq-plot. gf_qq() uses points and gf_qqstep() plots a step function through these points.

Usage

gf_qq(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  group,
  distribution = stats::qnorm,
  dparams = list(),
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "point",
  stat = "qq",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_qqline(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  group,
  distribution = stats::qnorm,
  dparams = list(),
  linetype = "dashed",
  alpha = 0.7,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
Arguments

object

When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula

A formula with shape ~ sample. Facets can be added using |

data

The data to be displayed in this layer. There are three options:

If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().

A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

...

Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute =
~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

- group
  - Used for grouping.
- distribution
  - Distribution function to use, if x not specified
- dparams
  - Additional parameters passed on to distribution function.
- xlab
  - Label for x-axis. See also gf_labs().
- ylab
  - Label for y-axis. See also gf_labs().
- title, subtitle, caption
  - Title, sub-title, and caption for the plot. See also gf_labs().
- geom, stat
  - Use to override the default connection between geom_histogram()/geom_freqpoly() and stat_bin().
- position
  - Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
- 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. It can also be a named logical vector to finely select the aesthetics to display.
- show.help
  - If TRUE, display some minimal help.
- inherit
  - A logical indicating whether default attributes are inherited.
- environment
  - An environment in which to look for variables not found in data.
- linetype
  - A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
- alpha
  - Opacity (0 = invisible, 1 = opaque).

Value

- a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.
gf_quantile

See Also
ggplot2::geom_qq()

Examples

gf_qq(~ rnorm(100))
data(penguins, package = "palmerpenguins")
gf_qq(~ bill_length_mm | species, data = penguins) %>% gf_qqline()
gf_qq(~ bill_length_mm | species, data = penguins) %>% gf_qqline(tail = 0.10)
gf_qq(~bill_length_mm, color = ~species, data = penguins) %>%
  gf_qqstep(~bill_length_mm, color = ~species, data = penguins)

gf_quantile

Formula interface to geom_quantile()

Description

This fits a quantile regression to the data and draws the fitted quantiles with lines. This is as a
continuous analogue to geom_boxplot().

Usage

gf_quantile(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  lineend = "butt",
  linejoin = "round",
  linemitre = 1,
  quantiles,
  formula,
  method,
  method.args,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "quantile",
  stat = "quantile",
  position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape \( y \sim x \). Faceting can be achieved by including | in the formula.

data
The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to \texttt{ggplot()}. A \texttt{data.frame}, or other object, will override the plot data. All objects will be fortified to produce a data frame. See \texttt{fortify()} for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a \texttt{data.frame}, and will be used as the layer data. A function can be created from a \texttt{formula} (e.g. \texttt{~ head(.x, 10)}).

...
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with \texttt{attribute = value}, (b) ggplot2 aesthetics to be mapped with \texttt{attribute = \texttt{~ expression}}, or (c) attributes of the layer as a whole, which are set with \texttt{attribute = value}.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.
group
Used for grouping.
linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth
A numerical line width or a formula used for mapping linewidth.
lineend
Line end style (round, butt, square).
linejoin
Line join style (round, mitre, bevel).
linemitre
Line mitre limit (number greater than 1).
quantiles
conditional quantiles of \( y \) to calculate and display
formula
formula relating \( y \) variables to \( x \) variables
method
Quantile regression method to use. Available options are "rq" (for \texttt{quantreg::rq()}) and "rqss" (for \texttt{quantreg::rqss()}).
method.args
List of additional arguments passed on to the modelling function defined by method.
xlab
Label for x-axis. See also \texttt{gf_labs()}. ylab
Label for y-axis. See also \texttt{gf_labs()}.
title, subtitle, caption
Title, sub-title, and caption for the plot. See also `gf_labs()`.

geom, stat
Use to override the default connection between `geom_quantile()` and `stat_quantile()`.

position
Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.

Value
a gg object

Specifying plot attributes
Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, B will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation
Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also
`ggplot2::geom_quantile()`

Examples
```r
gf_point((1 / hwy) ~ displ, data = mpg) %>%
gf_quantile((1 / hwy) ~ displ)
```
gf_raster  

Formula interface to geom_raster()

Description

Formula interface to geom_raster()

Usage

gf_raster(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha, 
  color, 
  fill, 
  group, 
  linetype, 
  linewidth, 
  hjust = 0.5, 
  vjust = 0.5, 
  interpolate = FALSE, 
  xlab, 
  ylab, 
  title, 
  subtitle, 
  caption, 
  geom = "raster", 
  stat = "identity", 
  position = "identity", 
  show.legend = NA, 
  show.help = NULL, 
  inherit = TRUE, 
  environment = parent.frame()
)

Arguments

object  When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula A formula with shape y ~ x or fill ~ x + y

data A data frame with the variables to be plotted.

...  Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
**gf_raster**

alpha Opacity (0 = invisible, 1 = opaque).

color A color or a formula used for mapping color.

fill A color for filling, or a formula used for mapping fill.

group Used for grouping.

linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth A numerical line width or a formula used for mapping linewidth.

hjust, vjust horizontal and vertical justification of the grob. Each justification value should be a number between 0 and 1. Defaults to 0.5 for both, centering each pixel over its data location.

interpolate If TRUE interpolate linearly, if FALSE (the default) don’t interpolate.

xlab Label for x-axis. See also gf_labs().

ylab Label for y-axis. See also gf_labs().

title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().

geom A character string naming the geom used to make the layer.

stat A character string naming the stat used to make the layer.

position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help If TRUE, display some minimal help.

inherit A logical indicating whether default attributes are inherited.

environment An environment in which to look for variables not found in data.

**Value**

a gg object

**Specifying plot attributes**

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, B will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

**Evaluation**

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.
See Also

ggplot2::geom_raster()

Examples

# Justification controls where the cells are anchored
D <- expand.grid(x = 0:5, y = 0:5)
D$z <- runif(nrow(D))
# centered squares
gf_raster(z ~ x + y, data = D)
gf_raster(y ~ x, fill = ~z, data = D)
# zero padding
gf_raster(z ~ x + y, data = D, hjust = 0, vjust = 0)
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape \( y_{\text{min}} + y_{\text{max}} \sim x_{\text{min}} + x_{\text{max}} \). Faceting can be achieved by including \( | \) in the formula.

data
A data frame with the variables to be plotted.

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

fill
A color for filling, or a formula used for mapping fill.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

xlab
Label for x-axis. See also gf_labs().

ylab
Label for y-axis. See also gf_labs().
title, subtitle, caption
Title, sub-title, and caption for the plot. See also gf_labs().

geom
A character string naming the geom used to make the layer.

stat
A character string naming the stat used to make the layer.

position
Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend
A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form \( A \ | \ B, B \) will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.
Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_rect()`

Examples

```r
gf_rect(1 + 2 ~ 3 + 4, alpha = 0.3, color = "red")
# use data = data.frame() so we get 1 rectangle and not 1 per row of faithful
# use inherit = FALSE because we are not reusing eruptions and waiting
gf_point(eruptions ~ waiting, data = faithful) %>%
gf_rect(1.5 + 3 ~ 45 + 68,
  fill = "red", alpha = 0.2,
  data = data.frame(), inherit = FALSE) %>%
gf_rect(3 + 5.5 ~ 68 + 100,
  fill = "green", alpha = 0.2,
  data = data.frame(), inherit = FALSE)
```

gf_relabel

Modify plot labeling

Description

Some packages like `expss` provide mechanisms for providing longer labels to R objects. These labels can be used when labeling plots and tables, for example, without requiring long or awkward variable names. This is an experimental feature and currently only supports `expss` or any other system that stores a label in the `label` attribute of a vector.

Usage

```r
gf_relabel(plot, labels = get_variable_labels(plot$data), ...)
```

## S3 method for class 'gf_ggplot'

`print(x, labels = get_variable_labels(x$data), ...)`

Arguments

- `plot` A `ggplot`.
- `labels` A named list of labels.
- `...` Additional named labels. See examples.
- `x` A `ggplot`.
**gf_ribbon**

*Value*

A plot with potentially modified labels.

**Examples**

```r
# labeling using a list
labels <- list(width = "width of foot (cm)", length = "length of foot (cm)",
               domhand = "dominant hand")
gf_point(length ~ width, color = ~domhand, data = mosaicData::KidsFeet) %>%
gf_relabel(labels)

# labeling using ...
gf_point(length ~ width, color = ~domhand, data = mosaicData::KidsFeet) %>%
gf_relabel(
  width = "width of foot (cm)",
  length = "length of foot (cm)",
  domhand = "dominant hand")

# Alternatively, we can store labels with data.
KF <- mosaicData::KidsFeet %>%
  set_variable_labels(
    length = 'foot length (cm)',
    width = 'foot width (cm')
  )
gf_point(length ~ width, data = KF)
gf_density2d(length ~ width, data = KF)
get_variable_labels(KF)
```

---

**gf_ribbon**

*Formula interface to geom_ribbon()*

**Description**

For each x value, `geom_ribbon()` displays a y interval defined by `ymin` and `ymax`. `geom_area()` is a special case of `geom_ribbon()`, where the `ymin` is fixed to 0 and `y` is used instead of `ymax`.

**Usage**

```r
gf_ribbon(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha = 0.3,
  xlab,
  ylab,
```
title,
subtitle,
caption,
geom = "ribbon",
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape ymin + ymax ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
...
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
alpha Opacity (0 = invisible, 1 = opaque).
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom The geometric object to use to display the data, either as a ggproto Geom subclass or as a string naming the geom stripped of the geom_ prefix (e.g. "point" rather than "geom_point")
stat The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")
position Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
gf_ridgeline

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help If TRUE, display some minimal help.

inherit A logical indicating whether default attributes are inherited.

environment An environment in which to look for variables not found in data.

See Also

ggplot2::geom_ribbon()

Examples

gf_ribbon()

gf_ribbon(low_temp + high_temp ~ date, data = mosaicData::Weather, fill = ~city, alpha = 0.4) %>%
gf_theme(theme = theme_minimal())
gf_linerange(
  low_temp + high_temp ~ date | city ~ .,
  color = ~high_temp,
  data = mosaicData::Weather
) %>%
gf_refine(scale_colour_gradientn(colors = rev(rainbow(5))))
gf_ribbon(low_temp + high_temp ~ date | city ~ ., data = mosaicData::Weather)
# Chaining in the data
## Not run:
mosaicData::Weather %>%
gf_ribbon(low_temp + high_temp ~ date, alpha = 0.4) %>%
gf_facet_grid(city ~ .)

## End(Not run)

---

gf_ridgeline

Formula interface to ggridges plots

Description

Formula interface to ggridges plots

Usage

gf_ridgeline(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  height,
  scale = 1,
)
min_height = 0,
color,
fill,
alpha,
group,
linetype,
linewidth,
point_size,
point_shape,
point_colour,
point_fill,
point_alpha,
point_stroke,
xlab,
ylab,
title,
subtitle,
caption,
geom = "ridgeline",
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_density_ridges(
  object = NULL,
gformula = NULL,
data = NULL,
...,
height,
scale = 1,
rel_min_height = 0,
color,
fill,
alpha,
group,
linetype,
linewidth,
point_size,
point_shape,
point_colour,
point_fill,
point_alpha,
point_stroke,
panel_scaling = TRUE,
gf_ridgeline

xlab,
ylab,
title,
subtitle,
caption,
geom = "density_ridges",
stat = "density_ridges",
position = "points_sina",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)
gf_density_ridges2(
  object = NULL,
gformula = NULL,
data = NULL,
..., 
height,
scale = 1,
rel_min_height = 0,
color,
fill,
alpha,
group,
linetype,
linewidth,
point_size,
point_shape,
point_colour,
point_fill,
point_alpha,
point_stroke,
panel_scaling = TRUE,
xlab,
ylab,
title,
subtitle,
caption,
geom = "density_ridges2",
stat = "density_ridges",
position = "points_sina",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)
gf_density_ridgeline_gradient(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  height,
  color,
  fill,
  alpha,
  group,
  linetype,
  linewidth,
  gradient_lwd = 0.5,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "ridgeline_gradient",
  stat = "identity",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_density_ridges_gradient(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  height,
  panel_scaling = TRUE,
  color,
  fill = ~stat(x),
  alpha,
  group,
  linetype,
  linewidth,
  gradient_lwd = 0.5,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "density_ridges_gradient",
  ...)
stat = "density_ridges",
position = "points_sina",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options: If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
height The height of each ridgeline at the respective x value. Automatically calculated and provided by ggridges::stat_density_ridges() if the default stat is not changed.
scale A scaling factor to scale the height of the ridgelines relative to the spacing between them. A value of 1 indicates that the maximum point of any ridgeline touches the baseline right above, assuming even spacing between baselines.
min_height A height cutoff on the drawn ridgelines. All values that fall below this cutoff will be removed. The main purpose of this cutoff is to remove long tails right at the baseline level, but other uses are possible. The cutoff is applied before any height scaling is applied via the scale aesthetic. Default is 0, so negative values are removed.
color A color or a formula used for mapping color.
fill A color for filling, or a formula used for mapping fill.
alpha Opacity (0 = invisible, 1 = opaque).
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
l linewidth A numerical line width or a formula used for mapping linewidth.
gf_ridgeline

point_shape, point_colour, point_size, point_fill, point_alpha, point_stroke

As in `ggridges::geom_ridgeline()`.

xlab Label for x-axis. See also `gf_labs()`.

ylab Label for y-axis. See also `gf_labs()`.

title, subtitle, caption

Title, sub-title, and caption for the plot. See also `gf_labs()`.

geom, stat Use to override the default connection between `geom_density()` and `stat_density()`.

position Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help If TRUE, display some minimal help.

inherit A logical indicating whether default attributes are inherited.

environment An environment in which to look for variables not found in `data`.

rel_min_height Lines with heights below this cutoff will be removed. The cutoff is measured relative to the overall maximum, so `rel_min_height = 0.01` would remove everything. Default is 0, so nothing is removed.

panel_scaling If TRUE, the default, relative scaling is calculated separately for each panel. If FALSE, relative scaling is calculated globally.

gradient_lwd A parameter to needed to remove rendering artifacts inside the rendered gradients. Should ideally be 0, but often needs to be around 0.5 or higher.

Details

Note that the `ggridges::stat_density_ridges()` makes joint density estimation across all datasets. This may not generate the desired result when using faceted plots. As an alternative, you can set `stat = "density"` to use `ggplot2::stat_density()`. In this case, it is required to add the aesthetic mapping `height = after_stat(density)` (see examples).

See Also

`ggridges::geom_density_ridges()`
`ggridges::geom_ridgeline()`
`ggridges::geom_density_ridges_gradient()`

Examples

data.frame(
  x = rep(1:5, 3), y = c(rep(0, 5), rep(1, 5), rep(3, 5)),
  height = c(0, 1, 3, 4, 0, 1, 2, 3, 5, 4, 0, 5, 4, 4, 1)
) %>%
gf_ridgeline(y ~ x, height = ~ height, group = ~y, fill = "lightblue", alpha = 0.7)
diamonds %>%
gf_density_ridges(cut ~ price,
gf_ridgeline

```r
scale = 2, fill = ~ cut, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)
diamonds %>%
gf_density_ridges(clarity ~ price | cut,
  scale = 2, fill = ~ clarity, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)
## Not run:
diamonds %>%
gf_density_ridges2(cut ~ price, scale = 2, fill = ~ cut, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)
## End(Not run)
## Not run:
diamonds %>%
gf_density_ridges(cut ~ price,
  scale = 2, fill = ~ cut, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)
## Not run:
diamonds %>%
gf_density_ridges2(cut ~ price, scale = 2, fill = ~ cut, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)
## End(Not run)
## Not run:
diamonds %>%
gf_density_ridges(cut ~ price,
  scale = 2, fill = ~ cut, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)
```

```
gf_density_ridges(clarity ~ price | cut, height = after_stat(density), stat = "density",
  scale = 2, fill = ~ clarity, alpha = 0.6, show.legend = FALSE) %>%
gf_theme(theme_ridges()) %>%
gf_refine(
  scale_y_discrete(expand = c(0.01, 0)),
  scale_x_continuous(expand = c(0.01, 0))
)

## End(Not run)
## Not run:
mosaicData::Weather %>%
gf_density_ridges_gradient(month ~ high_temp | city ~ ., fill = ~stat(x),
  group = ~ month, show.legend = FALSE, rel_min_height = 0.02) %>%
gf_refine(scale_fill_viridis_c(option = "B"), theme_bw())

## End(Not run)

gf_rug

**Formula interface to geom_rug()**

**Description**

gf_rugx() and gf_rugy() are versions that only add a rug to x- or y- axis. By default, these functions do not inherit from the formula in the original layer (because doing so would often result in rugs on both axes), so the formula is required.

**Usage**

gf_rug(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  sides = "bl",
  alpha,
  color,
  group,
  linetype,
  linewidth,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "rug",
  stat = "identity",
  position = "identity",
  show.legend = NA,
gf_rug

show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)

gf_rugx(
  object = NULL,
gformula = NULL,
data = NULL,
...
, sides = "b",
alpha,
color,
group,
linetype,
linewidth,
height = 0,
xlab,
ylab,
title,
subtitle,
caption,
geom = "rug",
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = FALSE,
environment = parent.frame()
)

gf_rugy(
  object = NULL,
gformula = NULL,
data = NULL,
...
, sides = "l",
alpha,
color,
group,
linetype,
linewidth,
width = 0,
xlab,
ylab,
title,
subtitle,
caption,
geom = "rug",
stat = "identity",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = FALSE,
environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x (gf_rug()) or ~ x (gf_rugx()) or ~ y (gf_rugy()).
data The data to be displayed in this layer. There are three options:
  If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
  A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
  A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
...
  Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value,
  (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
sides A string that controls which sides of the plot the rugs appear on. It can be set to a string containing any of "trbl", for top, right, bottom, and left.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom A character string naming the geom used to make the layer.
stat The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")
position

Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help

If TRUE, display some minimal help.

inherit

A logical indicating whether default attributes are inherited.

environment

An environment in which to look for variables not found in data.

height

amount of vertical jittering when position is jittered.

width

amount of horizontal jittering when position is jittered.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_rug()`

Examples

data(penguins, package = "palmerpenguins")
gf_point(bill_length_mm ~ bill_depth_mm, data = penguins) %>%
gf_rug(bill_length_mm ~ bill_depth_mm)

# There are several ways to control x- and y-rugs separately
gf_point(bill_length_mm ~ bill_depth_mm, data = penguins) %>%
gf_rugx(~bill_depth_mm, data = penguins, color = "red") %>%
gf_rugy(bill_length_mm ~ ., data = penguins, color = "green")

gf_point(bill_length_mm ~ bill_depth_mm, data = penguins) %>%
gf_rug(. ~ bill_depth_mm, data = penguins, color = "red", inherit = FALSE) %>%
gf_rug(bill_length_mm ~ ., data = penguins, color = "green", inherit = FALSE)

gf_point(bill_length_mm ~ bill_depth_mm, data = penguins) %>%
gf_rug(. ~ bill_depth_mm, data = penguins, color = "red", sides = "b") %>%
gf_rug(bill_length_mm ~ ., data = penguins, color = "green", sides = "l")

# jitter requires both an x and a y, but we can turn off one or the other with sides
gf_jitter(bill_length_mm ~ bill_depth_mm, data = penguins) %>%
gf_rug(color = "green", sides = "b", position = "jitter")

# rugs work with some 1-variable plots as well.
gf_histogram(~eruptions, data = faithful) %>%
gf_rug(~eruptions, data = faithful, color = "red") %>%
gf_rug(~eruptions, data = faithful, color = "navy", sides = "t")

# we can take advantage of inheritance to shorten the code
gf_histogram(~eruptions, data = faithful) %>%
gf_rug(color = "red") %>%
gf_rug(color = "navy", sides = "t")

# Need to turn off inheritance when using gf_dhistogram:
gf_dhistogram(~eruptions, data = faithful) %>%
gf_rug(~eruptions, data = faithful, color = "red", inherit = FALSE)

# using jitter with gf_histogram() requires manually setting the y value.
gf_dhistogram(~bill_depth_mm, data = penguins) %>%
gf_rug(0 ~ bill_depth_mm, data = penguins, color = "green", sides = "b", position = "jitter")

# the choice of y value can affect how the plot looks.
gf_dhistogram(~bill_depth_mm, data = penguins) %>%
gf_rug(0.5 ~ bill_depth_mm, data = penguins, color = "green", sides = "b", position = "jitter")

---

gf_segment

**Formula interface to geom_segment()**

**Description**

`geom_segment()` draws a straight line between points (x, y) and (xend, yend). `geom_curve()`
draws a curved line. See the underlying drawing function `grid::curveGrob()` for the parameters
that control the curve.

**Usage**

```r
gf_segment(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
```
Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula A formula with shape y + yend ~ x + xend.
data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
arow specification for arrow heads, as created by grid::arrow().
lineend  Line end style (round, butt, square).

xlab   Label for x-axis. See also gf_labs().

ylab   Label for y-axis. See also gf_labs().

title, subtitle, caption
Title, sub-title, and caption for the plot. See also gf_labs().

geom   A character string naming the geom used to make the layer.

stat   The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")

position Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

show.help If TRUE, display some minimal help.

inherit A logical indicating whether default attributes are inherited.

environment An environment in which to look for variables not found in data.

Value
a gg object

Specifying plot attributes
Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation
Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_segment()
Example

D <- data.frame(x1 = 2.62, x2 = 3.57, y1 = 21.0, y2 = 15.0)
gf_point(mpg ~ wt, data = mtcars) %>%
gf_curve(y1 + y2 ~ x1 + x2, data = D, color = "navy") %>%
gf_segment(y1 + y2 ~ x1 + x2, data = D, color = "red")

Description

Mapping with shape files

Usage

gf_sf(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  geometry,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  stat = "sf",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data: A data frame with the variables to be plotted.

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, (c) attributes of the layer as a whole, which are set with attribute = value, or (d) arguments for the geom, stat, or position function.

alpha: Opacity (0 = invisible, 1 = opaque).
color: A color or a formula used for mapping color.
fill: A color for filling, or a formula used for mapping fill.
group: Used for grouping.
linetype: A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth: A numerical line width or a formula used for mapping linewidth.
geometry: A column of class sfc containing simple features data. (Another option is that data may contain a column named geometry.) geometry is never inherited.
xlab: Label for x-axis. See also gf_labs().
ylab: Label for y-axis. See also gf_labs().
title, subtitle, caption: Title, sub-title, and caption for the plot. See also gf_labs().
stat: A character string naming the stat used to make the layer.
position: Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend: A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.
show.help: If TRUE, display some minimal help.
inherit: A logical indicating whether default attributes are inherited.
environment: An environment in which to look for variables not found in data.

Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.
Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

ggplot2::geom_line(), gf_point()

Examples

## Not run:
if (require(maps) && require(maptools) &&
    require(sf) && require(rgeos))
US <- sf::st_as_sf(maps::map("state", plot = FALSE, fill = TRUE))
gf_sf(fill = ~ factor(nchar(ID)), data = US) %>%
gf_refine(coord_sf())

# We can specify shape data and external data separately using geometry
MI <- sf::st_as_sf(maps::map("county", "michigan", plot = FALSE, fill = TRUE))
Migeom <- MI$geom
gf_sf(
  fill = ~ log10(population), data = MIpop %>% dplyr::arrange(county),
  geometry = ~Migeom, color = "white"
) %>%
gf_refine(coord_sf(), theme_bw())

# alternatively we can merge external data and shape data into one data frame.
MI %>%
dplyr::mutate(county = gsub("michigan," , "", ID)) %>%
dplyr::left_join(MIpop %>% dplyr::mutate(county = tolower(county))) %>%
gf_sf(fill = ~ population / 1e3) %>%
gf_refine(
  coord_sf(), theme_bw(),
  scale_fill_continuous(name = "population (thousands)", trans = "log10")
)

## End(Not run)

---

gf_sina  

**Formula interface to geom_sina()**

Description

The sina plot is a data visualization chart suitable for plotting any single variable in a multiclass dataset. It is an enhanced jitter strip chart, where the width of the jitter is controlled by the density distribution of the data within each class.
Usage

gf_sina(
    object = NULL,
    gformula = NULL,
    data = NULL,
    ..., 
    alpha,
    color,
    size,
    fill,
    group,
    xlab,
    ylab,
    title,
    subtitle,
    caption,
    geom = "point",
    stat = "sina",
    position = "identity",
    show.legend = NA,
    show.help = NULL,
    inherit = TRUE,
    environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.

data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

... Other arguments passed on to layer(). These are often aesthetics, used to set an aesthetic to a fixed value, like colour = "red" or size = 3. They may also be parameters to the paired geom/stat.

alpha Opacity (0 = invisible, 1 = opaque).

color A color or a formula used for mapping color.

size A numeric size or a formula used for mapping size.
A color for filling, or a formula used for mapping fill.

Group for grouping.

Label for x-axis. See also `gf_labs()`.

Label for y-axis. See also `gf_labs()`.

Title, sub-title, and caption for the plot. See also `gf_labs()`.

The geometric object to use to display the data, either as a ggproto Geom subclass or as a string naming the geom stripped of the `geom_` prefix (e.g. "point" rather than "geom_point")

The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the `stat_` prefix (e.g. "count" rather than "stat_count")

Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

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. It can also be a named logical vector to finely select the aesthetics to display.

If TRUE, display some minimal help.

A logical indicating whether default attributes are inherited.

An environment in which to look for variables not found in data.

Value

A `gg` object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggforce::geom_sina()`
gf_smooth

Examples

gf_sina(age ~ substance, data = mosaicData::HELPrt)

gf_smooth

Formula interface to geom_smooth()

Description

LOESS and linear model smoothers in ggformula.

Usage

gf_smooth(
    object = NULL,
    gformula = NULL,
    data = NULL,
    ...,
    method = "auto",
    formula = y ~ x,
    se = FALSE,
    method.args,
    n = 80,
    span = 0.75,
    fullrange = FALSE,
    level = 0.95,
    xlab,
    ylab,
    title,
    subtitle,
    caption,
    geom = "smooth",
    stat = "smooth",
    position = "identity",
    show.legend = NA,
    show.help = NULL,
    inherit = TRUE,
    environment = parent.frame()
)

gf_lm(
    object = NULL,
    gformula = NULL,
    data = NULL,
    ...,
    alpha = 0.3,
    lm.args = list(),
)
\begin{verbatim}
interval = "none",
level = 0.95,
fullrange = TRUE,
xlab,
ylab,
title,
ssubtitle,
caption,
geom = "lm",
stat = "lm",
position = "identity",
show.legend = NA,
show.help = NULL,
inherit = TRUE,
environment = parent.frame()
)
\end{verbatim}

Arguments

- **object**
  
  When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

- **gformula**
  
  A formula with shape \( y \sim x \). Faceting can be achieved by including \( | \) in the formula.

- **data**
  
  A data frame with the variables to be plotted.

- **...**
  
  Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = \( \sim \) expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

- **method**
  
  Smoothing method (function) to use, accepts either NULL or a character vector, e.g. "lm", "glm", "gam", "loess" or a function, e.g. MASS::rlm or mgcv::gam, stats::lm, or stats::loess. "auto" is also accepted for backwards compatibility. It is equivalent to NULL.

  For method = NULL the smoothing method is chosen based on the size of the largest group (across all panels). \texttt{stats::loess()} is used for less than 1,000 observations; otherwise \texttt{mgcv::gam()} is used with \( \text{formula} = y \sim s(x, \text{bs} = \text{"cs"}) \) with method = "REML". Somewhat anecdotally, \texttt{loess} gives a better appearance, but is \( O(N^2) \) in memory, so does not work for larger datasets.

  If you have fewer than 1,000 observations but want to use the same \texttt{gam()} model that method = NULL would use, then set method = "gam", formula = \( y \sim s(x, \text{bs} = \text{"cs"}) \).

- **formula**
  
  Formula to use in smoothing function, eg. \( y \sim x, y \sim \text{poly}(x, 2), y \sim \log(x) \). NULL by default, in which case method = NULL implies formula = \( y \sim x \) when there are fewer than 1,000 observations and formula = \( y \sim s(x, \text{bs} = \text{"cs"}) \) otherwise.

- **se**
  
  Display confidence interval around smooth? (TRUE by default, see level to control.)

- **method.args**
  
  List of additional arguments passed on to the modelling function defined by method.
n  Number of points at which to evaluate smoother.
span Controls the amount of smoothing for the default loess smoother. Smaller numbers produce wigglier lines, larger numbers produce smoother lines. Only used with loess, i.e. when method = "loess", or when method = NULL (the default) and there are fewer than 1,000 observations.
fullrange If TRUE, the smoothing line gets expanded to the range of the plot, potentially beyond the data. This does not extend the line into any additional padding created by expansion.
level Level of confidence interval to use (0.95 by default).
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
geom A character string naming the geom used to make the layer.
stat A character string naming the stat used to make the layer.
position Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
show.legend A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.
show.help If TRUE, display some minimal help.
inherit A logical indicating whether default attributes are inherited.
environment An environment in which to look for variables not found in data.
alpha Opacity (0 = invisible, 1 = opaque).
lm.args A list of arguments to stats::lm().
interval One of "none", "confidence" or "prediction".

Value

A gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form attribute = value or mapped using arguments of the form attribute = ~ expression.

In formulas of the form A | B, B will be used to form facets using facet_wrap() or facet_grid(). This provides an alternative to gf_facet_wrap() and gf_facet_grid() that is terser and may feel more familiar to users of lattice.

Evaluation

Evaluation of the ggplot2 code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.
See Also

```r
ggplot2::geom_smooth(), gf_spline()
```

Examples

```r
gf_smooth()
gf_lm()
gf_smooth(births ~ date, color = ~wday, data = mosaicData::Births78)
gf_smooth(births ~ date,
  color = ~wday, data = mosaicData::Births78,
  fullrange = TRUE
)
gf_smooth(births ~ date,
  color = ~wday, data = mosaicData::Births78,
  show.legend = FALSE, se = FALSE
)
gf_smooth(births ~ date,
  color = ~wday, data = mosaicData::Births78,
  show.legend = FALSE, se = TRUE
)
gf_lm(length ~ width,
  data = mosaicData::KidsFeet,
  color = ~biggerfoot, alpha = 0.2
)
```

```r
>%
gf_point()
gf_lm(length ~ width,
  data = mosaicData::KidsFeet,
  color = ~biggerfoot, fullrange = FALSE, alpha = 0.2
)
gf_point()
gf_lm(length ~ width,
  color = ~sex, data = mosaicData::KidsFeet,
  formula = y ~ poly(x, 2), linetype = "dashed"
)
```

```r
>%
gf_point()
gf_lm(length ~ width,
  color = ~sex, data = mosaicData::KidsFeet,
  formula = log(y) ~ x, backtrans = exp
)
```

```r
>%
gf_point()
gf_lm(hwy ~ displ,
  data = mpg,
  formula = log(y) ~ poly(x, 3), backtrans = exp,
  interval = "prediction", fill = "skyblue"
)
```

```r
>%
gf_lm(  
  formula = log(y) ~ poly(x, 3), backtrans = exp,
  interval = "confidence", color = "red"
  )
```
clotting <- data.frame(
  u = c(5,10,15,20,30,40,60,80,100),
  lot1 = c(118,58,42,35,27,25,21,19,18),
  lot2 = c(69,35,26,21,18,16,13,12,12)
)
gf_point(lot1 ~ u, data = clotting) %>%
gf_smooth(formula = y ~ log(x), method = "glm",
  method.args = list(family = Gamma))
gf_point(lot2 ~ u, data = clotting) %>%
gf_smooth(formula = y ~ log(x), color = "red", method = "glm",
  method.args = list(family = Gamma))

gf_spline

Formula interface to geom_spline()

Description

Fitting splines in ggformula.

Usage

gf_spline(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  weight,
  df,
  spar,
  tol,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "line",
  stat = "spline",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape y ~ x. Faceting can be achieved by including | in the formula.

data
A data frame with the variables to be plotted.

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

weight
An optional vector of weights. See smooth.spline().

df
desired equivalent degrees of freedom. See smooth.spline() for details.

spar
A smoothing parameter, typically in (0,1]. See smooth.spline() for details.

tol
A tolerance for sameness or uniqueness of the x values. The values are binned into bins of size tol and values which fall into the same bin are regarded as the same. Must be strictly positive (and finite). When NULL, IQR(x) * 10e-6 is used.

xlab
Label for x-axis. See also gf_labs().

ylab
Label for y-axis. See also gf_labs().

title, subtitle, caption
Title, sub-title, and caption for the plot. See also gf_labs().

geom
A character string naming the geom used to make the layer.

stat
A character string naming the stat used to make the layer.

position
Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend
A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.

Value

a gg object
Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`geom_spline()`, `gf_smooth()`, `gf_lm()`

Examples

```r
gf_spline(births ~ date, color = ~wday, data = mosaicData::Births78)
gf_spline(births ~ date, color = ~wday, data = mosaicData::Births78, df = 20)
gf_spline(births ~ date, color = ~wday, data = mosaicData::Births78, df = 4)
```

Description

This is a polar parameterisation of `geom_segment`. It is useful when you have variables that describe direction and distance.

Usage

```r
gf_spoke(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...
  angle,
  radius,
  alpha,
  color,
  group,
  linetype,
  linewidth,
)```
Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula A formula with shape y \sim x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
   If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot()
   A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
   A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. \sim \text{head}(.x, 10)).
...
   Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = \sim expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

angle The angle at which segment leaves the point (x,y).
radius The length of the segment.
alpha Opacity (0 = invisible, 1 = opaque).
color A color or a formula used for mapping color.
group Used for grouping.
linetype A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.
linewidth A numerical line width or a formula used for mapping linewidth.
xlab Label for x-axis. See also gf_labs().
ylab Label for y-axis. See also gf_labs().
title, subtitle, caption Title, sub-title, and caption for the plot. See also gf_labs().
**gf_spoke**

- **geom**: A character string naming the geom used to make the layer.
- **stat**: The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")
- **position**: Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.
- **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. It can also be a named logical vector to finely select the aesthetics to display.
- **show.help**: If TRUE, display some minimal help.
- **inherit**: A logical indicating whether default attributes are inherited.
- **environment**: An environment in which to look for variables not found in data.

**Value**

a gg object

**Specifying plot attributes**

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, B will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

**Evaluation**

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

**See Also**

`ggplot2::geom_spoke()`

**Examples**

```r
SomeData <- expand.grid(x = 1:10, y = 1:10)
SomeData$angle <- runif(100, 0, 2 * pi)
SomeData$speed <- runif(100, 0, sqrt(0.1 * SomeData$x))

gf_point(y ~ x, data = SomeData) %>%
gf_spoke(y ~ x, angle = ~angle, radius = 0.5)

gf_point(y ~ x, data = SomeData) %>%
gf_spoke(y ~ x, angle = ~angle, radius = ~speed)
```
Formula interface to geom_step()

Description
geom_path() connects the observations in the order in which they appear in the data. geom_line() connects them in order of the variable on the x axis. geom_step() creates a stairstep plot, highlighting exactly when changes occur. The group aesthetic determines which cases are connected together.

Usage

gf_step(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  alpha,
  color,
  group,
  linetype,
  linewidth,
  direction = "hv",
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "step",
  stat = "identity",
  position = "identity",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See `fortify()` for which variables will be created.

A function will be called with a single argument, the plot data. The return value must be a data frame, and will be used as the layer data. A function can be created from a formula (e.g. `~ head(.x, 10)`).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with `attribute = value`, (b) ggplot2 aesthetics to be mapped with `attribute = ~ expression`, or (c) attributes of the layer as a whole, which are set with `attribute = value`.

**alpha**
Opacity (0 = invisible, 1 = opaque).

**color**
A color or a formula used for mapping color.

**group**
Used for grouping.

**linetype**
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

**linewidth**
A numerical line width or a formula used for mapping linewidth.

**direction**
Direction of stairs: 'vh' for vertical then horizontal, 'hv' for horizontal then vertical, or 'mid' for step half-way between adjacent x-values.

**xlab**
Label for x-axis. See also `gf_labs()`.

**ylab**
Label for y-axis. See also `gf_labs()`.

**title, subtitle, caption**
Title, sub-title, and caption for the plot. See also `gf_labs()`.

**geom**
A character string naming the geom used to make the layer.

**stat**
The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")

**position**
Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

**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. It can also be a named logical vector to finely select the aesthetics to display.

**show.help**
If TRUE, display some minimal help.

**inherit**
A logical indicating whether default attributes are inherited.

**environment**
An environment in which to look for variables not found in data.

**Value**
a gg object
Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_step()`

Examples

```r
gf_step(births ~ date, data = mosaicData::Births78, color = ~wday)
```

# Roll your own Kaplan-Meier plot

```r
if (require(survival) && require(broom)) {
  # fit a survival model
  surv_fit <- survfit(coxph(Surv(time, status) ~ age + sex, lung))
  surv_fit
  # use broom::tidy() to create a tidy data frame for plotting
  surv_df <- tidy(surv_fit)
  head(surv_df)
  # now create a plot
  surv_df %>%
    gf_step(estimate ~ time) %>%
    gf_ribbon(conf.low + conf.high ~ time, alpha = 0.2)
}
```

---

**gf_text**

*Formula interface to geom_text() and geom_label()*

Description

Text geoms are useful for labeling plots. They can be used by themselves as scatterplots or in combination with other geoms, for example, for labeling points or for annotating the height of bars. `geom_text()` adds only text to the plot. `geom_label()` draws a rectangle behind the text, making it easier to read.
Usage

gf_text(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  label,
  alpha,
  angle,
  color,
  family,
  fontface,
  group,
  hjust,
  lineheight,
  size,
  vjust,
  parse = FALSE,
  nudge_x = 0,
  nudge_y = 0,
  check_overlap = FALSE,
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "text",
  stat = "identity",
  position = "nudge",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_label(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...,
  label,
  alpha,
  angle,
  color,
  family,
  fontface,
  group,
  hjust,
Arguments

- object: When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
- gformula: A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
- data: The data to be displayed in this layer. There are three options:
  - If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
  - A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
  - A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).
- ...: Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.
- label: The text to be displayed.
- alpha: Opacity (0 = invisible, 1 = opaque).
- angle: An angle for rotating the text.
- color: A color or a formula used for mapping color.
- family: A font family.
One of "plain", "bold", "italic", or "bold italic".

Used for grouping.

Numbers between 0 and 1 indicating how to justify text relative the the specified location.

Line height.

A numeric size or a formula used for mapping size.

If TRUE, the labels will be parsed into expressions and displayed as described in ?plotmath.

Horizontal and vertical adjustment to nudge labels by. Useful for offsetting text from points, particularly on discrete scales. Cannot be jointly specified with position.

If TRUE, text that overlaps previous text in the same layer will not be plotted. check_overlap happens at draw time and in the order of the data. Therefore data should be arranged by the label column before calling geom_text(). Note that this argument is not supported by geom_label().

Label for x-axis. See also gf_labs().

Label for y-axis. See also gf_labs().

Title, sub-title, and caption for the plot. See also gf_labs().

A character string naming the geom used to make the layer.

The statistical transformation to use on the data for this layer, either as a ggproto Geom subclass or as a string naming the stat stripped of the stat_ prefix (e.g. "count" rather than "stat_count")

Position adjustment, either as a string, or the result of a call to a position adjustment function. Cannot be jointly specified with nudge_x or nudge_y.

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. It can also be a named logical vector to finely select the aesthetics to display.

If TRUE, display some minimal help.

A logical indicating whether default attributes are inherited.

An environment in which to look for variables not found in data.

Amount of padding around label. Defaults to 0.25 lines.

Radius of rounded corners. Defaults to 0.15 lines.

Size of label border, in mm.

a gg object
Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form A | B, B will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of gformula. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_text()`

Examples

data(penguins, package = "palmerpenguins")
gf_text(bill_length_mm ~ bill_depth_mm,
data = penguins,
label = ~species, color = ~species, size = 2, angle = 30
)
penguins %>%
gf_point(bill_length_mm ~ bill_depth_mm, color = ~species, alpha = 0.5) %>%
gf_text(bill_length_mm ~ bill_depth_mm,
label = ~species, color = ~species,
size = 2, angle = 0, hjust = 0, nudge_x = 0.1, nudge_y = 0.1
)
if (require(dplyr)) {
data(penguins, package = "palmerpenguins")
penguins_means <-
penguins %>%
group_by(species) %>%
summarise(bill_length_mm = mean(bill_length_mm), bill_depth_mm = mean(bill_depth_mm))
gf_point(bill_length_mm ~ bill_depth_mm, data = penguins, color = ~species) %>%
gf_label(bill_length_mm ~ bill_depth_mm,
data = penguins_means,
label = ~species, color = ~species, size = 2, alpha = 0.7
)}
gf_theme

Themes for ggformula

Description
Themes for ggformula

Usage
gf_theme(object, theme, ...)

Arguments
object a gg object
theme a ggplot2 theme function like theme_minimal().
... If theme is missing, then these additional arguments are theme elements of the sort handled by ggplot2::theme().

Value
a modified gg object

gf_tile

Formula interface to geom_tile()

Description
geom_rect() and geom_tile() do the same thing, but are parameterised differently: geom_rect() uses the locations of the four corners (xmin, xmax, ymin and ymax), while geom_tile() uses the center of the tile and its size (x, y, width, height). geom_raster() is a high performance special case for when all the tiles are the same size.

Usage
gf_tile(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ...
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
Arguments

object
When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.

gformula
A formula with shape y ~ x. Faceting can be achieved by including | in the formula.

data
A data frame with the variables to be plotted.

...
Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha
Opacity (0 = invisible, 1 = opaque).

color
A color or a formula used for mapping color.

fill
A color for filling, or a formula used for mapping fill.

group
Used for grouping.

linetype
A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

linewidth
A numerical line width or a formula used for mapping linewidth.

xlab
Label for x-axis. See also gf_labs().

ylab
Label for y-axis. See also gf_labs().

title, subtitle, caption
Title, sub-title, and caption for the plot. See also gf_labs().

geom
A character string naming the geom used to make the layer.

stat
A character string naming the stat used to make the layer.

position
Either a character string naming the position function used for the layer or a position object returned from a call to a position function.

show.legend
A logical indicating whether this layer should be included in the legends. NA, the default, includes layer in the legends if any of the attributes of the layer are mapped.

show.help
If TRUE, display some minimal help.

inherit
A logical indicating whether default attributes are inherited.

environment
An environment in which to look for variables not found in data.
Value

a gg object

Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Positional attributes (a.k.a, aesthetics) are specified using the formula in gformula. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

See Also

`ggplot2::geom_tile()`

Examples

```r
D <- expand.grid(x = 0:5, y = 0:5)
D$z <- runif(nrow(D))
gf_tile(y ~ x, fill = ~z, data = D)
gf_tile(z ~ x + y, data = D)
```
gf_violin

Formula interface to geom_violin()

Description

A violin plot is a compact display of a continuous distribution. It is a blend of geom_boxplot() and geom_density(): a violin plot is a mirrored density plot displayed in the same way as a boxplot.

Usage

gf_violin(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
  alpha,
  color,
  fill,
  group,
  linetype,
  linewidth,
  weight,
  draw_quantiles = NULL,
  trim = TRUE,
  scale = "area",
  bw,
  adjust = 1,
  kernel = "gaussian",
  xlab,
  ylab,
  title,
  subtitle,
  caption,
  geom = "violin",
  stat = "ydensity",
  position = "dodge",
  show.legend = NA,
  show.help = NULL,
  inherit = TRUE,
  environment = parent.frame()
)

gf_violinh(
  object = NULL,
  gformula = NULL,
  data = NULL,
  ..., 
)
alpha, color, fill, group, linetype, linewidth, weight, draw_quantiles = NULL, trim = TRUE, scale = "area", bw, adjust = 1, kernel = "gaussian", xlab, ylab, title, subtitle, caption, geom = "violinh", stat = "xdensity", position = "dodgev", show.legend = NA, show.help = NULL, inherit = TRUE, environment = parent.frame() )

Arguments

object When chaining, this holds an object produced in the earlier portions of the chain. Most users can safely ignore this argument. See details and examples.
gformula A formula with shape y ~ x. Faceting can be achieved by including | in the formula.
data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

... Additional arguments. Typically these are (a) ggplot2 aesthetics to be set with attribute = value, (b) ggplot2 aesthetics to be mapped with attribute = ~ expression, or (c) attributes of the layer as a whole, which are set with attribute = value.

alpha Opacity (0 = invisible, 1 = opaque).
**gf_violin**

- **color**
  A color or a formula used for mapping color.

- **fill**
  A color for filling, or a formula used for mapping fill.

- **group**
  Used for grouping.

- **linetype**
  A linetype (numeric or "dashed", "dotted", etc.) or a formula used for mapping linetype.

- **linewidth**
  A numerical line width or a formula used for mapping linewidth.

- **weight**
  Useful for summarized data, weight provides a count of the number of values with the given combination of x and y values.

- **draw_quantiles**
  If not(NULL) (default), draw horizontal lines at the given quantiles of the density estimate.

- **trim**
  If TRUE (default), trim the tails of the violins to the range of the data. If FALSE, don’t trim the tails.

- **scale**
  If "area" (default), all violins have the same area (before trimming the tails). If "count", areas are scaled proportionally to the number of observations. If "width", all violins have the same maximum width.

- **bw**
  The smoothing bandwidth to be used. If numeric, the standard deviation of the smoothing kernel. If character, a rule to choose the bandwidth, as listed in `stats::bw.nrd()`.

- **adjust**
  A multiplicative bandwidth adjustment. This makes it possible to adjust the bandwidth while still using the a bandwidth estimator. For example, adjust = 1/2 means use half of the default bandwidth.

- **kernel**
  Kernel. See list of available kernels in `density()`.

- **xlab**
  Label for x-axis. See also `gf_labs()`.

- **ylab**
  Label for y-axis. See also `gf_labs()`.

- **title, subtitle, caption**
  Title, sub-title, and caption for the plot. See also `gf_labs()`.

- **geom, stat**
  Use to override the default connection between `geom_violin()` and `stat_ydensity()`.

- **position**
  Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use `position_jitter`), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

- **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. It can also be a named logical vector to finely select the aesthetics to display.

- **show.help**
  If TRUE, display some minimal help.

- **inherit**
  A logical indicating whether default attributes are inherited.

- **environment**
  An environment in which to look for variables not found in data.

**Value**

- a gg object
Specifying plot attributes

Positional attributes (a.k.a, aesthetics) are specified using the formula in `gformula`. Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form `attribute = value` or mapped using arguments of the form `attribute = ~ expression`.

In formulas of the form `A | B`, `B` will be used to form facets using `facet_wrap()` or `facet_grid()`. This provides an alternative to `gf_facet_wrap()` and `gf_facet_grid()` that is terser and may feel more familiar to users of `lattice`.

Evaluation

Evaluation of the `ggplot2` code occurs in the environment of `gformula`. This will typically do the right thing when formulas are created on the fly, but might not be the right thing if formulas created in one environment are used to create plots in another.

References


See Also

`ggplot2::geom_violin()`

Examples

```r
gf_violin(age ~ substance, data = mosaicData::HELPrct)
gf_violin(age ~ substance, data = mosaicData::HELPrct, fill = ~sex)
gf_violinh(substance ~ age, data = mosaicData::HELPrct)
gf_violinh(substance ~ age, data = mosaicData::HELPrct, fill = ~sex)
```

---

**ggformula**

*Formula interface to ggplot2*

Description

Formula interface to `ggplot2`

The ggformula system

The functions in `ggformula` provide a formula interface to `ggplot2` layer functions and a system for working with pipes to create multi-layer plots and to refine plots. For plots with just one layer, the formula interface is more compact than native `ggplot2` code and is consistent with modeling functions like `stats::lm()` that use a formula interface and with the numerical summary functions in the `mosaic` package.
Specifying plot attributes

Positional attributes (a.k.a aesthetics) are typically specified using a formula (see the \texttt{gformula} argument). Setting and mapping of additional attributes can be done through the use of additional arguments. Attributes can be set can be set using arguments of the form \texttt{attribute = value} or mapped using arguments of the form \texttt{attribute = ~ expression}. A (sometimes partial) list of available attributes can be obtained by executing plotting functions with no arguments.

In formulas of the form \( A \mid B \), \( B \) will be used to form facets using \texttt{facet\_wrap()} or \texttt{facet\_grid()}. This provides an alternative to \texttt{gf\_facet\_wrap()} and \texttt{gf\_facet\_grid()} that is terser and may feel more familiar to users of \texttt{lattice}.

Evaluation

Evaluation of the \texttt{ggplot2} code occurs in the environment specified by \texttt{environment}. This will typically do the right thing, but is exposed in case some non-standard behavior is desired. In earlier versions, the environment of the formula was used, but since some functions in the package do not require a formula, a separate argument is used now.

Examples

\begin{verbatim}
apropos("gf_")
gf_point()
\end{verbatim}

---

\texttt{layer\_factory} \hspace{1cm} Create a \texttt{ggformula} layer function

Description

Primarily intended for package developers, this function factory is used to create the layer functions in the \texttt{ggformula} package.

Usage

\begin{verbatim}
layer_factory(
  geom = "point",
  position = "identity",
  stat = "identity",
  pre = {
    },
  aes_form = y ~ x,
  extras = alist(),
  note = NULL,
  aesthetics = aes(),
  inherit.aes = TRUE,
  check.aes = TRUE,
  data = NULL,
  layer_fun = quo(ggplot2::layer),
  ...
)
\end{verbatim}
Arguments

- `geom`: The geom to use for the layer (may be specified as a string).
- `position`: The position function to use for the layer (may be specified as a string).
- `stat`: The stat function to use for the layer (may be specified as a string).
- `pre`: Code to run as a "pre-process".
- `aes_form`: A single formula or a list of formulas specifying how attributes are inferred from the formula. Use NULL if the function may be used without a formula.
- `extras`: An alist of additional arguments (potentially with defaults).
- `note`: A note to add to the quick help.
- `aesthetics`: Additional aesthetics (typically created using ggplot2::aes()) set rather than inferred from formula. `gf_dhistogram()` uses this to set the y aesthetic to `stat(density)`, for example.
- `inherit.aes`: A logical indicating whether aesthetics should be inherited from prior layers or a vector of character names of aesthetics to inherit.
- `check.aes`: A logical indicating whether a warning should be emitted when aesthetics provided don’t match what is expected.
- `data`: A data frame or NULL or NA.
- `layer_fun`: The function used to create the layer or a quosure that evaluates to such a function.
- `...`: Additional arguments.

Value

A function.

---

Mlpop  

*Population of Michigan counties*

Description

Population of Michigan counties

Usage

data(Mlpop)

Format

A data frame with populations of Michigan counties.

- `rank`  Population rank.
- `county` County name.
percs_by_group

Compute groupwise proportions and percents

Description

Transform a vector of counts and a vector of groups into a vector of proportions or percentages within groups.

Usage

percs_by_group(x, group)
props_by_group(x, group)

Arguments

x A vector of counts

group A vector to determine groups.

Examples

x <- c(20, 30, 30, 70)
g1 <- c("A", "A", "B", "B")
g2 <- c("A", "B", "A", "B")
props_by_group(x, g1)
percs_by_group(x, g1)
percs_by_group(x, g2)

StatAsh

ggproto classes for ggplot2

Description

These are typically accessed through their associated geom_*, stat_* or gf_* functions.

Usage

StatAsh
StatSpline
StatQqline
StatLm
GeomLm
StatAsh
StatFitdistr

See Also

stat_ash()
gf_ash()
stat_spline()
gf_spline()
stat_qq()
gf_qq()
stat_lm()
gf_lm()
geom_lm()
gf_lm()
stat_ash()
gf_ash()

---

stat_fitdistr

Description

This stat computes points for plotting a distribution function. Fitting is done using MASS::fitdistr() when analytic solutions are not available.

Usage

stat_fitdistr(
  mapping = NULL,
  data = NULL,
  geom = "path",
  position = "identity",
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE,
  dist = "dnorm",
  start = NULL,
  ...
)


stat_lm

Arguments

- **mapping**: Aesthetics created using `aes()` or `aes_string()`.
- **data**: A data frame.
- **geom**: A character string naming the geom used to make the layer.
- **position**: Either a character string naming the position function used for the layer or a position object returned from a call to a position function.
- **na.rm**: If TRUE, do not emit a warning about missing data.
- **show.legend**: A 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.
- **dist**: A character string indicating the distribution to fit. Examples include "dnorm", "dgamma", etc.
- **start**: A list of starting values used by `MASS::fitdistr()` when numerically approximating the maximum likelihood estimate.
- **...**: Additional arguments.

Value

A gg object

stat_lm

Linear Model Displays

Description

Adds linear model fits to plots. `geom_lm()` and `stat_lm()` are essentially equivalent. Use `geom_lm()` unless you want a non-standard geom.

Usage

```r
stat_lm(
  mapping = NULL,
  data = NULL,
  geom = "lm",
  position = "identity",
  interval = c("none", "prediction", "confidence"),
  level = 0.95,
  formula = y ~ x,
  lm.args = list(),
  backtrans = identity,
  ...
)
```

na.rm = FALSE,
show.legend = NA,
inherit.aes = TRUE
geom_lm(
  mapping = NULL,
  data = NULL,
  stat = "lm",
  position = "identity",
  interval = c("none", "prediction", "confidence"),
  level = 0.95,
  formula = y ~ x,
  lm.args = list(),
  backtrans = identity,
  ...
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE
)

Arguments

mapping Set of aesthetic mappings created by aes(). If specified and inherit.aes = TRUE (the default), it is combined with the default mapping at the top level of the plot. You must supply mapping if there is no plot mapping.

data The data to be displayed in this layer. There are three options:
If NULL, the default, the data is inherited from the plot data as specified in the call to ggplot().
A data.frame, or other object, will override the plot data. All objects will be fortified to produce a data frame. See fortify() for which variables will be created.
A function will be called with a single argument, the plot data. The return value must be a data.frame, and will be used as the layer data. A function can be created from a formula (e.g. ~ head(.x, 10)).

gem, stat Use to override the default connection between geom_lm and stat_lm.
position Position adjustment, either as a string naming the adjustment (e.g. "jitter" to use position_jitter), or the result of a call to a position adjustment function. Use the latter if you need to change the settings of the adjustment.

interval One of "none", "confidence" or "prediction".

level The level used for confidence or prediction intervals

formula a formula describing the model in terms of y (response) and x (predictor).

lm.args A list of arguments supplied to lm() when performing the fit.

backtrans a function that transforms the response back to the original scale when the formula includes a transformation on y.

... Other arguments passed on to layer(). These are often aesthetics, used to set an aesthetic to a fixed value, like colour = "red" or size = 3. They may also be parameters to the paired geom/stat.
stat_lm

na.rm

If FALSE, the default, missing values are removed with a warning. If TRUE, missing values are silently removed.

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. It can also be a named logical vector to finely select the aesthetics to display.

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()`.

Details

Stat calculation is performed by the (currently undocumented) `predictdf`. Pointwise confidence or prediction bands are calculated using the `predict()` method.

See Also

`lm()` for details on linear model fitting.

Examples

```r
library(ggplot2)
library(mosaicData)

# basic model
ggplot(data = mosaicData::KidsFeet, aes(y = length, x = width, color = sex)) +
  geom_lm() +
  geom_point()

# with prediction bands
ggplot(data = mosaicData::KidsFeet, aes(y = length, x = width, color = sex)) +
  geom_lm(interval = "prediction", color = "skyblue") +
  geom_lm(interval = "confidence") +
  geom_point() +
  facet_wrap(~sex)

# non-standard display
ggplot(data = mosaicData::KidsFeet, aes(y = length, x = width, color = sex)) +
  stat_lm(aes(fill = sex),
          color = NA, interval = "confidence", geom = "ribbon",
          alpha = 0.2
        ) +
  geom_point() +
  facet_wrap(~sex)

ggplot(mpg, aes(displ, hwy)) +
  geom_lm(
    formula = log(y) ~ poly(x, 3), backtrans = exp,
    interval = "prediction", fill = "skyblue"
  ) +
  geom_lm(
    formula = log(y) ~ poly(x, 3), backtrans = exp, interval = "confidence",
    color = "red"
  ) +
  geom_point()
```
stat_qqline

A Stat for Adding Reference Lines to QQ-Plots

Description

This stat computes quantiles of the sample and theoretical distribution for the purpose of providing reference lines for QQ-plots.

Usage

```r
stat_qqline(
  mapping = NULL,
  data = NULL,
  geom = "line",
  position = "identity",
  ..., 
  distribution = stats::qnorm,
  dparams = list(),
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE
)
```

Arguments

- **mapping**: An aesthetic mapping produced with `aes()` or `aes_string()`.
- **data**: A data frame.
- **geom**: A geom.
- **position**: A position object.
- **distribution**: A quantile function.
- **dparams**: A list of arguments for `distribution`.
- **na.rm**: A logical indicating whether a warning should be issued when missing values are removed before plotting.
- **show.legend**: A logical indicating whether legends should be included for this layer. If `NA`, legends will be include for each aesthetic that is mapped.
- **inherit.aes**: A logical indicating whether aesthetics should be inherited. When `FALSE`, the supplied mapping will be the only aesthetics used.

Examples

```r
data(penguins, package = "palmerpenguins")
ggplot(data = penguins, aes(sample = bill_length_mm)) +
  geom_qq() +
  stat_qqline(alpha = 0.7, color = "red", linetype = "dashed") +
  facet_wrap(~species)
```
stat_spline

Geoms and stats for spline smoothing

Description

Similar to geom_smooth, this adds spline fits to plots.

Usage

stat_spline(
  mapping = NULL,
  data = NULL,
  geom = "line",
  position = "identity",
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE,
  weight = NULL,
  df = NULL,
  spar = NULL,
  cv = FALSE,
  all.knots = FALSE,
  nknots = stats::.nknots.smspl,
  df.offset = 0,
  penalty = 1,
  control.spar = list(),
  tol = NULL,
  ...
)

geom_spline(
  mapping = NULL,
  data = NULL,
  stat = "spline",
  position = "identity",
  na.rm = FALSE,
  show.legend = NA,
  inherit.aes = TRUE,
  weight = NULL,
  df = NULL,
  spar = NULL,
  cv = FALSE,
  all.knots = FALSE,
  nknots = stats::.nknots.smspl,
  df.offset = 0,
  penalty = 1,
  control.spar = list(),
Arguments

mapping An aesthetic mapping produced with \texttt{aes()} or \texttt{aes_string()}.
data A data frame.
geom A geom.
position A position object.
na.rm A logical indicating whether a warning should be issued when missing values are removed before plotting.
show.legend A logical indicating whether legends should be included for this layer. If \texttt{NA}, legends will be included for each aesthetic that is mapped.
inherit.aes A logical indicating whether aesthetics should be inherited. When \texttt{FALSE}, the supplied mapping will be the only aesthetics used.
weight An optional vector of weights. See \texttt{smooth.spline(\ldots)}.
df desired equivalent degrees of freedom. See \texttt{smooth.spline(\ldots)} for details.
spar A smoothing parameter, typically in \((0,1]\). See \texttt{smooth.spline(\ldots)} for details.
cv A logical. See \texttt{smooth.spline(\ldots)} for details.
all.knots A logical. See \texttt{smooth.spline(\ldots)} for details.
nknots An integer or function giving the number of knots to use when \texttt{all.knots = \texttt{FALSE}}. See \texttt{smooth.spline(\ldots)} for details.
df.offset A numerical value used to increase the degrees of freedom when using GVC. See \texttt{smooth.spline(\ldots)} for details.
penalty the coefficient of the penalty for degrees of freedom in the GVC criterion. See \texttt{smooth.spline(\ldots)} for details.
control.spar An optional list used to control root finding when the parameter \texttt{spar} is computed. See \texttt{smooth.spline(\ldots)} for details.
tol A tolerance for sameness or uniqueness of the x values. The values are binned into bins of size \texttt{tol} and values which fall into the same bin are regarded as the same. Must be strictly positive (and finite). When \texttt{NULL}, \texttt{IQR(x) \times 10^{-6}} is used.
... Additional arguments
stat A stat.

Examples

\begin{verbatim}
if (require(mosaicData)) {
  ggplot(Births) + geom_spline(aes(x = date, y = births, colour = wday))
  ggplot(Births) + geom_spline(aes(x = date, y = births, colour = wday), nknots = 10)
}
\end{verbatim}
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