Package ‘ggvis’

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add_axis

Add a vega axis specification to a ggvis plot

Description

Axis specifications allow you to either override the default axes, or additional axes.

Usage

```r
add_axis(
  vis,
  type,
  scale = NULL,
  orient = NULL,
  title = NULL,
  title_offset = NULL,
  format = NULL,
  ticks = NULL,
```
values = NULL,
subdivide = NULL,
tick_padding = NULL,
tick_size_major = NULL,
tick_size_minor = tick_size_major,
tick_size_end = tick_size_major,
offset = NULL,
layer = "back",
grid = TRUE,
properties = NULL)
)

hide_axis(vis, scale)

Arguments

vis A ggvis object.
type The type of axis. Either x or y.
scale The name of the scale backing the axis component. Defaults to the scale type -
you will need to specify if you want (e.g.) a scale for a secondary y-axis.
orient The orientation of the axis. One of top, bottom, left or right. The orientation can
be used to further specialize the axis type (e.g., a y axis oriented for the right
edge of the chart) - defaults to bottom for x axes, and left for y axes.
title A title for the axis. By default, it uses the name of the field in the first data set
used by the scale. Use "" to suppress the title.
title_offset The offset (in pixels) from the axis at which to place the title.
format The formatting pattern for axis labels. Vega uses D3’s format pattern.
ticks A desired number of ticks. The resulting number may be different so that values
are "nice" (multiples of 2, 5, 10) and lie within the underlying scale’s range.
values Explicitly set the visible axis tick values.
subdivide If provided, sets the number of minor ticks between major ticks (the value 9
results in decimal subdivision).
tick_padding The padding, in pixels, between ticks and text labels.
tick_size_major, tick_size_minor, tick_size_end
The size, in pixels, of major, minor and end ticks.
offset The offset, in pixels, by which to displace the axis from the edge of the enclosing
group or data rectangle.
layer A string indicating if the axis (and any gridlines) should be placed above or
below the data marks. One of "front" or "back" (default).
grid A flag indicating if gridlines should be created in addition to ticks.
properties Optional mark property definitions for custom axis styling. Should be an object
created by axis_props, with properties for ticks, majorTicks, minorTicks, grid,
labels, title, and axis.
Details

More information about axes can be found in the "axes and legends" vignettes.

Compared to ggplot2

In ggplot2, axis (and legend) properties are part of the scales specification. In vega, they are separate, which allows the specification of multiple axes, and more flexible linkage between scales and axes.

See Also

Vega axis documentation: https://vega.github.io/vega/docs/axes/

Examples

```r
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~cyl) %>%
  layer_points() %>%
  add_axis("x", title = "Weight", orient = "top")

# Suppress axis with hide_axis
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~cyl) %>%
  layer_points() %>%
  hide_axis("x") %>%
  hide_axis("y")

mtcars %>% ggvis(x = ~wt, y = ~mpg) %>%
  layer_points() %>%
  add_axis("x", title = "Weight", ticks = 40,
  properties = axis_props(
    ticks = list(stroke = "red"),
    majorTicks = list(strokeWidth = 2),
    grid = list(stroke = "red"),
    labels = list(
      fill = "steelblue",
      angle = 50,
      fontSize = 14,
      align = "left",
      baseline = "middle",
      dx = 3
    ),
    title = list(fontSize = 16),
    axis = list(stroke = "#333", strokeWidth = 1.5)
  )
```

**add_data**

*Add dataset to a visualisation*

**Description**

Add dataset to a visualisation
Usage

```
add_data(vis, data, name = deparse2(substitute(data)), add_suffix = TRUE)
```

Arguments

- **vis**: Visualisation to modify.
- **data**: Data set to add.
- **name**: Data of data - optional, but helps produce informative error messages.
- **add_suffix**: Should a unique suffix be added to the data object’s ID? This should only be FALSE when the spec requires a data set with a specific name.

Examples

```
mtcars %>% ggvis(~mpg, ~wt) %>% layer_points()
NULL %>% ggvis(~mpg, ~wt) %>% add_data(mtcars) %>% layer_points()
```

---

**add_guide_axis**: Defunct function for adding an axis

Description

This function has been replaced with `add_axis`.

Usage

```
add_guide_axis(...)  
```

Arguments

- **...**: Other arguments.

---

**add_guide_legend**: Defunct function for adding a legend

Description

This function has been replaced with `add_legend`.

Usage

```
add_guide_legend(...)  
```

Arguments

- **...**: Other arguments.
add_legend

Add a vega legend specification to a ggvis plot

Description

Axis specifications allow you to either override the default legends, or supply additional legends.

Usage

add_legend(
  vis,
  scales = NULL,
  orient = "right",
  title = NULL,
  format = NULL,
  values = NULL,
  properties = NULL
)

hide_legend(vis, scales)

Arguments

vis A ggvis object.

scales The name of one or more scales for which to add a legend. Typically one of "size", "shape", "fill", "stroke", although custom scale names may also be used. Multiple names can also be used, like c("fill", "shape").

orient The orientation of the legend. One of "left" or "right". This determines how the legend is positioned within the scene. The default is "right".

title A title for the legend. By default, it uses the name the fields used in the legend. Use "" to suppress the title.

format The formatting pattern for axis labels. Vega uses D3’s format pattern.

values Explicitly set the visible legend values.

properties Optional mark property definitions for custom legend styling. Should be an object created by legend_props, with properties for title, label, symbols, gradient, legend.

Details

More information about axes can be found in the "axes and legends" vignettes.

Compared to ggplot2

In ggplot2, legend (and axis) properties are part of the scales specification. In vega, they are separate, which allows the specification of multiple legends, and more flexible linkage between scales and legends.
Examples

```r
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~cyl) %>%
layer_points() %>%
add_legend("fill", title = "Cylinders")

# Suppress legend with hide_legend
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~cyl) %>%
layer_points() %>%
hide_legend("fill")

# Combining two properties in one legend
mtcars %>%
ggvis(x = ~wt, y = ~mpg, fill = ~factor(cyl), shape = ~factor(cyl)) %>%
layer_points() %>%
add_legend(c("fill", "shape"))

# Control legend properties with a continuous legend, with x and y position
# in pixels.
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~cyl) %>%
layer_points() %>%
add_legend("fill", title = "Cylinders",
properties = legend_props(
  title = list(fontSize = 16),
  labels = list(fontSize = 12, fill = "#00F"),
  gradient = list(stroke = "red", strokeWidth = 2),
  legend = list(x = 500, y = 50)
))

# Control legend properties with a categorical legend, with x and y position
# in the scaled data space.
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~factor(cyl)) %>%
layer_points() %>%
add_legend("fill", title = "Cylinders",
properties = legend_props(
  title = list(fontSize = 16),
  labels = list(fontSize = 14, dx = 5),
  symbol = list(stroke = "black", strokeWidth = 2,
               shape = "square", size = 200),
  legend = list(
    x = scaled_value("x", 4.5),
    y = scaled_value("y", 30)
  )
))

# Control legend position using x_rel and y_rel which specify relative
# position, going from 0 to 1. (0, 0) is the bottom-left corner, and
# (1, 1) is the upper-right corner. The values control the position of
# the upper-left corner of the legend.
mtcars %>% ggvis(x = ~wt, y = ~mpg, fill = ~cyl) %>%
layer_points() %>%
```
add_props

```
add_relative_scales() %>%
  add_legend("fill", title = "Cylinders",
  properties = legend_props(
    legend = list(
      x = scaled_value("x_rel", 0.8),
      y = scaled_value("y_rel", 1)
    )
  )
)
```

---

**add_props**

*Add visual properties to a visualisation*

### Description

Add visual properties to a visualisation

### Usage

```r
add_props(vis, ..., .props = NULL, inherit = NULL, env = parent.frame())
```

### Arguments

- **vis**
  - Visualisation to modify.

- **...**
  - A set of name-value pairs. The name should be a valid vega property.
  - The first two unnamed components are taken to be `x` and `y`. Any additional unnamed components will raise an error.

- **.props**
  - When calling `props` from other functions, you’ll often have a list of quoted function functions. You can pass that function to the `.props` argument instead of messing around with `substitute`. In other words, `.props` lets you opt out of the non-standard evaluation that `props` does.

- **inherit**
  - If `TRUE`, the defaults, will inherit from properties from the parent layer. If `FALSE`, it will start from nothing.

- **env**
  - The environment in which to evaluate variable properties.

### Examples

```r
mtcars %>% ggvis(~wt, ~mpg) %>% layer_points()
mtcars %>% ggvis(~wt, ~mpg) %>% add_props(~wt, ~mpg) %>% layer_points()
mtcars %>% ggvis(~wt) %>% add_props(y = ~mpg) %>% layer_points()
```
add_relative_scales  Add x_rel and y_rel scales

Description

This function adds scales named x_rel and y_rel, each of which has a domain of 0 to 1, and the range is the plot’s width or height. These scales are useful for positioning visual elements relative to the plotting area. For example, with legends.

Usage

add_relative_scales(vis)

Arguments

vis A ggvis object.

See Also

add_legend for a usage example.

add_tooltip  Add tooltips to a plot.

Description

Add tooltips to a plot.

Usage

add_tooltip(vis, html, on = c("hover", "click"))

Arguments

vis Visualisation to add tooltips to.

html A function that takes a single argument as input. This argument will be a list containing the data in the mark currently under the mouse. It should return a string containing HTML or NULL to hide tooltip for the current element.

on Should tooltips appear on hover, or on click?
Examples

```r
## Run these examples only in interactive R sessions
if (interactive()) {

all_values <- function(x) {
  if(is.null(x)) return(NULL)
  paste0(names(x), " : ", format(x), collapse = "<br />")
}

base <- mtcars %>% ggvis(x = ~wt, y = ~mpg) %>%
  layer_points() %>%
  add_tooltip(all_values, "hover") %>%
  add_tooltip(all_values, "click")

# The data sent from client to the server contains only the data columns that
# are used in the plot. If you want to get other columns of data, you should
# to use a key to line up the item from the plot with a row in the data.
mtc <- mtcars
mtc$id <- 1:nrow(mtc) # Add an id column to use ask the key

all_values <- function(x) {
  if(is.null(x)) return(NULL)
  row <- mtc[mtc$id == x$id, ]
  paste0(names(row), " : ", format(row), collapse = "<br />")
}

mtc %>% ggvis(x = ~wt, y = ~mpg, key := ~id) %>%
  layer_points() %>%
  add_tooltip(all_values, "hover")
}
```

---

**auto_group**

*Automatically group data by grouping variables*

**Description**

Use `auto_group` to group up a dataset on all categorical variables specified by props, and have each piece rendered by the same mark.

**Usage**

```r
auto_group(vis, exclude = NULL)
```

**Arguments**

- `vis` The ggvis visualisation to modify.
- `exclude` A vector containing names of props to exclude from auto grouping. It is often useful to exclude `c("x", "y")`, when one of those variables is categorical.
See Also

To manually specify grouping variables, see group_by.

Examples

```r
# One line
mtcars %>% ggvis(~disp, ~mpg, stroke = ~factor(cyl)) %>% layer_paths()

# One line for each level of cyl
mtcars %>% ggvis(~disp, ~mpg, stroke = ~factor(cyl)) %>% group_by(cyl) %>%
         layer_paths()

# The grouping column can already be stored as a factor
mtcars2 <- mtcars
mtcars2$cyl <- factor(mtcars2$cyl)
mtcars2 %>% ggvis(~disp, ~mpg, stroke = ~cyl) %>% auto_group() %>%
         layer_paths()
```

---

`axis_props`

Create an `axis_props` object for controlling axis properties.

Description

The items in each of the lists can be a literal value, like 5 or "blue", or they can be a `scaled_value` object.

Usage

```r
axisProps(
  ticks = NULL,
  majorTicks = NULL,
  minorTicks = NULL,
  grid = NULL,
  labels = NULL,
  title = NULL,
  axis = NULL
)
```

Arguments

- `ticks`: A named list of line properties for ticks.
- `majorTicks`: A named list of line properties for major ticks.
- `minorTicks`: A named list of line properties for minor ticks.
- `grid`: A named list of line properties for grid lines.
- `labels`: A named list of text properties for axis labels.
- `title`: A named list of text properties for the axis title.
- `axis`: A named list of line properties for the axis line.
Description

Bands are used to set the width or height on categorical scales - a band represent the height or width allocated for one level of a factor.

Usage

band(offset = NULL, mult = NULL)

is.prop_band(x)

Arguments

offset, mult  Additive and multiplicate offsets used to adjust the band size. For example, use mult = 0.9 to make a bar take up 90% of the space allocated for its category.

x  object to test for band-ness

Examples

df <- data.frame(label = c("a", "b", "c"), n = c(10, 9, 4))

base <- df %>% ggvis(~label, y2 = 0, y = ~n)
base %>% layer_rects(width = band())
base %>% layer_rects(width = band(offset = -1))
base %>% layer_rects(width = band(mult = 0.9))

# A nominal scale with padding is more symmetrical than band with a mult
base %>% layer_rects(width = band(mult = 0.75))
base %>% layer_rects(width = band(mult = 0.75)) %>%
  scale_nominal("x", padding = 0.25, points = FALSE)

cocaine  Cocaine seizures in the US.

Description

This dataset comes from STRIDE, the System to Retrieve Information from Drug Evidence. It contains all concaine seizures in the US from 2007 that have a known weight.

Usage

cocaine
compute_align

**Format**
Data frame with 3380 observations of 5 variables.

**Variables**
- **state**: State where seizure occurred.
- **potency**: Purity of cocaine, as percentage (100% = pure cocaine, 0% = all filler)
- **weight**: Weight, in grams.
- **month**: Month in which seizure occurred.
- **price**: Estimated value in USD.

**Use**
Use of this data requires your agreement to refer to your analyses as "unvalidated DEA data and to claim authorship and responsibility for any inferences and/or conclusions you may draw from this information."

---

**compute_align**

*Align positions using length.*

**Description**
This compute function is often used in conjunction with `compute_count`, when used on data with a continuous x variable. By default, the computed width will be equal to the resolution of the data, or, in other words the smallest difference between two values in the data.

**Usage**

```r
compute_align(x, var, length = NULL, align = 0.5, dir = "x")
```

**Arguments**
- **x**: Dataset-like object to align. Built-in methods for data frames, grouped data frames and ggvis visualisations.
- **var**: Name of variable to compute width of.
- **length**: An absolute length to use. If NULL (the default), the width will be equivalent to the resolution of the data.
- **align**: Where does the existing variable fall on the new bins? 0 = left edge, 0.5 = center, 1 = right edge.
- **dir**: Direction, i.e. "x" or "y". Used to generate variable names in output.

**Details**
An absolute width for each x can be specified by using the `width` argument. If `width` is NULL (the default), it will use the resolution of the data as the width.
Value

The original data frame, with additional columns:

'dir'min_ left boundary of bin
'dir'max_ right boundary of bin
'dir'len_ width of bin

See Also

compute_bin For counting cases within ranges of a continuous variable.
compute_count For counting cases at specific values of a variable.

Examples

```r
ggplot(mtcars) + geom_histogram(bins = 3, position = 'identity')
motors %>% compute_bin(~hp) %>% compute_align(~x_)
```

# Use a specific width
```r
gp <- ggplot(pressure) + geom_histogram(bins = 3, position = 'identity')
gp + compute_bin(~temperature) %>% compute_align(~x_, length = 5)
```

# It doesn't matter whether you transform inside or outside of a vis
```r
ggplot(mtcars) %>% geom_histogram(aes(x = disp))
ggplot(mtcars) %>% geom_histogram(bins = 3, position = 'identity') %>% compute_align(~x_, length = .5)
```

# Varying align
```r
ggplot(mtcars) %>% geom_histogram(bins = 3, position = 'identity') %>% compute_align(~x_, length = .5, align = input_slider(0, 1))
```

---

**compute_bin**  
*Bin data along a continuous variable*

Description

Bin data along a continuous variable
Usage

```r
compute_bin(
  x,
  x_var,
  w_var = NULL,
  width = NULL,
  center = NULL,
  boundary = NULL,
  closed = c("right", "left"),
  pad = FALSE,
  binwidth
)
```

Arguments

- **x**: Dataset-like object to bin. Built-in methods for data frames, grouped data frames and ggvis visualisations.
- **x_var, w_var**: Names of x and weight variables. The x variable must be continuous.
- **width**: The width of the bins. The default is `NULL`, which yields 30 bins that cover 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.
- **center**: The center of one of the bins. Note that if center is above or below the range of the data, things will be shifted by an appropriate number of widths. To center on integers, for example, use `width=1` and `center=0`, even if 0 is outside the range of the data. At most one of `center` and `boundary` may be specified.
- **boundary**: A boundary between two bins. As with `center`, things are shifted when `boundary` is outside the range of the data. For example, to center on integers, use `width=1` and `boundary = 0.5`, even if 1 is outside the range of the data. At most one of `center` and `boundary` may be specified.
- **closed**: One of "right" or "left" indicating whether right or left edges of bins are included in the bin.
- **pad**: If `TRUE`, adds empty bins at either end of x. This ensures frequency polygons touch 0. Defaults to `FALSE`.
- **binwidth**: Deprecated; use `width` instead.

Value

A data frame with columns:

- **count_**: the number of points
- **x_**: mid-point of bin
- **xmin_**: left boundary of bin
- **xmax_**: right boundary of bin
- **width_**: width of bin
compute_boxplot

See Also

compute_count For counting cases at specific locations of a continuous variable. This is useful when the variable is continuous but the data is granular.

Examples

mtcars %>% compute_bin(~mpg)
mtcars %>% compute_bin(~mpg, width = 10)
mtcars %>% group_by(cyl) %>% compute_bin(~mpg, width = 10)

# It doesn't matter whether you transform inside or outside of a vis
mtcars %>% compute_bin(~mpg) %>% ggvis(~x_, ~count_) %>% layer_paths()
mtcars %>% ggvis(~ x_, ~ count_) %>% compute_bin(~mpg) %>% layer_paths()

# Missing values get own bin
mtcars2 <- mtcars
mtcars2$mpg[sample(32, 5)] <- NA
mtcars2 %>% compute_bin(~mpg, width = 10)

# But are currently silently dropped in histograms
mtcars2 %>% ggvis(~mpg) %>% layer_histograms(~mpg)

compute_boxplot Calculate boxplot values

Description

Calculate boxplot values

Usage

compute_boxplot(x, var = NULL, coef = 1.5)

Arguments

x Dataset-like object to compute boxplot values. There are built-in methods for data frames, grouped data frames, and ggvis visualisations.

var Name of variable for which to compute boxplot values. The variable must be continuous.

coef The maximum length of the whiskers as multiple of the inter-quartile range. Default value is 1.5.

Value

A data frame with columns:

min_ Lower whisker = smallest observation greater than or equal to lower hinge - 1.5 * IQR
lower_ Lower hinge (25th percentile)  
median_ Median (50th percentile)  
upper_ Upper hinge (75th percentile)  
max_ Upper whisker = largest observation less than or equal to upper hinge + 1.5 * IQR  
outliers_ A vector of values that are outside of the min and max

See Also  
layer_boxplots

Examples  
mtcars %>% compute_boxplot(~mpg)  
mtcars %>% group_by(cyl) %>% compute_boxplot(~mpg)

compute_count Count data at each location

Description  
Count data at each location

Usage  
compute_count(x, x_var, w_var = NULL)

Arguments  
x Dataset-like object to count. Built-in methods for data frames, grouped data frames and ggvis visualisations.  
x_var, w_var Names of x and weight variables.

Value  
A data frame with columns:  
count_ the number of points  
x_ the x value where the count was made

The width of each "bin" is set to the resolution of the data – that is, the smallest difference between two x values.

See Also  
compute_bin For counting cases within ranges of a continuous variable.  
compute_align For calculating the "width" of data.
compute_density

Examples

```r
mtcars %>% compute_count(~cyl)

# Weight the counts by car weight value
mtcars %>% compute_count(~cyl, ~wt)

# If there's one weight value at each x, it effectively just renames columns.
pressure %>% compute_count(~temperature, ~pressure)

# Also get the width of each bin
pressure %>% compute_count(~temperature, ~pressure) %>% compute_align(~x_)
```

```r
# It doesn't matter whether you transform inside or outside of a vis
mtcars %>% compute_count(~cyl, ~wt) %>%
  compute_align(~x_) %>%
  ggvis(x = ~xmin_, x2 = ~xmax_, y = ~count_, y2 = 0) %>%
  layer_rects()

mtcars %>%
  ggvis(x = ~xmin_, x2 = ~xmax_, y = ~count_, y2 = 0) %>%
  compute_count(~cyl, ~wt) %>%
  compute_align(~x_) %>%
  layer_rects()
```

---

compute_density  Compute density of data.

Description

Compute density of data.

Usage

```r
compute_density(
  x,
  x_var,
  w_var = NULL,
  kernel = "gaussian",
  trim = FALSE,
  n = 256L,
  na.rm = FALSE,
  ...
)
```

Arguments

- `x`: Dataset (data frame, grouped_df or ggvis) object to work with.
- `x_var`: Names of variables to use for x position, and for weights.
- `w_var`: Names of variables to use for x position, and for weights.
- `kernel`: Smoothing kernel. See `density` for details.
trim  If TRUE, the default, density estimates are trimmed to the actual range of the data. If FALSE, they are extended by the default 3 bandwidths (as specified by the cut parameter to density).

n  Number of points (along x) to use in the density estimate.

na.rm  If TRUE missing values will be silently removed, otherwise they will be removed with a warning.

...  Additional arguments passed on to density.

Value

A data frame with columns:

pred_ regularly spaced grid of n locations

resp_ density estimate

Examples

mtcars %>% compute_density(~mpg, n = 5)
mtcars %>% group_by(cyl) %>% compute_density(~mpg, n = 5)
mtcars %>% ggvis(~mpg) %>% compute_density(~mpg, n = 5) %>%
layer_points(~pred_, ~resp_)

compute_model_prediction

Create a model of a data set and compute predictions.

Description

Fit a 1d model, then compute predictions and (optionally) standard errors over an evenly spaced grid.

Usage

compute_model_prediction(
  x,
  formula,
  ..., 
  model = NULL,
  se = FALSE,
  level = 0.95,
  n = 80L,
  domain = NULL,
  method
)

compute_smooth(x, formula, ..., span = 0.75, se = FALSE)
compute_model_prediction

Arguments

x  
Dataset-like object to model and predict. Built-in methods for data frames, grouped data frames and ggvis visualisations.

formula  
Formula passed to modelling function. Can use any variables from data.

...  
arguments passed on to model function

model  
Model fitting function to use - it must support R’s standard modelling interface, taking a formula and data frame as input, and returning predictions with predict. If not supplied, will use loess for <= 1000 points, otherwise it will use gam. Other modelling functions that will work include lm, glm and rlm.

se  
include standard errors in output? Requires appropriate method of predict_grid, since the interface for returning predictions with standard errors is not consistent across modelling frameworks.

level  
the confidence level of the standard errors.

n  
the number of grid points to use in the prediction

domain  
If NULL (the default), the domain of the predicted values will be the same as the domain of the prediction variable in the data. It can also be a two-element numeric vector specifying the min and max.

method  
Deprecated. Please use model instead.

span  
Smoothing span used for loess model.

Details

compute_model_prediction fits a model to the data and makes predictions with it. compute_smooth is a special case of model predictions where the model is a smooth loess curve whose smoothness is controlled by the span parameter.

Value

A data frame with columns:

resp_ regularly spaced grid of n locations
pred_ predicted value from model
pred_lwr_ and pred_upr_    upper and lower bounds of confidence interval (if se = TRUE)
pred_se_ the standard error (width of the confidence interval) (if se = TRUE)

Examples

# Use a small value of n for these examples
mtcars %>% compute_model_prediction(mpg ~ wt, n = 10)
mtcars %>% compute_model_prediction(mpg ~ wt, n = 10, se = TRUE)
mtcars %>% group_by(cyl) %>% compute_model_prediction(mpg ~ wt, n = 10)

# compute_smooth defaults to loess
mtcars %>% compute_smooth(mpg ~ wt)
# Override model to suppress message or change approach
mtcars %>% compute_model_prediction(mpg ~ wt, n = 10, model = "loess")
mtcars %>% compute_model_prediction(mpg ~ wt, n = 10, model = "lm")

# Set the domain manually
mtcars %>
  compute_model_prediction(mpg ~ wt, n = 20, model = "lm", domain = c(0, 8))

# Plot the results
mtcars %>% compute_model_prediction(mpg ~ wt) %>%
  ggvis(~pred_, ~resp_) %>%
  layer_paths()
mtcars %>% ggvis() %>%
  compute_model_prediction(mpg ~ wt) %>%
  layer_paths(~pred_, ~resp_)

compute_stack Stack overlapping data.

Description
Stack overlapping data.

Usage
compute_stack(x, stack_var = NULL, group_var = NULL)

Arguments
  x A data object
  stack_var A string specifying the stacking variable.
  group_var A string specifying the grouping variable.

Value
A data frame with columns:
stack_upr_ the lower y coordinate for a stack bar
stack_lwr_ the upper y coordinate for a stack bar

Examples
mtcars %>% cbind(count = 1) %>% compute_stack(~count, ~cyl)

# Shouldn't use or affect existing grouping
mtcars %>% cbind(count = 1) %>% group_by(am) %>% compute_stack(~count, ~cyl)

# If given a ggvis object, will use x variable for stacking by default
mtcars %>% ggvis(x = ~cyl, y = ~wt) %>%
compute_stack(stack_var = ~wt, group_var = ~cyl) %>%
layer_rects(x = ~cyl - 0.5, x2 = ~cyl + 0.5, y = ~stack_upr_,
y2 = ~stack_lwr_)

# Collapse across hair & eye colour data across sex
hec <- as.data.frame(xtabs(Freq ~ Hair + Eye, HairEyeColor))
hec %>% compute_stack(~Freq, ~Hair)

# Without stacking - bars overlap
hec %>% ggvis(~Hair, ~Freq, fill = ~Eye, fillOpacity := 0.5) %>%
layer_rects(y2 = 0, width = band())

# With stacking
hec %>% ggvis(x = ~Hair, y = ~Freq, fill = ~Eye, fillOpacity := 0.5) %>%
compute_stack(~Freq, ~Hair) %>%
layer_rects(y = ~stack_lwr_, y2 = ~stack_upr_, width = band())

# layer_bars stacks automatically:
hec %>% ggvis(~Hair, ~Freq, fill = ~Eye, fillOpacity := 0.5) %>%
group_by(Eye) %>%
layer_bars(width = 1)

---

**compute_tabulate**  
*Count data at each location of a categorical variable*

**Description**

Count data at each location of a categorical variable

**Usage**

```r
compute_tabulate(x, x_var, w_var = NULL)
```

**Arguments**

- `x`  
  Dataset-like object to count. Built-in methods for data frames, grouped data frames and ggvis visualisations.

- `x_var, w_var`  
  Names of x and weight variables.

**Value**

A data frame with columns:

- `count_`  
  the number of points

- `x_`  
  value of bin
See Also

- `compute_bin` For counting cases within ranges of a continuous variable.
- `compute_count` For counting cases at specific locations of a continuous variable. This is useful when the variable is continuous but the data is granular.

Examples

```r
library(dplyr)
# The tabulated column must be countable (not numeric)
## Not run: mtcars %>% compute_tabulate(~cyl)
mtcars %>% mutate(cyl = factor(cyl)) %>% compute_tabulate(~cyl)

# Or equivalently:
mtcars %>% compute_tabulate(~factor(cyl))

# If there's one weight value at each x, it effectively just renames columns.
pipeline %>% compute_tabulate(~factor(temperature), ~pressure)

# It doesn't matter whether you transform inside or outside of a vis
mtcars %>% compute_tabulate(~factor(cyl)) %>%
  ggvis(x = ~x_, y = ~count_, y2 = 0) %>%
  layer_rects(width = band())

mtcars %>%
  ggvis(x = ~x_, y = ~count_, y2 = 0) %>%
  compute_tabulate(~factor(cyl)) %>%
  layer_rects(width = band())

# compute_tabulate is used automatically in layer_bars when no y prop
# is supplied.
mtcars %>% ggvis(x = ~factor(cyl)) %>% layer_bars()
```

---

**explain**

*Explain details of an object*

**Description**

This is a generic function which gives more details about an object than `print`, and is more focused on human readable output than `str`.

**See Also**

`dplyr::explain` for more information.

**Examples**

```r
p <- mtcars %>% ggvis(x = ~cyl) %>% layer_bars()
explain(p)
```
explain.ggvis

Print out the structure of a ggvis object in a friendly format

Description
Print out the structure of a ggvis object in a friendly format

Usage
## S3 method for class 'ggvis'
explain(x, ...)

Arguments
x Visualisation to explain
... Needed for compatibility with generic. Ignored by this method.

export_png

Export a PNG or SVG from a ggvis object

Description
This requires that the external program vg2png is installed. This is part of the vega node.js module.

Usage
export_png(vis, file = NULL)
export_svg(vis, file = NULL)

Arguments
vis A ggvis object.
file Output file name. If NULL, defaults to "plot.svg" or "plot.png".

See Also
https://github.com/trifacta/vega for information on installing vg2png and vg2svg.

Examples
## Not run:
mtcars %>% ggvis(x = ~wt) %>% export_png()
## End(Not run)
**get_data**  
*Get data from a ggvis object*

**Description**

This function is useful for inspecting the data in a ggvis object.

**Usage**

```r
get_data(vis)
```

**Arguments**

- **vis**: A ggvis object.

**Examples**

```r
p <- cocaine %>% ggvis(~price) %>% layer_bars()
get_data(p)
```

---

**ggvis**  
*Visualise a data set with a ggvis graphic.*

**Description**

`ggvis` is used to turn a dataset into a visualisation, setting up default mappings between variables in the dataset and visual properties. Nothing will be displayed until you add additional layers.

**Usage**

```r
ggvis(data = NULL, ..., env = parent.frame())
```

**Arguments**

- **data**: A data object.
- **...**: Property mappings. If not named, the first two mappings are taken to be `x` and `y`. Common properties are `x`, `y`, `stroke`, `fill`, `opacity`, `shape`.
- **env**: Environment in which to evaluate properties.
Examples

# If you don’t supply a layer, ggvis uses layer_guess() to guess at
# an appropriate type:
mtcars %>% ggvis(~mpg, ~wt)
mtcars %>% ggvis(~mpg, ~wt, fill = ~cyl)
mtcars %>% ggvis(~mpg, ~wt, fill := "red")
mtcars %>% ggvis(~mpg)

# ggvis has a functional interface: every ggvis function takes a ggvis
# an input and returns a modified ggvis as output.
layer_points(ggvis(mtcars, ~mpg, ~wt))

# To make working with this interface more natural, ggvis imports the
# pipe operator from magrittr. x %>% f(y) is equivalent to f(x, y) so
# we can rewrite the previous command as
mtcars %>% ggvis(~mpg, ~wt) %>% layer_points()

# For more complicated plots, add a line break after %>%
mtcars %>%
ggvis(~mpg, ~wt) %>%
layer_points() %>%
layer_smooths()

---

ggvisControlOutput  

Create a ggvis control output element in UI

Description

This is effectively the same as uiOutput, except that on the client side it may call some plot resizing
functions after new controls are drawn.

Usage

ggvisControlOutput(outputId, plotId = NULL)

Arguments

outputId  The output variable to read the value from.
plotId  An optional plot ID or vector of plot IDs. The plots will have their .onControlOutput functions called after the controls are drawn.

Details

ggvisControlOutput is intended to be used with bind_shiny on the server side.

Examples

ggvisControlOutput("plot1")
ggvis_message  Send a message to ggvis running on client

Description

This will be sent to the client and passed to a handler in ggvis.messages on the client side. The handler is specified by type.

Usage

ggvis_message(session, type, data = NULL, id = NULL)

Arguments

- session: A session object.
- type: A string representing the type of the message.
- data: An object (typically a list) containing information for the client.
- id: A unique identifier for ggvis message handler (optional).

---

group_by  Divide data into groups.

Description

Divide data into groups.

Arguments

- x: a visualisation
- ...: variables to group by.
- add: By default, when add = FALSE, group_by will override existing groups. To instead add to the existing groups, use add = TRUE
**handle_brush**  
*Handle brush events on a visualisation.*

Description

Currently for brush events to be triggered on a visualisation, you must use a `.brush` property. This limitation will be lifted in the future.

Usage

```r
display_brush(vis, on_move = NULL, fill = "black")
```

Arguments

- **vis**
  Visualisation to listen to.
- **on_move**
  Callback function with arguments:
  - **items** A data frame containing information about the items under the plot. An empty data.frame if no points under the brush.
  - **page_loc** Location of the brush with respect to the page
  - **plot_loc** Location of the brush with respect to the plot
  - **session** The session, used to communicate with the browser
- **fill** Colour of the brush.

Examples

```r
# Display tooltip when objects are brushed
mtcars %>%
ggvis(x = ~wt, y = ~mpg, size.brush := 400) %>%
layer_points() %>%
handle_brush(function(items, page_loc, session, ...) {
  show_tooltip(session, page_loc$r + 5, page_loc$t, html = nrow(items))
})
```

**handle_click**  
*Handle mouse actions on marks.*

Description

Handle mouse actions on marks.

Usage

```r
display_click(vis, on_click = NULL)
```

```r
display_hover(vis, on_mouse_over = NULL, on_mouse_out = NULL)
```
Arguments

```r
vis Visualisation to listen to.

on_click, on_mouse_over
Callback function with arguments:
 data A data frame with one row
 location A named list with components x and y
 session The session, used to communicate with the browser

on_mouse_out
Callback function with argument:
 session The session, used to communicate with the browser
```

Examples

```r
location <- function(location, ...) cat(location$x, "x", location$y, "\n")
mtcars %>% ggvis(~mpg, ~wt) %>% layer_points() %>%
 handle_click(location)
mtcars %>% ggvis(~mpg, ~wt) %>% layer_points() %>%
 handle_hover(function(...) cat("over\n"), function(...) cat("off\n"))
mtcars %>% ggvis(~mpg, ~wt) %>% layer_points() %>%
 handle_hover(function(data, ...) str(data))
```

Description

Handlers and interactive inputs for plot sizing.

Usage

```r
handle_resize(vis, on_resize)
plot_width(vis)
plot_height(vis)
```

Arguments

```r
vis Visualisation to listen to.

on_resize Callback function with arguments:
 width,height Width and height in pixels
 padding A named list of four components giving the padding in each direction
 session The session, used to communicate with the browser
```
Examples

# This example just prints out the current dimensions to the console
mtcars %>% ggvis(~mpg, ~wt) %>%
  layer_points() %>%
  handle_resize(function(width, height, ...) cat(width, "x", height, "\n"))

# Use plot_width() and plot_height() to dynamically get the plot size
# inside the plot.
mtcars %>% ggvis(~mpg, ~wt) %>% layer_text(text := plot_width())
mtcars %>% ggvis(~mpg, ~wt) %>% layer_text(text := plot_height())

input_checkbox

Create an interactive checkbox.

Description

Create an interactive checkbox.

Usage

input_checkbox(
  value = FALSE,  
  label = "",  
  id = rand_id("checkbox_"),  
  map = identity
)

Arguments

value Initial value (TRUE or FALSE).
label Display label for the control, or NULL for no label.
id A unique identifier for this input. Usually generated automatically.
map A function with single argument x, the value of the control on the client. Returns
modified value.

See Also

Other interactive input: input_select(), input_slider(), input_text()

Examples

input_checkbox(label = "Confidence interval")
input_checkbox(label = "Confidence interval", value = TRUE)

# Used in layer_smooths
mtcars %>% ggvis(~wt, ~mpg) %>%
layer_smooths(se = input_checkbox(label = "Confidence interval"))

# Used with a map function, to convert the boolean to another type of value
model_type <- input_checkbox(label = "Use flexible curve", map = function(val) if(val) "loess" else "lm")
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_model_predictions(model = model_type)

---

**input_select** Create interactive control to select one (or more options) from a list.

**Description**

- `input_radiobuttons` only ever selects one value
- `input_checkboxgroup` can always select multiple values
- `input_select` can select only one if `multiple = FALSE`, otherwise the user can select multiple by using modifier keys

**Usage**

```r
input_select(
  choices,
  selected = NULL,
  multiple = FALSE,
  label = "",
  id = rand_id("select_"),
  map = identity,
  selectize = FALSE
)

input_radiobuttons(
  choices,
  selected = NULL,
  label = "",
  id = rand_id("radio_"),
  map = identity
)

input_checkboxgroup(
  choices,
  selected = NULL,
  label = "",
  id = rand_id("radio_"),
  map = identity
)```
Arguments

- **choices**: List of values to select from. If elements of the list are named, then that name — rather than the value — is displayed to the user. It's also possible to group related inputs by providing a named list whose elements are (either named or unnamed) lists, vectors, or factors. In this case, the outermost names will be used as the group labels (leveraging the `<optgroup>` HTML tag) for the elements in the respective sublist. See the example section for a small demo of this feature.

- **selected**: The initially selected value (or multiple values if `multiple = TRUE`). If not specified then defaults to the first value for single-select lists and no values for multiple select lists.

- **multiple**: Is selection of multiple items allowed?

- **label**: Display label for the control, or `NULL` for no label.

- **id**: A unique identifier for this input. Usually generated automatically.

- **map**: A function with single argument `x`, the value of the control on the client. Returns a modified value.

- **selectize**: Whether to use `selectize.js` or not.

See Also

Other interactive input: `input_checkbox()`, `input_slider()`, `input_text()`

Examples

```r
# Dropdown
input_select(c("a", "b", "c"))
input_select(c("a", "b", "c"), multiple = TRUE)
input_select(c("a", "b", "c"), selected = "c")

# If you want to select variable names, you need to convert
# the output of the input to a name with map so that they get
# computed correctly
input_select(names(mtcars), map = as.name)

# Radio buttons
input_radiobuttons(choices = c("Linear" = "lm", "LOESS" = "loess"),
                   label = "Model type")
input_radiobuttons(choices = c("Linear" = "lm", "LOESS" = "loess"),
                   selected = "loess",
                   label = "Model type")

# Used in layer_model_predictions
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_model_predictions(model = input_radiobuttons(
    choices = c("Linear" = "lm", "LOESS" = "loess"),
    selected = "loess",
    label = "Model type"))

# Checkbox group
mtcars %>% ggvis(x = ~wt, y = ~mpg) %>%
```
layer_points(
    fill := input_checkboxgroup(
        choices = c("Red" = "r", "Green" = "g", "Blue" = "b"),
        label = "Point color components",
        map = function(val) {
            rgb(0.8 * "r" %in% val, 0.8 * "g" %in% val, 0.8 * "b" %in% val)
        }
    )
)

---

**input_slider**

Create an interactive slider.

**Description**

Create an interactive slider.

**Usage**

```r
input_slider(
    min,
    max,
    value = (min + max)/2,
    step = NULL,
    round = FALSE,
    format = NULL,
    locale = "us",
    ticks = TRUE,
    animate = FALSE,
    sep = ",",
    pre = NULL,
    post = NULL,
    label = "",
    id = rand_id("slider_"),
    map = identity
)
```

**Arguments**

- **min, max**
  
  The minimum and maximum values (inclusive) that can be selected.

- **value**
  
  The initial value of the slider, either a number, a date (class Date), or a date-time (class POSIXt). A length one vector will create a regular slider; a length two vector will create a double-ended range slider. Must lie between `min` and `max`.

- **step**
  
  Specifies the interval between each selectable value on the slider. Either `NULL`, the default, which uses a heuristic to determine the step size or a single number. If the values are dates, `step` is in days; if the values are date-times, `step` is in seconds.
input_text

Create an interactive text or numeric input box.

Description

input_numeric only allows numbers and comes with a spin box control. input_text allows any type of input.

Usage

input_text(value, label = "", id = rand_id("text_"), map = identity)

input_numeric(value, label = "", id = rand_id("numeric_"), map = identity)
Arguments

value Initial value.
label Display label for the control, or NULL for no label.
id A unique identifier for this input. Usually generated automatically.
map A function with single argument x, the value of the control on the client. Returns a modified value.

See Also

Other interactive input: `input_checkbox()`, `input_select()`, `input_slider()`

Examples

```r
fill_text <- input_text(label = "Point color", value = "red")
mtcars %>% ggvis(~wt, ~mpg, fill := fill_text) %>% layer_bars()

size_num <- input_numeric(label = "Point size", value = 25)
mtcars %>% ggvis(~wt, ~mpg, size := size_num) %>% layer_points()
```

is.broker

Determine if an object is a broker object

Description

Determine if an object is a broker object

Usage

is.broker(x)

Arguments

x An object to test.
layer_bars

Display data with bars (a barchart).

Description

This will add bars to a plot. The exact behavior is complicated because the term bar chart is used to describe four important variations on a theme. The action of layer_bars depends on two factors: whether or not a y prop has been specified, and whether the x props is continuous or categorical.

Usage

layer_bars(vis, ..., stack = TRUE, width = NULL)

Arguments

vis Visualisation to modify
... Visual properties used to override defaults.
stack If there are multiple bars to be drawn at an x location, should the bars be stacked? If FALSE, the bars will be overplotted on each other.
width Width of each bar. When x is continuous, this controls the width in the same units as x. When x is categorical, this controls the width as a proportion of the spacing between items (default is 0.9).

Visualisations

If no y prop has been specified, then this will count the number of entries at each unique x value. There will be one bar at each unique x value, and the y value (or height) of each bar will represent the count at that x value.

If a y prop has been specified, then those y values will be used as weights for a weighted count at each unique x value. If no x values appear more than once in the data, then the end result is a plot where the height of the bar at each x value is simply the y value. However, if an x value appear more than once in the data, then this will sum up the y values at each x.

If the x variable is continuous, then a continuous x axis will be used, and the width of each bar is by default equal to the resolution of the data – that is, the smallest difference between any two x values.

If the x variable is categorical, then a categorical x axis will be used. By default, the width of each bar is 0.9 times the space between the items.

See Also

layer_histograms For bar graphs of counts at each unique x value, in contrast to a histogram’s bins along x ranges.
compute_count and compute_tabulate for more information on how data is transformed.
Examples

# Discrete x: bar graph of counts at each x value
cocaine %>% ggvis(~state) %>% layer_bars()

# Continuous x: bar graph of counts at unique locations
cocaine %>% ggvis(~month) %>% layer_bars()

# Use y prop to weight by additional variable. This is also useful
# if you have pretabulated data
cocaine %>% ggvis(~state, ~weight) %>% layer_bars()
cocaine %>% ggvis(~month, ~weight) %>% layer_bars()

# For continuous x, layer_bars is useful when the variable has a few
# unique values that you want to preserve. If you have many unique
# values and you want to bin, use layer_histogram
cocaine %>% ggvis(~price) %>% layer_bars()
cocaine %>% ggvis(~price) %>% layer_histograms(width = 100)

# If you have unique x values, you can use layer_bars() as an alternative
# to layer_points()
perspective %>% ggvis(~temperature, ~pressure) %>% layer_points()
perspective %>% ggvis(~temperature, ~pressure) %>% layer_bars()

# When x is continuous, width controls the width in x units
perspective %>% ggvis(~temperature, ~pressure) %>% layer_bars(width = 10)
# When x is categorical, width is proportional to spacing between bars
perspective %>% ggvis(~factor(temperature), ~pressure) %>%
  layer_bars(width = 0.5)

# Stacked bars
# If grouping var is continuous, you need to manually specify grouping
ToothGrowth %>% group_by(dose) %>%
  ggvis(x = ~supp, y = ~len, fill = ~dose) %>% layer_bars()
# If grouping var is categorical, grouping is done automatically
cocaine %>% ggvis(x = ~state, fill = ~as.factor(month)) %>%
  layer_bars()

layer_boxplots

Description

This will add boxplots to a plot. The action of layer_boxplots depends on whether the x prop is continuous or categorical.

Usage

layer_boxplots(vis, ..., coef = 1.5, width = NULL)
layer_densities

Arguments

vis Visualisation to modify

... Visual properties used to override defaults.

coeef The maximum length of the whiskers as multiple of the inter-quartile range. Default value is 1.5.

width Width of each bar. When x is continuous, this controls the width in the same units as x. When x is categorical, this controls the width as a proportion of the spacing between items (default is 0.9).

Details

The upper and lower "hinges" correspond to the first and third quartiles (the 25th and 75th percentiles). This differs slightly from the method used by the boxplot function, and may be apparent with small samples. See boxplot.stats for more information on how hinge positions are calculated for boxplot.

The upper whisker extends from the hinge to the highest value that is within 1.5 * IQR of the hinge, where IQR is the inter-quartile range, or distance between the first and third quartiles. The lower whisker extends from the hinge to the lowest value within 1.5 * IQR of the hinge. Data beyond the end of the whiskers are outliers and plotted as points (as specified by Tukey).

See Also

compute_boxplot for more information on how data is transformed.

Examples

library(dplyr)

mtcars %>% ggvis(~factor(cyl), ~mpg) %>% layer_boxplots()
# Set the width of the boxes to half the space between tick marks
mtcars %>% ggvis(~factor(cyl), ~mpg) %>% layer_boxplots(width = 0.5)

# Continuous x: boxes fill width between data values
mtcars %>% ggvis(~cyl, ~mpg) %>% layer_boxplots()
# Setting width=0.5 makes it 0.5 wide in the data space, which is 1/4 of the distance between data values in this particular case.
mtcars %>% ggvis(~cyl, ~mpg) %>% layer_boxplots(width = 0.5)

# Smaller outlier points
mtcars %>% ggvis(~factor(cyl), ~mpg) %>% layer_boxplots(size := 20)
Description

transform_density is a data transformation that computes a kernel density estimate from a dataset. layer_density combines transform_density with mark_path and mark_area to display a smooth line and its standard error.

Usage

layer_densities(
  vis,
  ...,  
  kernel = "gaussian",
  adjust = 1,
  density_args = list(),
  area = TRUE
)

Arguments

vis The visualisation to modify
... Visual properties, passed on to props.
kernel Smoothing kernel. See density for details.
adjust Multiple the default bandwidth by this amount. Useful for controlling wiggliness of density.
density_args Other arguments passed on to compute_density and thence to density.
area Should there be a shaded region drawn under the curve?

Examples

# Basic density estimate
faithful %>% ggvis(~waiting) %>% layer_densities()
faithful %>% ggvis(~waiting) %>% layer_densities(area = FALSE)

# Control bandwidth with adjust
faithful %>% ggvis(~waiting) %>% layer_densities(adjust = .25)
faithful %>% ggvis(~waiting) %>%
  layer_densities(adjust = input_slider(0.1, 5))

# Control stroke and fill
faithful %>% ggvis(~waiting) %>%
  layer_densities(stroke := "red", fill := "red")

# With groups
PlantGrowth %>% ggvis(~weight, fill = ~group) %>% group_by(group) %>%
  layer_densities()
PlantGrowth %>% ggvis(~weight, stroke = ~group) %>% group_by(group) %>%
  layer_densities(strokeWidth := 3, area = FALSE)
layer_guess

Guess the right type of layer based on current properties.

Description

layer_guess provides the magic behind the default behaviour of ggvis.

Usage

layer_guess(vis, ...)

Arguments

vis
The visualisation to add the new layer to.

...
Other arguments passed on individual layers.

Defaults

- Continuous x, layer_histograms
- Categorical x, layer_bars
- Continuous x and y, layer_points

Examples

# A scatterplot:
mtcars %>% ggvis(~mpg, ~wt)
mtcars %>% ggvis(~mpg, ~wt) %>% layer_guess()

# A histogram:
mtcars %>% ggvis(~mpg)
mtcars %>% ggvis(~mpg) %>% layer_guess()

layer_histograms

Display binned data

Description

Display binned data
Usage

layer_histograms(
  vis,
  ..., width = NULL,
  center = NULL,
  boundary = NULL,
  closed = c("right", "left"),
  stack = TRUE,
  binwidth
)

layer_freqpolys(
  vis,
  ..., width = NULL,
  center = NULL,
  boundary = NULL,
  closed = c("right", "left"),
  binwidth
)

Arguments

vis Visualisation to modify
...
width The width of the bins. The default is NULL, which yields 30 bins that cover 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.
center The center of one of the bins. Note that if center is above or below the range of the data, things will be shifted by an appropriate number of widths. To center on integers, for example, use width=1 and center=0, even if 0 is outside the range of the data. At most one of center and boundary may be specified.
boundary A boundary between two bins. As with center, things are shifted when boundary is outside the range of the data. For example, to center on integers, use width = 1 and boundary = 0.5, even if 1 is outside the range of the data. At most one of center and boundary may be specified.
closed One of "right" or "left" indicating whether right or left edges of bins are included in the bin.
stack If TRUE, will automatically stack overlapping bars.
binwidth Deprecated; use width instead.

See Also

layer_bars For bar graphs of counts at each unique x value, in contrast to a histogram’s bins along x ranges.
Examples

# Create histograms and frequency polygons with layers
mtcars %>% ggvis(~mpg) %>% layer_histograms()
mtcars %>% ggvis(~mpg) %>% layer_histograms(width = 2)
mtcars %>% ggvis(~mpg) %>% layer_freqpolys(width = 2)

# These are equivalent to combining compute_bin with the corresponding
# mark
mtcars %>% compute_bin(~mpg) %>% ggvis(~x_, ~count_) %>% layer_paths()

# With grouping
mtcars %>% ggvis(~mpg, fill = ~factor(cyl)) %>% group_by(cyl) %>%
  layer_histograms(width = 2)
mtcars %>% ggvis(~mpg, stroke = ~factor(cyl)) %>% group_by(cyl) %>%
  layer_freqpolys(width = 2)

layer_lines

Layer lines on a plot.

Description

layer_lines differs from layer_paths in that layer_lines sorts the data on the x variable, so
the line will always proceed from left to right, whereas layer_paths will draw a line in whatever
order appears in the data.

Usage

layer_lines(vis, ...)

Arguments

vis Visualisation to modify.
...
Visual properties.

See Also

layer_paths

Examples

mtcars %>% ggvis(~wt, ~mpg, stroke = ~factor(cyl)) %>% layer_lines()

# Equivalent to
mtcars %>% ggvis(~wt, ~mpg, stroke = ~factor(cyl)) %>%
group_by(cyl) %>% dplyr::arrange(wt) %>% layer_paths()
layer_model_predictions

Overlay model predictions or a smooth curve.

Description

layer_model_predictions fits a model to the data and draw it with layer_paths and, optionally, layer_ribbons. layer_smooths is a special case of layering model predictions where the model is a smooth loess curve whose smoothness is controlled by the span parameter.

Usage

layer_model_predictions(
  vis,
  ..., 
  model,
  formula = NULL,
  model_args = NULL,
  se = FALSE,
  domain = NULL
)

layer_smooths(vis, ..., span = 0.75, se = FALSE)

Arguments

vis Visualisation to modify

... Visual properties. Stroke properties control only affect line, fill properties only affect standard error band.

model Name of the model as a string, e.g. "loess", "lm", or "MASS::rlm". Must be the name of a function that produces a standard model object with a predict method. For layer_smooth this is always "loess".

formula Model formula. If not supplied, guessed from the visual properties, constructing y ~ x.

model_args A list of additional arguments passed on to the model function.

se Also display a point-wise standard error band? Defaults to FALSE because interpretation is non-trivial.

domain If NULL (the default), the domain of the predicted values will be the same as the domain of the prediction variable in the data. It can also be a two-element numeric vector specifying the min and max.

span For layer_smooth, the span of the loess smoother.
Examples

```r
mtcars %>% ggvis(~wt, ~mpg) %>% layer_smooths()
mtcars %>% ggvis(~wt, ~mpg) %>% layer_smooths(se = TRUE)

# Use group by to display multiple smoothes
mtcars %>% ggvis(~wt, ~mpg) %>% group_by(cyl) %>% layer_smooths()

# Control appearance with props
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_smooths(se = TRUE, stroke := "red", fill := "red", strokeWidth := 5)

# Control the wiggliness with span. Default is 0.75
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  layer_smooths(span = 0.2)
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  layer_smooths(span = 1)

# Map to an input to modify interactively
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  layer_smooths(span = input_slider(0.2, 1))

# Use other modelling functions with layer_model_predictions
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  layer_model_predictions(model = "lm") %>%
  layer_model_predictions(model = "MASS::rlm", stroke := "red")

# Custom domain for predictions
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  layer_model_predictions(model = "lm", domain = c(0, 8))
mtcars %>% ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  layer_model_predictions(model = "lm",
                          domain = input_slider(0, 10, value = c(1, 4)))

# layer_smooths() is just compute_smooth() + layer_paths()
# Run loess or other model outside of a visualisation to see what variables
# you get.
mtcars %>% compute_smooth(mpg ~ wt)
mtcars %>% compute_model_prediction(mpg ~ wt, model = "lm")

mtcars %>%
  ggvis(~wt, ~mpg) %>%
  layer_points() %>%
  compute_smooth(mpg ~ wt) %>%
  layer_paths(~pred_, ~resp_, strokeWidth := 2)
```

Description

Interactive inputs bound to arrow keys.
Usage

left_right(min, max, value = (min + max)/2, step = (max - min)/40)

up_down(min, max, value = (min + max)/2, step = (max - min)/40)

Arguments

min                A minimum value.
max                A maximum value.
value             The initial value before any keys are pressed. Defaults to half-way between min
                  and max.
step              How much each key press changes value. Defaults to 40 steps along range

Examples

size <- left_right(1, 801, value = 51, step = 50)
opacity <- up_down(0, 1, value = 0.9, step = 0.05)

mtcars %>% ggvis(~mpg, ~wt, size := size, opacity := opacity) %>%
layer_points()

legend_props

Create an axis_props object for controlling legend properties.

Description

The items in each of the lists can be a literal value, like 5 or "blue", or they can be a scaled_value
object.

Usage

legend_props(
  title = NULL,
  labels = NULL,
  symbols = NULL,
  gradient = NULL,
  legend = NULL
)

Arguments

title          A named list of text properties for the legend title.
labels         A named list of text properties for legend labels.
symbols        A named list of line properties for symbols (for discrete legend items).
gradient       A named list of line properties a continuous color gradient.
legend         A named list of line properties for the overall legend. The x and y position can
                be set here, which will override automatic positioning.
linked_brush

Create a linked brush object.

Description
A linked brush has two sides: input and output

Usage
linked_brush(keys, fill = "red")

Arguments
keys vector of all possible keys, if known.
fill brush colour

Value
A list with components:
input A function that takes a visualisation as an argument and adds an input brush to that plot
selected A reactive providing a logical vector that describes which points are under the brush
fill A reactive that gives the fill colour of points under the brush

Note
linked_brush is very new and is likely to change substantially in the future

Examples
lb <- linked_brush(keys = 1:nrow(mtcars), "red")

# Change the colour of the points
mtcars %>%
ggvis(~disp, ~mpg) %>%
layer_points(fill := lb$fill, size.brush := 400) %>%
lb$input()

# Display one layer with all points and another layer with selected points
library(shiny)
mtcars %>%
ggvis(~disp, ~mpg) %>%
layer_points(size.brush := 400) %>%
lb$input() %>%
layer_points(fill := "red", data = reactive(mtcars[lb$selected(), ]))
marks

Vega marks.

Description

These functions create mark objects, corresponding to vega marks. Marks are leaves in the plot tree, and control the details of the final rendering. Marks are equivalent to the basic geoms in ggplot2 (e.g. point, line, polygon), where ggvis layers correspond to combinations of geoms and statistical transforms.

Usage

`emit_points(vis, props)`

`layer_points(vis, ..., data = NULL)`

`emit_images(vis, props)`

`layer_images(vis, ..., data = NULL)`

`emit_arcs(vis, props)`

`layer_arcs(vis, ..., data = NULL)`

`emit_ribbons(vis, props)`

`layer_ribbons(vis, ..., data = NULL)`

`emit_paths(vis, props)`

`layer_paths(vis, ..., data = NULL)`

`emit_rects(vis, props)`

`layer_rects(vis, ..., data = NULL)`

`emit_text(vis, props)`

`layer_text(vis, ..., data = NULL)`

Arguments

vis          Visualisation to modify
props, ...  A `props` object, named according to the properties listed below.
data        An optional dataset, if you want to override the usual data inheritance for this mark.
Details

Note that by supplying a fill property to `mark_path` will produce a filled property. `mark_point` is an alias to `mark_symbol`.

Properties

You can set the following mark properties:

- x The first (typically left-most) x-coordinate.
- x2 The second (typically right-most) x-coordinate.
- width The width of the mark (if supported).
- y The first (typically top-most) y-coordinate.
- y2 The second (typically bottom-most) y-coordinate.
- height The height of the mark (if supported).
- opacity The overall opacity.
- fill The fill color.
- fillOpacity The fill opacity
- stroke The stroke color.
- strokeWidth The stroke width, in pixels.
- strokeOpacity The stroke opacity.
- size [symbol] The pixel area of the symbol. For example in the case of circles, the radius is determined in part by the square root of the size value.
- shape [symbol] The symbol shape to use. One of circle (default), square, cross, diamond, triangle-up, or triangle-down (symbol only)
- innerRadius [arc] The inner radius of the arc, in pixels.
- outerRadius [arc] The outer radius of the arc, in pixels.
- startAngle [arc] The start angle of the arc, in radians.
- endAngle [arc] The end angle of the arc, in radians.
- tension [area, line] Depending on the interpolation type, sets the tension parameter.
- url [image] The URL from which to retrieve the image.
- align [image, text] The horizontal alignment of the object. One of left, right, center.
- baseline [image, text] The vertical alignment of the object. One of top, middle, bottom.
- text [text] The text to display.
- dx [text] The horizontal margin, in pixels, between the text label and its anchor point. The value is ignored if the align property is center.
- dy [text] The vertical margin, in pixels, between the text label and its anchor point. The value is ignored if the baseline property is middle.
- angle [text] The rotation angle of the text, in degrees.
• font [text] The typeface to set the text in (e.g., Helvetica Neue).
• fontSize [text] The font size, in pixels.
• fontWeight [text] The font weight (e.g., bold).
• fontStyle [text] The font style (e.g., italic).

To each property, you can assign any property object (prop) either locally (i.e. in the mark), or in a parent layer.

---

**padding**  
*Define padding.*

**Description**

Define padding.

**Usage**

padding(top = NULL, right = NULL, bottom = NULL, left = NULL)

**Arguments**

top, right, bottom, left  
Amount of padding on each border. Can either be a single number, "auto", or "strict"

**Examples**

p <- mtcars %>% ggvis(~wt, ~mpg) %>% layer_points()  
p %>% set_options(padding = padding())  
p %>% set_options(padding = padding(10, 10, 10, 10))

---

**prop**  
*Create a property.*

**Description**

Properties are used to describe the visual properties of marks. You create a single property definition with prop, and manage sets of named properties with props (which also provides shortcuts for creating the most common kind of properties)
Usage

```r
prop(
    property,
    x,
    scale = NULL,
    offset = NULL,
    mult = NULL,
    env = parent.frame(),
    event = NULL,
    label = NULL
)
```

`is.prop(x)`

`is.prop_constant(x)`

`is.prop_variable(x)`

`is.prop_reactive(x)`

Arguments

- **property**: A property, like "x", "x2", "y", "fill", and so on.
- **x**: The value of the property. This can be an atomic vector (a constant), a name or quoted call (a variable), a single-sided formula (a constant or variable depending on its contents), or a delayed reactive (which can be either variable or constant).
- **scale**: If `NULL`, automatically determine behavior by the kind of value (constant, variable, or reactive). If `TRUE` use the default scale associated with property. If `FALSE`, do not scale the value. Otherwise supply a string to select a custom scale. If `x` is an interactive input, then this defaults to the scale parameter of the input.
- **offset, mult**: Additive and multiplicate pixel offset used to adjust scaled values. These are useful if you want to place labels offset from points.
- **env**: If `x` is a quoted call this provides the environment in which to look for variables not in the data. You should not need this in ordinary operation.
- **event**: An event to which this property applies. One of "update", "enter", "exit", "hover", "brush".
- **label**: A label for this prop to use for reporting errors.

See Also

- `props` to manage multiple properties and to succinctly create the most common types.

Examples

```r
prop("x", 1)
prop("x", -1)
```
prop("fill", quote(cyl))
prop("fill", ~cyl)
prop("x", input_slider(0, 100))

# If you have a variable name as a string
var <- "cyl"
prop("x", as.name(var))

# Use a custom scale
prop("y", quote(cyl), scale = "y-2")

# Don't scale variable (i.e. it already makes sense in the visual space)
prop("fill", ~colour, scale = FALSE)

# Use a constant, but scaled
prop("x", 5, scale = TRUE)

# Use other events
prop("y", quote(cyl), scale = "y-2")

---

**props**

Manage a list of properties.

**Description**

props() provides a tool for concise creation of prop objects using a set of conventions designed to capture the most common use cases. If you need something less common, you'll need to use prop to access all possible options.

**Usage**

props(..., .props = NULL, inherit = TRUE, env = parent.frame())

is.ggvis_props(x)

**Arguments**

... A set of name-value pairs. The name should be a valid vega property. The first two unnamed components are taken to be x and y. Any additional unnamed components will raise an error.

.props When calling props from other functions, you'll often have a list of quoted function functions. You can pass that function to the .props argument instead of messing around with substitute. In other words, .props lets you opt out of the non-standard evaluation that props does.

inherit If TRUE, the defaults, will inherit from properties from the parent layer If FALSE, it will start from nothing.

env The environment in which to evaluate variable properties.

x an object to test for props-ness.
Heuristics

If the values are not already objects of class prop, props uses the following heuristics to when creating the prop:

- atomic vectors, e.g. \( x = 1 \): scaled = FALSE
- an interactive input, e.g. \( x = \text{input\_slider} \): scaled = FALSE
- a formula containing a single value, e.g. \( x \sim 1 \): scaled = TRUE
- a formula containing a name or expression, \( x \sim \text{mpg} \): scaled = TRUE

Non-standard evaluation

props uses non-standard evaluation in a slightly unusual way: if you provide a formula input, the LHS of the formula will provide the name of the component. In otherwise, \( \text{props}(x = y \sim 1) \) is the same as \( \text{props}(y \sim 1) \).

You can combine variables from the dataset and variables defined in the local environment: expressions will be evaluated in the environment which the formula was defined.

If you have the name of a variable in a string, see the props vignette for how to create the needed property mapping.

Enter, exit, hover, and update events

There are four different property events that the marks can use. These can, for example, be used to change the appearance of a mark when the mouse cursor is hovering over it: when the mark is hovered over, it uses the hover event, and when the mark isn’t hovered over, it uses the update event

- enter: This event is used by marks when they are added to a plot.
- update: This event is used by marks after they have entered, and also after they have been hovered over.
- exit: This event is used by marks as they are removed from a plot.
- hover: This event is used when the mouse cursor is over the mark.

You can specify the event for a property, by putting a period and the event after the property name. For example, \( \text{props}(\text{fill}.\text{update} := \"black\", \text{fill}.\text{hover} := \"red\") \) will make a mark have a black fill normally, and red fill when it is hovered over.

The default event is update, so if you run \( \text{props}(\text{fill} := \"red\") \), this is equivalent to \( \text{props}(\text{fill}.\text{update} := \"red\") \).

In practice, the enter and exit events are useful only when the update has a duration (and is therefore not instantaneous). The update event can be thought of as the "default" state.

Key property

In addition to the standard properties, there is a special optional property called key. This is useful for plots with dynamic data and smooth transitions: as the data changes, the key is used to tell the plot how the new data rows should be matched to the old data rows. Note that the key must be an unscaled value. Additionally, the key property doesn’t have a event, since it is independent of enter, update, exit, and hover events.
Properties

You can set the following mark properties:

- x The first (typically left-most) x-coordinate.
- x2 The second (typically right-most) x-coordinate.
- width The width of the mark (if supported).
- y The first (typically top-most) y-coordinate.
- y2 The second (typically bottom-most) y-coordinate.
- height The height of the mark (if supported).
- opacity The overall opacity.
- fill The fill color.
- fillOpacity The fill opacity
- stroke The stroke color.
- strokeWidth The stroke width, in pixels.
- strokeOpacity The stroke opacity.
- size [symbol] The pixel area of the symbol. For example in the case of circles, the radius is determined in part by the square root of the size value.
- shape [symbol] The symbol shape to use. One of circle (default), square, cross, diamond, triangle-up, or triangle-down (symbol only)
- innerRadius [arc] The inner radius of the arc, in pixels.
- outerRadius [arc] The outer radius of the arc, in pixels.
- startAngle [arc] The start angle of the arc, in radians.
- endAngle [arc] The end angle of the arc, in radians.
- tension [area, line] Depending on the interpolation type, sets the tension parameter.
- url [image] The URL from which to retrieve the image.
- align [image, text] The horizontal alignment of the object. One of left, right, center.
- baseline [image, text] The vertical alignment of the object. One of top, middle, bottom.
- text [text] The text to display.
- dx [text] The horizontal margin, in pixels, between the text label and its anchor point. The value is ignored if the align property is center.
- dy [text] The vertical margin, in pixels, between the text label and its anchor point. The value is ignored if the baseline property is middle.
- angle [text] The rotation angle of the text, in degrees.
- font [text] The typeface to set the text in (e.g., Helvetica Neue).
- fontSize [text] The font size, in pixels.
- fontWeight [text] The font weight (e.g., bold).
- fontStyle [text] The font style (e.g., italic).

To each property, you can assign any property object (prop) either locally (i.e. in the mark), or in a parent layer.
Examples

# Set to constant values
props(x := 1, y := 2)
# Map to variables in the dataset
props(x = ~mpg, y = ~cyl)
# Set to a constant value in the data space
props(x = 1, y = 1)
# Use an interactive slider
props(opacity := input_slider(0, 1))

# To control other settings (like custom scales, mult and offset)
# use a prop object
props(prop("x", "old", scale = "x", offset = -1))

# Red when hovered over, black otherwise (these are equivalent)
props(fill := "black", fill.hover := "red")
props(fill.update := "black", fill.hover := "red")

# Use a column called id as the key (for dynamic data)
props(key := ~id)

# Explicitly create prop objects. The following are equivalent:
props(fill = ~cyl)
props(fill.update = ~cyl)
props(prop("fill", ~cyl))
props(prop("fill", ~cyl, scale = "fill", event = "update"))

# Prop objects can be programmatically created and added:
property <- "fill"
expr <- parse(text = "wt/mpg")[[1]]
p <- prop(property, expr)
props(p)

# Using .props
props(.props = list(x = 1, y = 2))
props(.props = list(x = ~mpg, y = ~cyl))
props(.props = list(quote(x := ~mpg)))

prop_domain

Description
Property domain.

Usage
prop_domain(x, data)
Arguments

x    property to dispatch on
data  name of data set

resolution  Compute the "resolution" of a data vector.

Description

The resolution is is the smallest non-zero distance between adjacent values. If there is only one unique value, then the resolution is defined to be one.

Usage

resolution(x, zero = TRUE)

Arguments

x    numeric vector
zero  should a zero value be automatically included in the computation of resolution

Details

If x is an integer vector, then it is assumed to represent a discrete variable, and the resolution is 1.

Examples

resolution(1:10)
resolution((1:10) - 0.5)
resolution((1:10) - 0.5, FALSE)
resolution(c(1,2, 10, 20, 50))
resolution(as.integer(c(1, 10, 20, 50)))  # Returns 1

scaled_value  Create a scaled_value object

Description

These are for use with legends and axes.

Usage

scaled_value(scale, value)

Arguments

scale    The name of a scale, e.g., "x", "fill".
value    A value which will be transformed using the scale.
scales

Add a scale to a ggvis plot

Description

This creates a scale object for a given scale and variable type, and adds it to a ggvis plot. The scale object is populated with default settings, which depend on the scale (e.g. fill, x, opacity) and the type of variable (e.g. numeric, nominal, ordinal). Any settings that are passed in as arguments will override the defaults.

Arguments

vis        A ggvis object.
scale      The name of a scale, such as "x", "y", "fill", "stroke", etc.
type      A variable type. One of "numeric", "nominal", "ordinal", "logical", "datetime".
...       other arguments passed to the scale function. See the help for `scale_numeric`, `scale_ordinal` and `scale_datetime` for more details. For example, you might supply `trans = "log"` to create a log scale.
name      If NULL, the default, the scale name is the same as scale. Set this to a custom name to create multiple scales for stroke or fill, or (god forbid) a secondary y scale.

Scale selection

ggvis supports the following types of scales. Typical uses for each scale type are listed below:

- numeric For continuous numeric values.
- nominal For character vectors and factors.
- ordinal For ordered factors (these presently behave the same as nominal).
- logical For logical (TRUE/FALSE) values.
- datetime For dates and date-times.

Each type has a corresponding function: `scale_numeric`, `scale_nominal`, and so on.
The scale types for ggvis are mapped to scale types for Vega, which include "ordinal", "quantitative", and "time". See `ggvis_scale` for more details.

Given a scale and type, the range is selected based on the combination of the scale and type. For example, you get a different range of colours depending on whether the data is numeric, ordinal, or nominal. Some scales also set other properties. For example, nominal/ordinal position scales also add some padding so that points are spaced away from plot edges.

Not all combinations have an existing default scale. If you use a combination that does not have an existing combination, it may suggest you’re displaying the data in a suboptimal way. For example, there is no default for a numeric shape scale, because there’s no obvious way to map continuous values to discrete shapes.
Examples

```r
p <- mtcars %>%
ggvis(x = ~wt, y = ~mpg, fill = ~factor(cyl), stroke = ~hp) %>%
layer_points()

p %>% scale_numeric("x")
p %>% scale_numeric("stroke")
p %>% scale_nominal("fill")

# You can also supply additional arguments or override the defaults
p %>% scale_numeric("x", trans = "log")
p %>% scale_numeric("stroke", range = c("red", "blue"))
```

scale_datetime

Add a date-time scale to a ggvis object.

Description

A date/time scale controls the mapping of date and time variables to visual properties.

Usage

```r
scale_datetime(
  vis,
  property,
  domain = NULL,
  range = NULL,
  reverse = NULL,
  round = NULL,
  utc = NULL,
  clamp = NULL,
  nice = NULL,
  expand = NULL,
  name = property,
  label = NULL,
  override = NULL
)
```

Arguments

- `vis`: A ggvis object.
- `property`: The name of a property, such as "x", "y", "fill", "stroke", etc.
- `domain`: The domain of the scale, representing the set of data values. For ordinal scales, a character vector; for quantitative scales, a numeric vector of length two. Either value (but not both) may be NA, in which case domainMin or domainMax is set. For dynamic scales, this can also be a reactive which returns the appropriate type of vector.
The range of the scale, representing the set of visual values. For numeric values, the range can take the form of a two-element array with minimum and maximum values. For ordinal data, the range may by an array of desired output values, which are mapped to elements in the specified domain. The following range literals are also available: "width", "height", "shapes", "category10", "category20".

If true, flips the scale range.

If true, rounds numeric output values to integers. This can be helpful for snapping to the pixel grid.

If TRUE, uses UTC times. Default is FALSE.

If TRUE, values that exceed the data domain are clamped to either the minimum or maximum range value. Default is FALSE.

If specified, modifies the scale domain to use a more human-friendly value range. Should be a string indicating the desired time interval; legal values are "second", "minute", "hour", "day", "week", "month", or "year".

A multiplier for how much the scale should be expanded beyond the domain of the data. For example, if the data goes from 10 to 110, and expand is 0.05, then the resulting domain of the scale is 5 to 115. Set to 0 and use nice=FALSE if you want exact control over the domain.

Name of the scale, such as "x", "y", "fill", etc. Can also be an arbitrary name like "foo".

Label for the scale. Used for axis or legend titles.

Should the domain specified by this ggvis_scale object override other ggvis_scale objects for the same scale? Useful when domain is manually specified. For example, by default, the domain of the scale will contain the range of the data, but when this is TRUE, the specified domain will override, and the domain can be smaller than the range of the data. If FALSE, the domain will not behave this way. If left NULL, then it will be treated as TRUE whenever domain is non-NULL.

See Also

scales, scale_numeric, https://vega.github.io/vega/docs/scales/#time

Other scales: scale_numeric(), scale_ordinal()

Examples

set.seed(2934)
dat <- data.frame(
  time = as.Date("2013-07-01") + 1:100,
  value = seq(1, 10, length.out = 100) + rnorm(100)
)
p <- dat %>% ggvis(~time, ~value) %>% layer_points()

# Start and end on month boundaries
p %>% scale_datetime("x", nice = "month")
dist <- data.frame(times = as.POSIXct("2013-07-01", tz = "GMT") +
                   rnorm(200) * 60 * 60 * 24 * 7)
p <- dist %>% ggvis(x = ~times) %>% layer_histograms()
p

# Start and end on month boundaries
p %>% scale_datetime("x", nice = "month")
p %>% scale_datetime("x", utc = TRUE)

---

**scale_numeric**

Add a numeric scale to a ggvis object.

**Description**

A numeric (quantitative) scale controls the mapping of continuous variables to visual properties.

**Usage**

```
scale_numeric(
  vis,
  property,
  domain = NULL,
  range = NULL,
  reverse = NULL,
  round = NULL,
  trans = NULL,
  clamp = NULL,
  exponent = NULL,
  nice = NULL,
  zero = NULL,
  expand = NULL,
  name = property,
  label = NULL,
  override = NULL
)
```

**Arguments**

- **vis**
  A ggvis object.
- **property**
  The name of a visual property, such as "x", "y", "fill", "stroke". Note both x and x2 use the "x" scale (similarly for y and y2). fillOpacity, opacity and strokeOpacity use the "opacity" scale.
- **domain**
  The domain of the scale, representing the set of data values. For ordinal scales, a character vector; for quantitative scales, a numeric vector of length two. Either value (but not both) may be NA, in which case domainMin or domainMax is set. For dynamic scales, this can also be a reactive which returns the appropriate type of vector.
The range of the scale, representing the set of visual values. For numeric values, the range can take the form of a two-element array with minimum and maximum values. For ordinal data, the range may be an array of desired output values, which are mapped to elements in the specified domain. The following range literals are also available: "width", "height", "shapes", "category10", "category20".

If true, flips the scale range.

If true, rounds numeric output values to integers. This can be helpful for snapping to the pixel grid.

A scale transformation: one of "linear", "log", "pow", "sqrt", "quantile", "quantize", "threshold". Default is "linear".

If TRUE, values that exceed the data domain are clamped to either the minimum or maximum range value. Default is FALSE.

Sets the exponent of the scale transformation. For pow transform only.

If TRUE, modifies the scale domain to use a more human-friendly number range (e.g., 7 instead of 6.96). Default is FALSE.

If TRUE, ensures that a zero baseline value is included in the scale domain. This option is ignored for non-quantitative scales. Default is FALSE.

A multiplier for how much the scale should be expanded beyond the domain of the data. For example, if the data goes from 10 to 110, and expand is 0.05, then the resulting domain of the scale is 5 to 115. Set to 0 and use nice=FALSE if you want exact control over the domain. If left NULL, behavior will depend on the scale type. For positional scales (x and y), expand will default to 0.05. For other scales, it will default to 0.

Name of the scale, such as "x", "y", "fill", etc. Can also be an arbitrary name like "foo".

Label for the scale. Used for axis or legend titles.

Should the domain specified by this ggvis_scale object override other ggvis_scale objects for the same scale? Useful when domain is manually specified. For example, by default, the domain of the scale will contain the range of the data, but when this is TRUE, the specified domain will override, and the domain can be smaller than the range of the data. If FALSE, the domain will not behave this way. If left NULL, then it will be treated as TRUE whenever domain is non-NULL.

The default values for most of the arguments is NULL. When the plot is created, these NULL values will be replaced with default values, as indicated below.

See Also

scales, scale_ordinal, https://vega.github.io/vega/docs/scales/#quantitative

Other scales: scale_datetime(), scale_ordinal()
scale_ordinal

Add a ordinal, nominal, or logical scale to a ggvis object.

Description

Ordinal, nominal, and logical scales are all categorical, and are treated similarly by ggvis.

Usage

scale_ordinal(
  vis,
  property,
  domain = NULL,
  range = NULL,
  reverse = NULL,
  round = NULL,
  points = NULL,
  padding = NULL,
  sort = NULL,
  name = property,
  label = NULL,
  override = NULL
)

scale_nominal(
  vis,
  property,
  domain = NULL,
  range = NULL,
scale_ordinal

reverse = NULL,
round = NULL,
points = NULL,
padding = NULL,
name = property,
label = NULL,
override = NULL
)

scale_logical(
    vis,
      property,
      domain = NULL,
      range = NULL,
      reverse = NULL,
      round = NULL,
      points = NULL,
padding = NULL,
      sort = NULL,
      name = property,
      label = NULL,
      override = NULL
)

Arguments

vis A ggvis object.
property The name of a property, such as "x", "y", "fill", "stroke", etc.
domain The domain of the scale, representing the set of data values. For ordinal scales, a character vector; for quantitative scales, a numeric vector of length two. Either value (but not both) may be NA, in which case domainMin or domainMax is set. For dynamic scales, this can also be a reactive which returns the appropriate type of vector.
range The range of the scale, representing the set of visual values. For numeric values, the range can take the form of a two-element array with minimum and maximum values. For ordinal data, the range may by an array of desired output values, which are mapped to elements in the specified domain. The following range literals are also available: "width", "height", "shapes", "category10", "category20".
reverse If true, flips the scale range.
round If true, rounds numeric output values to integers. This can be helpful for snapping to the pixel grid.
points If TRUE (default), distributes the ordinal values over a quantitative range at uniformly spaced points. The spacing of the points can be adjusted using the padding property. If FALSE, the ordinal scale will construct evenly-spaced bands, rather than points. Note that if any mark is added with a band() prop, then the scale for that prop will automatically have points set to FALSE.
padding

Applies spacing among ordinal elements in the scale range. The actual effect depends on how the scale is configured. If the points parameter is true, the padding value is interpreted as a multiple of the spacing between points. A reasonable value is 1.0, such that the first and last point will be offset from the minimum and maximum value by half the distance between points. Otherwise, padding is typically in the range [0, 1] and corresponds to the fraction of space in the range interval to allocate to padding. A value of 0.5 means that the range band width will be equal to the padding width. For positional (x and y) scales, the default padding is 0.1. For other scales, the default padding is 0.5.

sort

If TRUE, the values in the scale domain will be sorted according to their natural order. Default is FALSE.

name

Name of the scale, such as "x", "y", "fill", etc. Can also be an arbitrary name like "foo".

label

Label for the scale. Used for axis or legend titles.

override

Should the domain specified by this ggvis_scale object override other ggvis_scale objects for the same scale? Useful when domain is manually specified. For example, by default, the domain of the scale will contain the range of the data, but when this is TRUE, the specified domain will override, and the domain can be smaller than the range of the data. If FALSE, the domain will not behave this way. If left NULL, then it will be treated as TRUE whenever domain is non-NULL.

See Also


Other scales: scale_datetime(), scale_numeric()

Examples

p <- PlantGrowth %>% ggvis(~group, ~weight) %>% layer_points()

p

p %>% scale_nominal("x", padding = 0)
p %>% scale_nominal("x", padding = 1)

p %>% scale_nominal("x", reverse = TRUE)

p <- ToothGrowth %>% group_by(supp) %>%
  ggvis(~len, fill = ~supp) %>%
  layer_histograms(width = 4, stack = TRUE)

# Control range of fill scale
p %>% scale_nominal("fill", range = c("pink", "lightblue"))

# There's no default range when the data is categorical but the output range
# is continuous, as in the case of opacity. In these cases, you can
# manually specify the range for the scale.
mtcars %>% ggvis(x = ~wt, y = ~mpg, opacity = ~factor(cyl)) %>%
  layer_points() %>%
  scale_nominal("opacity", range = c(0.2, 1))
set_options

Description

Set options for a ggvis plot

Usage

set_options(
  vis,
  width = NULL,
  height = NULL,
  keep_aspect = NULL,
  resizable = NULL,
  padding = NULL,
  duration = NULL,
  renderer = NULL,
  hover_duration = NULL
)

Arguments

vis          Visualisation to modify
width, height Width and height of plot, in pixels. Default is 600x400. width or height can also be "auto", in which case the plot will size to fit in the containing div. This is useful only in a Shiny app or custom HTML output. Note that height="auto" should only be used when the plot is placed within a div that has a fixed height; if not, automatic height will not work, due to the way that web browsers do vertical layout.
keep_aspect  Should the aspect ratio be preserved? The default value is FALSE, or the value of getOption("ggvis.keep_aspect"), if it is set.
resizable    If TRUE, allow the user to resize the plot. The default value is TRUE, or the value of getOption("ggvis.resizable"), if it is set. Not compatible when width or height is "auto".
padding      A padding object specifying padding on the top, right, left, and bottom. See padding.
duration     Duration of transitions, in milliseconds.
renderer     The renderer to use in the browser. Can be "canvas" or "svg" (the default).
hover_duration The amount of time for hover transitions, in milliseconds.

See Also

ggetOption and options, for getting and setting global options.
default_options to see the default options.
Examples

```r
mtcars %>%
ggvis(~wt, ~mpg) %>%
layer_points() %>%
set_options(width = 300, height = 200, padding = padding(10, 10, 10, 10))

# Display the default options
str(default_options())
```

---

**set_scale_label**  
*Set the label for a scale*

**Description**

Set the label for a scale

**Usage**

```r
set_scale_label(vis, scale, label)
```

**Arguments**

- `vis`: A ggvis object.
- `scale`: The name of a scale, like "x".
- `label`: Text to use for the label.

---

**shiny-ggvis**  
*Connect a ggvis graphic to a shiny app.*

**Description**

Embedding ggvis in a shiny app is easy. You need to make a place for it in your `ui.r` with `ggvisOutput`, and tell your `server.r` where to draw it with `bind_shiny`. It’s easiest to learn by example: there are many shiny apps in `demo/apps/` that you can learn from.

**Usage**

```r
bind_shiny(
  vis,
  plot_id,
  controls_id = NULL,
  ...,
  session = shiny::getDefaultReactiveDomain()
)
```
bind_shiny_ui(vis, controls_id, session = shiny::getDefaultReactiveDomain())

ggvisOutput(plot_id = rand_id("plot_id"))

Arguments

vis    A ggvis object, or a reactive expression that returns a ggvis object.
plot_id unique identifier to use for the div containing the ggvis plot.
controls_id Unique identifier for controls div.
... Other arguments passed to as.vega.
session A Shiny session object.

Client-side

In your UI, use ggvisOutput() in ui.r to insert an html placeholder for the plot.

If you’re going to be using interactive controls generated by ggvis, use renderUI() to add a placeholder. By convention, if the id of plot placeholder is called "plot", call the controls placeholder "plot_ui".

Server-side

When you run ggvis plot interactively, it is automatically plotted because it triggers the default print method. In shiny apps, you need to explicitly render the plot to a specific placeholder with bind_shiny:

p %>% bind_shiny("plot")

If the plot has controls, and you’ve reserved space for them in the UI, supply the name of the placeholder as the third argument:

p %>% bind_shiny("plot", "plot_ui")

Examples

```r
# Simplest possible app:
library(shiny)
runApp(list(
  ui = bootstrapPage(
    ggvisOutput("p"),
    uiOutput("p_ui")
  ),
  server = function(..., session) {
    mtcars %>%
      ggvis(~wt, ~mpg) %>%
      layer_points() %>%
      layer_smooths(span = input_slider(0, 1)) %>%
      bind_shiny("p", "p_ui")
  })
)```

```
show_spec

Print out the vega plot specification

Description
Print out the vega plot specification

Usage
show_spec(vis, pieces = NULL)

Arguments
vis Visualisation to print
pieces Optional, a character or numeric vector used to pull out selected pieces of the spec

Examples
base <- mtcars %>% ggvis(~mpg, ~wt) %>% layer_points()  
base %>% show_spec()  
base %>% show_spec("scales")

show_tooltip

Send a message to the client to show or hide a tooltip

Description
Send a message to the client to show or hide a tooltip

Usage
show_tooltip(session, l = 0, t = 0, html = "")
hide_tooltip(session)

Arguments
session A Shiny session object.
l Pixel location of left edge of tooltip (relative to page)
t Pixel location of top edge of tooltip (relative to page)
html HTML to display in the tooltip box.
Create a page with a sidebar

Description
This creates a page with a sidebar, where the sidebar moves to the bottom when the width goes below a particular value.

Usage

```r
sidebarBottomPage(sidebarPanel, mainPanel, shiny_headers = TRUE)
```

```r
sidebarBottomPanel(...)
```

```r
mainTopPanel(...)
```

Arguments

- `sidebarPanel` The `sidebarBottomPanel` containing input controls.
- `mainPanel` The `mainTopPanel` containing the main content.
- `shiny_headers` Should Shiny headers be embedded in the page? This should be TRUE for interactive/dynamic pages, FALSE for static pages.
- `...` Additional tags.

Examples

```r
sidebarBottomPage(sidebarBottomPanel(), mainTopPanel())
```

---

singular

Use singular when you want constant x or y position.

Usage

```r
singular()
```

```r
scale_singular(
    vis,
    property,
    name = property,
    label = name,
    points = TRUE,
```
Arguments

vis
A ggvis object.

property
The name of a property, such as "x", "y", "fill", "stroke", etc.

name
Name of the scale, such as "x", "y", "fill", etc. Can also be an arbitrary name like "foo".

label
Label for the scale. Used for axis or legend titles.

points
If TRUE (default), distributes the ordinal values over a quantitative range at uniformly spaced points. The spacing of the points can be adjusted using the padding property. If FALSE, the ordinal scale will construct evenly-spaced bands, rather than points. Note that if any mark is added with a band() prop, then the scale for that prop will automatically have points set to FALSE.

domain
The domain of the scale, representing the set of data values. For ordinal scales, a character vector; for quantitative scales, a numeric vector of length two. Either value (but not both) may be NA, in which case domainMin or domainMax is set. For dynamic scales, this can also be a reactive which returns the appropriate type of vector.

override
Should the domain specified by this ggvis_scale object override other ggvis_scale objects for the same scale? Useful when domain is manually specified. For example, by default, the domain of the scale will contain the range of the data, but when this is TRUE, the specified domain will override, and the domain can be smaller than the range of the data. If FALSE, the domain will not behave this way. If left NULL, then it will be treated as TRUE whenever domain is non-NULL.

Examples

```
mtcars %>% ggvis("", ~mpg) %>%
  layer_points() %>%
  scale_nominal("x") %>%
  add_axis("x", title = "", tick_size_major = 0)
```

# OR
```
mtcars %>% ggvis("", ~mpg) %>%
  layer_points() %>%
  scale_singular("x")
```

# OR, even simpler
```
mtcars %>% ggvis(singular(), ~mpg) %>% layer_points()
```

# In the other direction:
```
mtcars %>% ggvis(~mpg, singular()) %>% layer_points()
```
vector_type

Determine the "type" of a vector

Description

The `vector_type` collapses down the class of base vectors into something useful more for visualisation, yielding one of "datetime", "numeric", "ordinal", "nominal" or "logical".

Usage

`vector_type(x)`

Arguments

- `x`: a vector

See Also

- `default_scale`, which uses this when picking the default scale.

vega_data_parser

Determine the vega data type for a vector

Description

This is used to specify the data type so that the appropriate parser is used when Vega receives the data.

Usage

`vega_data_parser(x)`

Arguments

- `x`: A vector.
waggle

Waggle back and forth between two numbers

Description
Waggle back and forth between two numbers

Usage
waggle(min, max, value = (min + max)/2, step = (max - min)/50, fps = 10)

Arguments
- **min**: A minimum value.
- **max**: A maximum value.
- **value**: Starting value. Defaults to half-way between min and max.
- **step**: How much value changes at each frame. Defaults to 50 steps between min and max so it takes 5 seconds to waggle once.
- **fps**: number of frames per second.

Examples
```r
span <- waggle(0.2, 1)
mtcars %>% ggvis(~mpg, ~wt) %>%
  layer_points() %>%
  layer_smooths(span = span)
```

go

zero_range

Determine if range of vector is close to zero, with a specified tolerance

Description
The machine epsilon is the difference between 1.0 and the next number that can be represented by the machine. By default, this function uses epsilon * 100 as the tolerance. First it scales the values so that they have a mean of 1, and then it checks if the difference between them is larger than the tolerance.

Usage
zero_range(x, tol = .Machine$double.eps * 100)

Arguments
- **x**: numeric range: vector of length 2
- **tol**: A value specifying the tolerance. Defaults to .Machine$double.eps * 100.
Value

logical TRUE if the relative difference of the endpoints of the range are not distinguishable from 0.

Examples

es <- .Machine$double.eps
zero_range(c(1, 1 + eps))  # TRUE
zero_range(c(1, 1 + 99 * eps))  # TRUE
zero_range(c(1, 1 + 101 * eps))  # FALSE - Crossed the tol threshold
zero_range(c(1, 1 + 2 * eps), tol = eps)  # FALSE - Changed tol

# Scaling up or down all the values has no effect since the values
# are rescaled to 1 before checking against tol
zero_range(100000 * c(1, 1 + eps))  # TRUE
zero_range(100000 * c(1, 1 + 200 * eps))  # FALSE
zero_range(.00001 * c(1, 1 + eps))  # TRUE
zero_range(.00001 * c(1, 1 + 200 * eps))  # FALSE

# NA values
zero_range(c(1, NA))  # NA
zero_range(c(1, NaN))  # NA

# Infinite values
zero_range(c(1, Inf))  # FALSE
zero_range(c(-Inf, Inf))  # FALSE
zero_range(c(Inf, Inf))  # TRUE

### Pipe graphics

Description

Like dplyr, ggvis also uses the pipe function, %>% to turn function composition into a series of
imperative statements.

Arguments

lhs, rhs   A visualisation and a function to apply to it

Examples

# Instead of
layer_points(ggvis(mtcars, ~mpg, ~wt))
# you can write
mtcars %>% ggvis(~mpg, ~wt) %>% layer_points()
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